

Research Notes

Vol. II

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8/26/53

Beginning again.

1. Purification of HFT stocks.

Testing by picking single colonies - mixed in HFT and med. 1 loop at dist 50 cm 20 second and spotting loop on indicator.

N16 - 10 colonies picked 7/10 colonies give strong reaction
stocks made of #1 (1, 2, 3, 5, 7, 8, 10)

N17 ① 10 colonies picked 10/10 gave no reaction

② 10 colonies picked 10/10 gave no reaction

9/16/53 all remaining gal⁻ in 1 tested - N12 suggestive. Re v

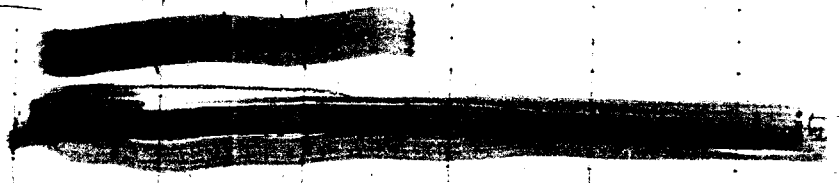
2. Use of med. HFT to find auxotrophic mutation which will be transmissible.

in 7' HFT gal⁻ 1 used - Del 1-100 from original probably.

① cells from unselected low culture sedimented - dispd. in 1 ml saline 0.5 ml cell + 1.0 ml HFT - mixed and diluted out cells plated - a dilution of cell grown 12 hrs + and then per run on grown cell.

a. cells used gal⁻ prototyp gal⁺ - prototyp - no evidence of gal⁺ transmissible plating - 1 probably too dilute? both cases.

b. per run.



1924 lysate made - apparent clearing after 45" exposure - viscosity to the lysate -

- 0.1 ml of lysate / 578 on EM13(0) - gave 354 plaques / $\frac{1}{2}$ plot = 2.8×10^4 / ml

Against 2785
 no odd - 9
 0.1 1924 - 17

Plaques appear to be of two sizes?

2281 vs 750 λ to see if HET property is 90% periodicity a property of λ or in general.

2 Transductions
 2281. no odd 5
 0.1 750 λ - 364

1810 no odd 2
 750 λ - 2 119

24 pop picked from each for purification and HET examination
 apparently none found HET

Checking on 2251

$2805 = (2319) \times 2251$ on EM13 λ .

	(-)
7.	244
2.	197
3.	319
4.	183
5.	208
6.	283
	<hr/> 1434

0.4 / 1434 (-)

1673 gal - produced by mat. Previous reference page 206

1673 gal - #5 = 2312 - possibly a non transducible locus

1. Recheck - not transduced by NI, N16, S18, S92(+), K-12, λ 9/17/52
 in confirmation and extension of previous work. HFT λ - no transduction by K-12

2. Made λ - lysate made - tested 9/13/52

	750	2175	811	2070 = gal
0.1 ml lys 1/2 plate	37/4	7/8	74/17	121/3
control 1/2 plate				

some suggestion that lysate inhibitory to spnt. pop in phage 1/2 of plate.

By \pm test a new locus - B₂ lysate test gal?

9/11/52

1673 gal - #11 = W2318 - another possible non transducible locus.

<u>Rekt.</u>	<u>λ Source</u>	<u>Result</u>	<u>Comment</u>
1.	NI (1-10)	no trans.	area of lysate = spnt ridge
2.	N16 (1-10)	" "	" " " " c. 20000 plaques.
3.	S18 (1-10)	" "	area " " " " spnt ridge
4.	2175 & 750(+)	" "	" " " " c. 30 plaques.

2062 - attempted phage detection - HFT λ 's on D(0)

1.	NI and	no colonies	↓ since pr - gal probably > 1% transduction likely
2.	N16	" "	
3.	N7	" "	
4.	S18	" "	
5.	2175 & 750(+)	" "	

9/13/53

1573 gal - #11

2318 X 811

controls ok

- 1. 32
 - 2. 45
 - 3. 49
 - 4. 52
 - 5. 62
 - 6. 76
- 286 (1) 0/286

9/16/53

2251 lysate - Is crossability of 2251 (578) really so or an S^r ductum?

Check against 578 on B gal and B gal SM

- 1. B gal - 1/8 plate = 65 pop = 520
- 2. B gal SM = 0 = 0
- 3. Replica B gal on B gal SM - no conspicuous growth - Couple minute colonies

2238 lysate #2 tested against 2175, 578

- 2175 1. control half plate = 7
- 2. lysate " " " " = 48 - pupae picked and checked.

- 578 1. control half plate = 7
- 2. lysate " " " " = 8 (lysis)

2312 lysate - (2312 = 1673 gal - #5) tested against 2175

this lysate also does not appear to transmit 2175⁻ previously found. also - does not appear to + other gal. - see pg. 220
lysate appears inhibitory to 2175. Rechecked in adipt. type cft.

Broth Control = 19
Associated lysis = 28 12 per cent - see stable (+) after 2.

2312 used X⁺ - This is second time - Culture (X⁺) used for lysate above discarded by mistake.

9/17/53

2312 X 811 - controls ok EMB gel

S gel	(+)	(-)
1.	0	30
2.	0	14
3.	0	24
4.	0	18
5.	0	28
	0	114

See also page 206

1/14
0/114
1/128

2238 transductions see page 221
2175222Y

24 phages picked

1st 5	2nd	3rd
8/24 +	7/24 +	9/24 +

518 by 2238A

Spread plates

1. no add	19
2. 2238A	26

811 by 2238A

1. no add	46
2. 2238A	55

} don't go?

750 by 2238A

1. no add	4
2. 2238A	24

~~all spread plates negative~~
2nd 3rd
11/24 + 13/24 +

2312 X 902 - controls ok
EMB gel

4 plates (-) = 118 = 66 = 528 X 4 = 2112
(+) = confirmed on slt on
EMB gel.

18/
2112(-)

9/20/53

ST8. Adsorption of HFT 2⁻ (3rd batch)

1st (+) Total cells 1.3 x 10¹⁰
 39 64 x 4 = 256 dil: 10⁶ x 10

= $\frac{64 \times 256}{1312 \times 3900} = 3\%$ transferred

2nd (+) 64 x 4 = 256
 7 276 / 7.00 = 3.9%

2nd 0.03 / 7.00 = 2.7%

8 (+) colonies picked and streaked out 2 times
 Pop. colonies were picked and cross brushed on ST8, and streaked out

Colony	/ST8	simult. streaked		4 Colonies picked from each subsequent subculture / ST8			
		(+)	(-)	(+)	(-)	(+)	(-)
1.	str. lytic	used	+	+	+	+	+
2.	"	"	+	+	+	+	+
3.	"	"	+	+	+	+	+
4.	non lytic	"	-	-	-	-	-
5.	str. lytic	"	+	+	+	+	+
6.	wk. lytic	"	+	+	+	+	+
7.	sh. lytic	"	+	+	+	+	+
8.	wk lytic	"	+	+	+	+	+

all of shil. segregating

9/24/53

Assay of lysates - Lumenly

K-12 lysate 5/30/53

<u>750</u>	<u>Lysate</u>	<u>No. plaques</u>	Δ
1.	0	2	0
2.	0.025	23	21
3.	0.05	51	49
4.	0.1	114	112
5.	0.15	185	183
6.	0.2	144	142
7.	0.3	132	
0.	0.5	70	

Comparison

λ	750	811
0.025	21	21
0.05	49	42
0.10	112	78
0.15	183	119
0.20	142	102

811

			Δ
1.	0	99	0
2.	0.025	104	5
3.	0.05	141	42
4.	0.1	177	78
5.	0.15	218	119
6.	0.2	261	162
7.	0.3	231	132

1027 - A second try - Lwoff attempt

1. Supernat. broth before inad. -
2. Centrifuge (after 3 hours inc.) of inad. culture -

checked 1516

7 plaques.

Does sub-transduce only and vice versa?

In case usual technique allows inhib. of trans. between adsorption type experiment done - "no add" plate run to efflu. elypt broth used

750	no add	1
	811 λ -5	9
811	no add	162
	750 λ -2	201
578	no add	740
	750 λ -2	73
1924	no add	33
	750 λ -2	43

Number of Papillae Per Plate

300
200
100

Assay of K-12 Lysate
on W750 Gal.²

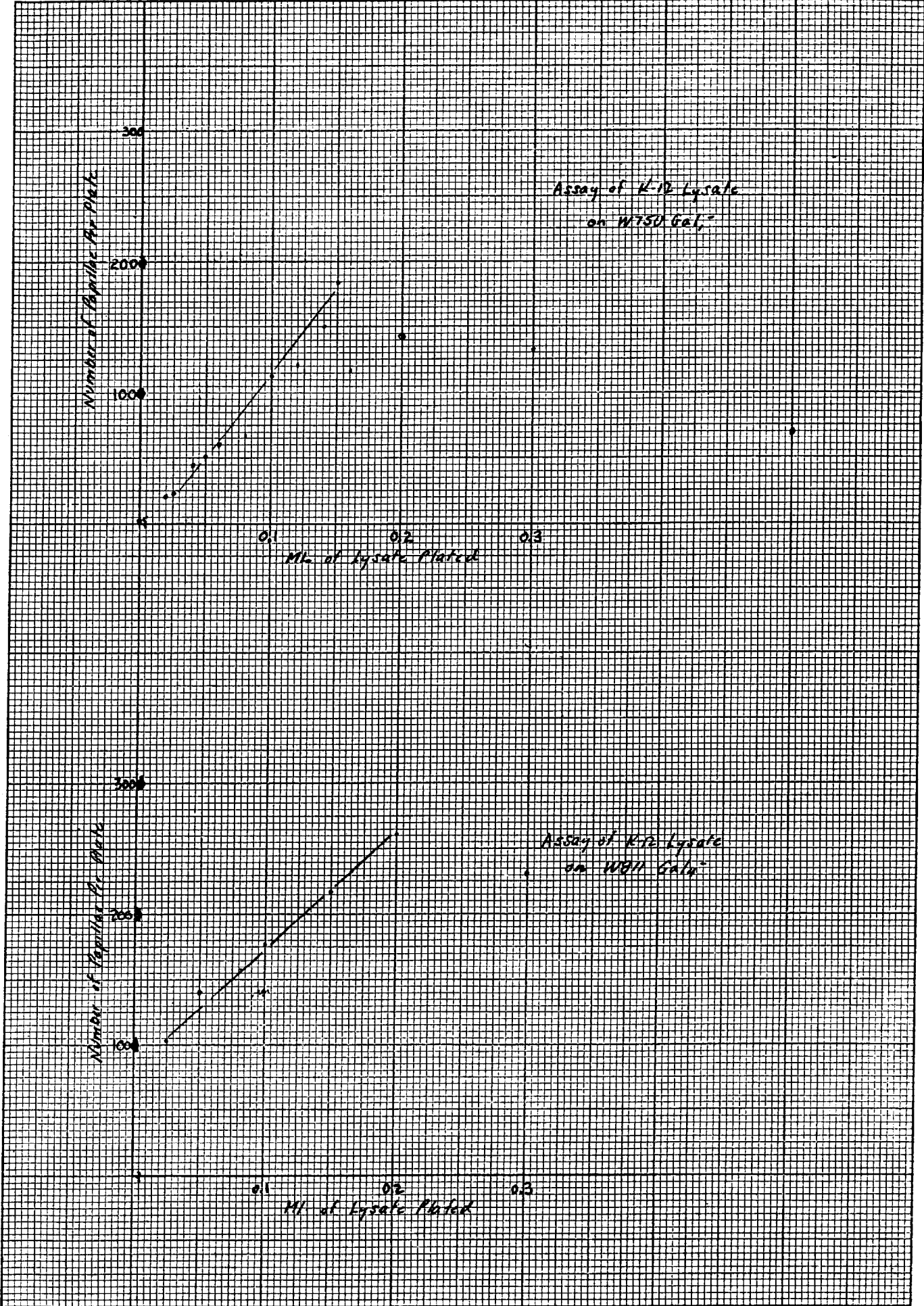
0.1 0.2 0.3
Ml. of Lysate Plated

Number of Papillae Per Plate

300
200
100

Assay of K12 Lysate
on W811 Gal.²

0.1 0.2 0.3
Ml. of Lysate Plated



9/24/53

2312 - a double minus - gal_2^- and gal_1^- outside the region of transduction? Cross and examine if possible to see if two classes of gal^- - can be recovered. gal_2^- , gal_1^- in which gal_1^- does ~~not~~ transduce gal_2^- -

1655x 2312 controls etc.

	(+)	(-)
1.	2	0
2.	3	0
3.	2	0
4.	7	0
5.	3	0
	<u>17</u>	<u>0</u>

Repeat using pr addition to medium -

8th adsorpt. in of K-12 λ 5/30 - 26×10^{10} (see W. H. G. (4)) + orig p 224

1. 0.8ml cell + 0.8ml K-12 λ \rightarrow $10^7, 10^6, 10^5, 10^4$ O.D. = 272, 347 = $310 \times 10 \times 10^7 = 3.1 \times 10^{10}$ cells/ml
 O.D. = 108 papillae
 net $\frac{41}{67}$ birth control = 41
2. 0.7ml cell + 0.7ml K-12 λ
 = 150 papillae
 net $\frac{108}{42}$ 67
3. 0.6ml cell + 0.6ml K-12 λ
 = 132 papillae
 net $\frac{109}{42}$ 109
- no adsorption 91

67
78

2297 E K-12 λ

1/2 plate = control
 1/2 " = 0.1ml K-12 λ

36
66
27

3.6
x6
216
216
3.6

1st ab. = 3.1×10^{10} adsorpt
 2nd ab. = 2.7×10^{10} adsorpt
 (3.8 x 10¹⁰)

5.8 x 10¹⁰ adsorpt
 3.1 x 10¹⁰ cells =
 are 2.9 x 10¹⁰ cells

slight diff. if at all in vial - suggestion of inhibition by side. λ around edge.

9/27/53

Adsorption K-12/5780

750 cell assay $10^2, 10^4, 10^6, 10^7 = 122, 127$ $125 \times 10 \times 10^7 = 1.25 \times 10^{10}$

centrifuged resuspended in orig. volume

- 1. 1.0 ml cells + 1.0 ml K-12 = 0.1 ml = 45
 5% 0.1 ml both exposed = 3
 net = 42
- 2. 0.9 ml + 0.9 ml 0.1 = 159 net. 156
- 3. 0.8 + 0.8 0.1 = 159 156

275 cell assay $10^2, 10^4, 10^6, 10^7 = 99, 130 = 115 \times 10 \times 10^7 = 1.15 \times 10^{10}$

- 1. 1.0 ml + 1.0 ml K-12 = 52 both control = 20 net = 32
- 2. 0.9 ml + 0.9 ml K-12 = 103 net = 83
- 3. 0.8 ml + = 97 net = 77

Cross

1655 x 2312 Repeat Protein added created 1655 on cross for controls etc.

	(+)	(-)	
1.	2	0	
2.	2	0	
3.	0	0	
4.	1	0	
5.	2	2(?)	} check
6.	1	1	
	<u>8</u>	<u>3</u>	

Strains of 2312/+, 2076/+, } made after testing 8 colonies of each to see if are lysogenic.

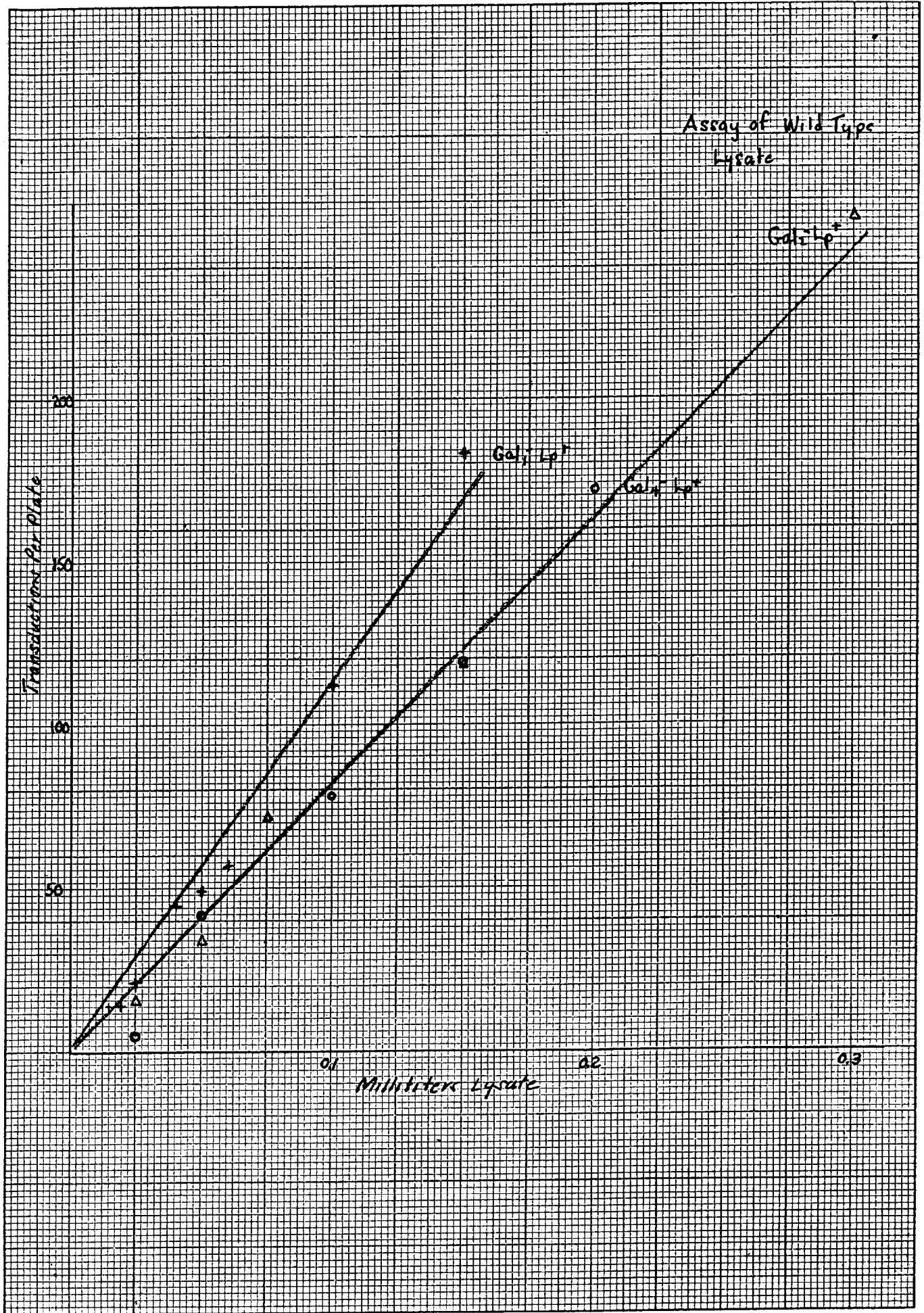
1436 look its st

✓ for gal allele

<u>1/11</u>	<u>1/16</u>	<u>st</u>
++	++	00

apparently still gal-





9/28/53

Adaptation K. P. 1/30
 150 *Adaptation* +

150 transducer with 90% to isolate NI HFT - also variable no

1. no add half of plate = 1
 2. 0.1 ml 912g = 126 pick 24 and test

2175 ~~Adaptation~~ Linearity of assay 16-12 5/30

cell assay 10^6 568, 450 = $509 \times 10 \times 10^6$ $5.09 \times 10^9/m$

	no. pup	Δ
1. no add	18	0
2. 0.025M	34	16
3. 0.05	52	34
3. 0.075	90	72
4. 0.100		
5. 0.15	137	119
6. 0.2		
7. 0.3	273	255

2175 - ductin versus variable cell no.

cell assay 10^7 = 257, 27, = $237 \times 10 \times 10^7$ = 2.37×10^{10}

Cell mass	Dilution	Sp. pup	trans. plate	Δ
2.4×10^9	und.	23	83	60
1.2×10^9	1-2	22	106	84
6×10^8	1-4	13	104	91
3×10^8	1-8	18	129	111
2.4×10^8	1-10 (?)	17	112	85
1.2×10^8	1-20	14	135	121
6×10^7	1-40	14	115	101
3×10^7	1-80 (?)	14	92	
2.4×10^7	1-100	19	109	90
1.2×10^7	1-200	15	93	78
6×10^6	1-400	16	87	71
3×10^6	1-800	8	83	75
2.4×10^6	1-1000	11	84	73

578 Lytic ?

c. 6×10^9 - 10^{10} cells 578 exposed to 1.0 ml K-16 5/30 twice

centrifuged after each exposure - supernat discarded
 after second adapt. - Resuspended in MSB (10ml)
 with anaerobe - 5 hours - non-turbid

Inoculated 150 (B₀) after ~~centrifugation~~ centrifugation

control $\frac{1}{2}$ = 10
 0.1 ml lytic = 12
 1/2175 on EM8 goe }
 1/2062 D(0) }
 no pup }
 either half of plate }
 $10^8 = 3310$ 3.3×10^6
 $10^7 = 150 \times 16 = 2400 \times 10^7 = 2.4 \times 10^{10}$

10/1/53

2281 transductions

- 1. 902 λ = 0
- 2. K-12 λ = 21
- 3. 750 λ = 252
- 4. 2238 λ = 19
- 5. 871 λ = 97

} picked and checked -

2281 K12	2281 750	2281 871
12/18 + 4" alt.	24/24 + 4" alt.	20/24 + 4" alt.
19 ² /1	22/24 + 3	23/24

1485 lytic λ

7. c. 5×10^2 10^{10} cells exposed for adaptation (ultraviolet) to 871 λ -9 twice centrifuged resuspended in NSB - Equivalent no cells both treated, used as control - incubated with aeration - c. 4.5 hours. After 5 hours phage tube transparent but dark, control turbid.

Assay

$$\left. \begin{aligned} 10^7 \text{ dil} &= \frac{1}{16} \times 157 = 2400 \times 10^7 = 2.4 \times 10^{10} \\ 10^8 \text{ dil} &= 294 = 294 \times 10^8 = 2.9 \times 10^{10} \end{aligned} \right\}$$

Transducing action

	control % plate	% plate = 0.1 m
750	3	2
578	9	8
2175	7	8

10/6/53

Segments from 2281 - Picked after 2 (+) picking and checked.

2281 transcribed by K-12

Segment	Sp	S18	N12	Alk
1.	r	+	0	2-
2.	r	0 or +		
3.	s	+	+	contaminated with +?
4.	r	+	0	2-
5.	r	+	0	2-
6.	s	-	+	?
7.	s	+	+	contaminated with +?
8.	s	+	0	2-
9.	s	+	0	..
10.	s	+	0	..
11.	r	+	0	..
12.	s	0	0	?

7S III 8
5r III 4

$$\frac{5+9}{12+16} = \frac{14}{28} = 50$$

Added to 233
for correlation of lysogenicity
and transduction in table summary

10/20/53
What are?
Dumbly segments

2281 ± 750 (-) seq -

disregard

Seq #	LP	/N7	/N16	Amplitude
1.	r	0	0	?
2.	r	0	0	?
3.	r	0	+	1-
4.	r	0	+	?
5.	s	0	+	1-
6.				
7.	r	0	0	?
8.	r	+3	+3	?
9.	r	+ 15	0	2-
10.	r	0	+	1-
11.	r	0	0	?
12.	r	0	0	?
13.	r	+ 5	+	?
14.	r	0	0	?
15.	r	0	0	?
16.	r	0	0	?
17.	r	0	0	?
18.	r	0	0	?
19.	r	0	0	?
20.	r	0	0	?
21.	r	0	+	1-
22.	r	0	0	?
23.	r	0	0	?
24.	r	0	0	?

cont'd?

mix
up
1st time

15
22 v

This
is a
probably not
a case
of
G₁ - G₂

$$\frac{22+16}{23+21} = \frac{38}{44}$$

22014011

Seq

Seq #	hp	NIL	Sit	Xble
1.	v	0 3	+	2
2.	r	0 3	+	2
3.	r	+	o	4
4.	r	+	o	4
5.	v	0	+	2
6.	s	0 4 5	+	2
7.	s	0 4 5	+	2
8.	v	0 4 5	+	2
9.	r	0 4 5	+	2
10.	r	0 4 5	+	2
11.	s	0 4 5	+	2
12.	s	0 4 5	+	2
13.	r	0 +	→	-
14.	r	0 4	+	2
15.	r	+	o	4
16.	r	0 4 5	+	2
17.	r	0 4 5	+	2
18.	v	0 4 5	+	2
19.	r	0 4 5	+	2
20.	v	0 4 5	+	2

discard

33
10v

↑
Calc

↑
Calc

16 2
34

$$\frac{16 + 17}{19 + 21} = \frac{33}{40}$$

750C902

In Search of HFT

once related in HFT test

W24 6/18/54

Culture and lysate has referred to as NA-4

	N/15	N/16	Milke	1/2175	1/750
1	0	+	gal ₁ -	0	0
2	0	+		↓	
3	0	+		↓	
4	0	+		+	
5	+	0	gal ₂ -	↓	
6	-	+	gal ₁ -	↓	
7	0	+		↓	
8	gal +	↓		↓	+
9	0	+		↓	0
10	0	+		↓	
11	0	+		↓	
12	0	+		↓	
13	0	+		↓	
14	0	+		↓	
15	gal +	↓		↓	+
16	0	+		↓	0
17	0	+		↓	
18	0	+		↓	
19	0	+		↓	
20	0	+		↓	
21	0	+		↓	
22	gal +	-		↓	

lysate made and behaves as HFT in 2175, 518 - (none as in 2175)
 6/20/53 Second lysate made also HFT gal₁ - Strain made - cultured in shake bottle as W2346

W2346

18 gal₁
1 gal₂

~~W2346~~

W2346 = 4 }
7 }
9 }
11 }
6/20/54 failed to survive stroke

10/10/53
Linearity -

750 - K-12d 5/30 pup

1.	no add	3	0	
2.	0.02	24	21	105
3.	0.04	48	45	113
4.	0.06	60	57	
5.	0.08	68	65	
6.	0.1	87	84	
7.	0.12	123	120	
8.	0.14	152	148	
9.	0.16	117	114	
				507
				101

low-remain probin (0.2 - 0.12) of pupae

2175

1.	no add	21	0	
2.	0.02	32	11	55
3.	0.04	64	40	108
4.	0.06	74	53	89
5.	0.08	62		
6.	0.1	118	97	
7.	0.12	101	97	
8.	0.14	101	81	
9.	0.16	159	138	
				348
				87

low-remain probin of 0.2 no pupae

811

1.	no add	37	0	Converted to 0.12
2.	0.02	52	15	x5 = 75
3.	0.04	79	42	x2.5 = 104
4.	0.06	81	44	x1.6 = 73
5.	0.08	135	98	x1.2 = 122
6.	0.1	157	120	120
7.	0.12	127	97	80
8.	0.14	189	152	108
9.	0.16	161	144	182
				97.4

W945 (relative of W902) Test to see if L_p^R by attempt. direction
 one plate
 1. no add $\frac{1}{2} = 12$
 2. K-12d 5/30 $= 2$ } looks like inhibitory effect

W1436 ✓ in nutrition

D(0)	no growth	
D(0)+TB	no growth	
D(0)+LB	slight growth	+
D(0)+TLB	growth	++
D(0)+TLB	growth	+++

10/13/53

Comparison of 2238 and 2297

	Control half	0.1m 1/2 half
1. 2238	62	53
2. 2297	7	2

2281 dishes - Repeated as of course segregants
control bottles are all barren

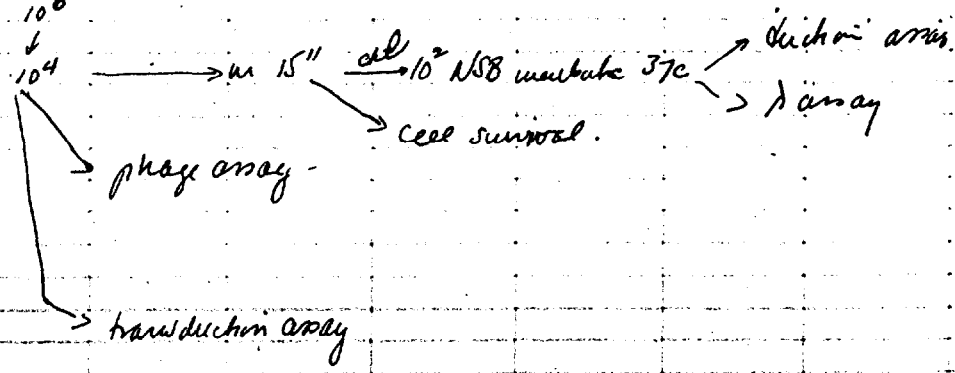
1. 7501 =	214	-	21/24
2. K-121	46	-	18/24
3. P11 λ =	98	-	24/24

N16, NA4 Yield of HFT1 from

Procedure - Serial cultures - centrifuged and resusp. in 1.1 ml saline
c. 10¹⁰ cells/ml

Dilutions

10⁸
↓
10⁶
↓
10⁴



N16

Pre Inoculation - No's given for mod. dose. Post - Inoculation

	<u>N16</u>	<u>NA4</u>	<u>N16</u>	<u>NA4</u>
Plaque Assay				
Cell Assay	1.34 x 10 ³	2.9 x 10 ³	3.37 x 10 ²	3.87 x 10 ²
T-dishes Assay				

deduct work

see 1 232 for notes
24 plates $\frac{1}{2}$ + 4

10/21/53

2281 + K12 Batch II Repeat exam. because of $L_{p.1}$ seq.

Seq	Nb	SIF	$L_{p.1}$	Amala
1.	0	+	S	2-
2.	0	+	S	.
3.	0	+	S	.
4.	0	+	S	.
5.	0	+	S	.
6.	0	+	S	.
7.	0	+	S	.
8.	+	+	T	?
9.	0	+	T	2-
10.	0	+	T	.
11.	0	+	T	.
12.	0	+	T	.
13.	0	+	T	.
14.	0	+	T	.
15.	0	+	S	.
16.	0	+	S	.
			9 S 7 T	16 2- <u>17</u>

re-examine 2- in re-examination after purification.

56% Sp^s

For crosses x 90r

233-3
-7
-13
-16

① 8 plates per x

> 4665
> 4992
106
> 455v

Control str. do except 90r line c. 10 (+)
at 90r for control?
due to lack of neutralization of is cult.?
oh

② cross > 3140 (+)

③ 276 $\frac{9 \text{ plates}}{40r}$ all for control

(-) Counts made on smallest plate, therefore (-) #'s are minima. Variation among plates $\pm 10\%$

233-B a Freckel? cross E 902 F+

summary in sheet

10/21/53

2

2281 t 750 Batch II Repeat for tp characterization

Seq	N16	N1	tp	active
1	0	+	r	2-
2	0	+	r	
3	0	+	r	
4	0	+	r	
5	0	+	r	
6	0	+	s	
7	0	+	r	
8	0	+	r	
9	+	+	r	?
10	+	0	r	1-
11	0	+	r	2-
12	0	+	r	
13	0	+	r	
14	0	+	r	
15	0	+	r	
16	0	+	r	
17	0	+	r	
18	0	+	r	
19	0	+	r	
20	0	+	r	
21	0	+	r	

reexamined - 2 reexam. after purification

Official Exp.

2281 t 750 p 236 B

16 r 5 s
23% s
20 2-
1 1-
1

~~2281 t 750~~
6/1
4.6
1.4
0

10/21/53

2281 + 811 Batch II - Repeat for k_p ✓

Seq	nil	sif	k_p	asile
1	0	+	r s	2 ⁻
2	0	+	r s	"
3	0	+	r s	"
4	0	+	r s	2 ⁻
5	0	+	r s	"
6	0	+	r s	"
7	0	+	r s	"
8	0	+	r s	"
9	0	+	r s	"
10	0	+	r s	"
11	0	+	r s	"
12	0	+	r s	"
13	0	+	r s	"
14	0	+	r s	2 ⁻
15	0	+	r s	"
16	0	+	r s	"
17	0	+	r s	"
18	0	+	r s	"
19	0	+	r s	"
20	0	+	r s	"
21	0	+	r s	"
22	0	+	r s	"
23	0	+	r s	2 ⁻

→

→

discard

2⁵ reexamination after purification

45
17r
19% 5
21
~~20~~ 2⁻
12

10/27/53

Budd III

2281E12 Repeat - as in 226B

Papillae	2281	seg shield out	Segment	N16	N1	Notes
1	lys	mixed	(mixed +)	-	-	mixed
2	lys	..	mixed +	-	-	mixed +
3	lys	..	(mixed +)	-	-	mixed +
4	lys	..	(+)	-	-	lys
5	1/2 non lys	-	-	(+)
6	lys	-	+	lys +
7	lys	-	+	"
8	1/2 non lys	a	..	-	+	lys +
9	lys	mixed	..	-	+	lys +
10	lys	-	+	lys +
11	lys	-	+	lys +
12	lys	-	+	lys +
13	lys	..	mixed (+)	-	+	lys +
14	lys	-	+	lys +
15	1/2 non lys	a	pure seg +	-	+	lys +
16	lys	mixed	..	-	+	lys +
17	lys	-	+	lys +
18	lys	-	+	lys +
19	lys	-	+	lys +
20	lys	-	+	lys +
21	lys	-	+	lys +
22	1/2 non lys	..	non seg	-	+	lys +
23	lys	-	+	lys +

discarded by accident

official

2281E12 Ept. 233

11 Lpt +
2 Lpt

24 1/3

11 Lpt +
2 Lpt

official
2 got

10/26/53

Batch III - 100, 2(+) picked, 3(+) tested / 1485, shaded out in job and a (-) picked and test

2281 ± 750

Popillae	1485	opp streaked w/ mixed	segregant	new segregant	N1	N16
1	lys	mixed	lys		+	0
2	"	"	"		"	"
3	"	"	"		"	"
4	"	"	"		"	"
5	"	"	"		"	"
6	"	"	"		"	"
7	"	"	"		"	"
8	↓ un lys.	"	un lys	segregant	"	"
9	lys.	"	lys	lys	"	"
10	"	"	"	"	"	"
11	"	"	"	"	"	"
12	"	"	"	"	"	"
13	"	"	"	"	"	"
14	↓ un lys	"	un lys	---	lost	→
15	lys	"	lys	"	+	0
16	"	"	"	"	"	"
17	"	"	"	"	"	"
18	↓ un lys	part (b)	un lys	---	lost	→
19	↓ un lys	(+) (-) internal	un lys	---	lost	→
20	lys	mixed	lys	"	+	0
21	"	"	"	"	"	"
22	"	"	"	"	"	"
23	---	---	Dark un lys	---	+	0
24	"	"	lys	---	"	"

20
19 Lp
1 Lp

3/2 un lys.

$MH = gl_2 = 20$

Crossing test x 900 I (-) part II
4 906 131 = 1037
8 > 10 572
12 unmutated Est
16 " " 110
21 110
6 > 100 x 8 = 2800

6/20/54 on strike stand

out 9/13/53

10/26/53

2281 ± 811 ← A 2281 ± 750 on

236B

Population	1485 lys	PT out mixed	Segregants lys	new- Segregants lys	116 0	517 +	Mile 2-
1							
2							
3							
4							
5							
6							
7							
8							
9							
10							
11							
12							
13		slow, (-) ?			0	0	?
14							
15					0	+	2- → on 1 in lys
16		mixed					
17							
18							
19							
20							
21							
22							
23					+	0	4-
24							

1/3 non lys

Weighty ole? No

Disregard any disordered lys classification. apt

For cross x 9 or 1485

$$\begin{array}{r} \times 902 \\ 6 \quad 0/117 \end{array}$$

$$10 \times 13718 = 21096$$

$$14(15000) \times 10 = 210000 \text{ (nearly 100% growth)}$$

$$23 \quad 300/39144$$

6/10/54

Stocks of these made and saved remainder discarded on 6/22/54

out 7/13

" "

2/9/54	811	2175	cross
lysats	1/1	1/1	4-
23	1/1	1/1	-- ?
14	1/1	1/1	

W2350

10/27/53 Examination of the Segregants from HFT medium

In each case pop., and 3 (+) colonies picked in purification

1. 700 has been by N16 g_{002}

(a) Examination of segregants after purification

1.	-	+	1-
2.	-	+	1-
3.	-	+	1-
4.	mixed		
5.	-	+	1-
6.	-	+	1-
7.	+	-	2-
8.	-	+	1-
9.	mixed		
10.	-	-	? double?
11.	+	-	2-
12.	-	+	1-
13.	-	+	1-
14.	-	+	1-
15.	-	+	1-

15 segregants = ~~100%~~ $\frac{9}{18} = 50\%$

part E N1, N16, possibly by S18 - a double? ²
 lysate of this in $\frac{2175}{700} = \frac{3}{1}$ (exp/cult)
 This culture also found to be h_{p1}
 Cross X 902 - controls ok.
 1 plate = 207 x 8 = 1656 (+) (w/ +)
 This is W2586

2. 750E S18 g_{004}

(a) after purification 9/18 = 50% segregants

(b) observations of segregants

1.	+	+
2.	+	+
3.	mixed = (b)	
4.	+	+
5.	-	+
6.	-	-
7.	-	+
8.	-	+
9.	-	+

10/27/53 HFT duhan

(3) 2175 transduced by NI

(a) after purification

	NI	NI6	locus
1.	-	-	?
2.	-	+	1-
3.	-	+	1-
4.	-	+	1-
5.	-	+	1-
6.	-	+	1-
7.	+	-	2-
8.	-	-	?
9.	-	+	1-
10.	+	-	2-
			6 1-
			3 2-
			1 4?

steel segregating = all 16 all int 4

~~no steel segregating~~ on subsequent test int + NI6, E by NI, SIF = gal,

(7) in redheads - transduced by SIF - a double

(4) 2175 transduced by SIF

(a) after purification

	NI6	all int SIF	locus
1.	-	mixed	
2.	+	-	4-
3.	+	-	4-
4.	-	+	2-
5.	-	+	2-
6.	-	+	2-
7.	+	-	4-
8.	+	-	4-
9.	-	+	2-
10.	-	+	2-
11.	-	+	2-
12.	-	+	2-
13.	-	+	2-
			8 (8)
			4 (4)

steel segregating

10/27/53 HFT kawduhais

③ 811 transduced by N16 gal⁻
 ② after purification 15/18 still segregating
 ④ Etadumaki

	N16	S1F
1.	+	-
2.	+	-
3.	+	-
4.	+	-
5.	+	-
6.	+	-
7.	+	-
8.	+	-
9.	+	-
10.	+	-
11.	+	-
12.	+	-
13.	+	-
14.	+	-
15.	+	-

10/29/53

1765 Grd- mutants - Induced in O gal E p.u. c. 10 sec. approx

Inoc #	SIF	516	511	516B2	Control	Arche?
1	mut (+)					
2	0	0	0	0	Lp ^r or clw?	
3						
4	mut (+)	+	+	+		
5		+	+	+		
6		+	+	+	growth for time	
7		+	+	+	Lp ^r	
8		+	+	+	Lp ^r	
9		-	-	-	Lp ^r	
10		+	+	+	Lp ^r	
11		+	+	+	Lp ^r	
12		+	+	+	Lp ^r	
13		+	+	+	Lp ^r	
2		-	-	-	Lp ^r or clw?	
4	mut (+)					
5		+	+	+		new
6		+	+	+	growth for time	col ₁ -
7	mut (+)	-	-	-	Lp ^r or clw	mutated?
8		+	+	+		new
9	mut (+)	-	-	-	Lp ^r or clw	mutated?
10		+	+	+		new
11		-	+	-	Lp ^r	col ₄ x 1
12		+	-	+	Lp ^r	col ₂ -
13		+	-	+	Lp ^r	col ₂ -
		4-	2-	1-		

Stocks

- 2645

- 2646
- 2647
- ~~2648~~
- 2648
- ~~2649~~
- 2650
- 2651
- 2652

embryo - 7
- mut. plus - 9

Runs

#	X	518	-	1255	2405	3639	4810	6050 (-)	no (+)
	X	750	-	581	722	1176	1771	2683 (-)	no (+)

7/14/54

{ Cultures of above entered }
in stock

11/2/53

Lytic λ Recombination

λ 1475 - lytic lambda reported previously. self-d genome on
 (became untransmitted, recombined - EML says
 transduces diploid and 247
 control half 0.1 ml half

W750
 W2175
 W518

2	0
6	2
13	8

Again says no transducing activity of
 lytic λ.

HFT, λ - Does it grow lytically?

1. Fresh 518 cells from aerated culture sedimented - e.g. 10" and
 exposed to 2. 1.0 ml portions of 518 - in adsorption technique -
 sedimented and resuspended in NSP - incubated 3 hours -
 centrifuged (with much cleaning) and chloroformed.

2. Tested against

1. W750
 2. 2175
 3. W518

control half of plate	0.1 ml half of plate
1	90
10	130
12	7

no plaques when
 263 small plaques

Results suggest no reproduction of HFT λ on W518
 Apparently transducing of 700, 2175 identical non-adsorbable
 plaques

Lytic λ - Wild type genome on W518

assayed previously on 2175 with no action
 Reason

	control 1/2	0.1 ml 1/2
W750	1	15
W518	6	38
W2175	10	13

apparent action on 250 518, not on 2175
 ✓ in stabilites

In each case on the 1st shearing in plates
 were found - considered as corroboratory evidence
 that action was genuine transduction.
 The low number of transductions is compatible
 with the assumption that the t were due to
 desiccated or unadsorbed plaques in the prep.

W1210 λ

	control 1/2	0.1 ml 1/2
1. W750	1	92
2. W518	4	128
3. W2175	18	10

20/24 (83%) ~~plaque~~ after 3. (1) picked

Effect of h_p^n on

	sol ₁ control 1/2	transductions 0.1 ml 1/2
1. 2238	9	16
2. 2297	3	45

11/6/53

- Reversals -

750 gal + R - 1

Actin of lysate control = 0, 0.1 ml = 648

2175 gal + R - 1, 2

Actin of lysate 1. control = 10, 0.1 ml = 96
2. control = 6, 0.1 ml = 558

54 gal + R on 750 - A means of distinguishing gal₁ from gal₂

	control	0.1 ml lysate
1. gal + R - 1	1	5
2. " " - 2	1	3
3. " " - 5	0	146
4. " " - 8	1	153

11/8/53

2251 X 750 - Since the principles (in part D) in which case X 2257 have been made, haven't you checked if 2257 distinct from 750?

- in controls done -

	(-)	(+)
1.	173	0
2.	651	0
3.	989	0
4.	1463	0
5.	1781	0
6.	2092	0
7.	2405	0
8.	2743	0
9.	3197	0
10.	3606	0
	3606	0

$$3.6 \times 10^3 \times \frac{20.27 \times 10^{-3}}{1.00} = 0.027\% (+)$$

$$\frac{72}{250}$$

previous animals of

$$gal_2 \leftrightarrow gal_1 = < 0.13$$

✓ in shaking on EM13 gal

10/8/53

578 correlation of transduction with lysogenicity

N16 used
e. 10^{10} cells used - exposed to about 10^7 N16

1. Cell assay - 10^8 dil = 98, 121, 141, = 360% = 120×10^6 = 1.2×10^{10}

2. N16 assay (on 578). 10^7 dil = 7, 4 = $11/2 = 6 \times 10^7$ Assay probably low -

3. streak plates \rightarrow 2 day (+) colonies
 A picked $\left\{ \begin{array}{l} 1 \text{ possibly (+)} \\ 1 \text{ partially lysed col. neither used} \end{array} \right.$ (neither used) total no. of colonies suggests little if any killing of cells by phage

B 12 picked on 2nd day some population colonies - appear untransduced on 578

C 24 mini colonies picked against 578
 2 day (+) both non lys λ
 12 picked all non lys not done -
 24 tested 1 λ^+ 1 λ^- 23 - 3 of intermediate sensitivity -

11/17/53 On a repeat exp. 1 (+) obtained. Analysis of 4 (+) \rightarrow λ resistant, non lysogenic
 4 (-) exp \rightarrow λ sensitive

750 protophage transduced by 902-1

24 (-) picked after purification - exp irradiated and plated against 275, 710
 2 segregants appear to be gal⁻ HFT = #14, #19
 streaks made

W1765 -

negative transduction to obtain gal⁻ in a line compatible with W750 -

1. Three colonies obtained on second streaking from HFT gal⁻ (NI) and 1765 on B(-)

2. Checked against HFT 1, HFT 2, HFT 4. Not transduced by HFT 1 - therefore suspected gal⁻

3. Crossed with irradiated W750 on B gal - control ok on 15 EMS gal plate a total of 152 (-) colonies - has this culture gone F (1765 gal⁻)
 News - using non irradiated 750

Conflict?

W2373

11/11/53

578 E 12/10 Analysis of the segregant.

Segregant	tp	N16	S18	Allele
1	r	+	0	4
2	r	+	0	4
3	r	+	0	4
4	r	+	0	4
5	s	+	0	4
6	r	+	0	4
7	..	+	0	4
8	..	0	+	2
9	..	+	0	4
10	..	+	0	4
11	..	+	0	4
12	..	+	0	4
13	..	+	0	4
14	..	+	0	4
15	..	+	0	4
16	..	Contam.	0 (+)	-
17	..	+	-	4
18	..	+	-	4
19	..	+	-	4
20	..	-	+	2
				<u>17</u> 4
				2 2

Crosses
 8 x 902, 1436,
 20 x " , 1436,
 5 x 902, 1436
 9 x " "

6/18/54
 Stocker of
 Fun
 Capt. discarded

hysets: 4/10/54 / 811 / 2175 Cross
 - 8 1/1 1/0
 - 20 7/1 1/0 2-

11/12/53

75061210 (1) reg analysis -

Serient #	LP	N1	N16	Made
1	r	0	+	1
2	.	0	+	1
3	.	+	0	2- ✓
4	.	0	+	1-
5	.	0	+	1-
6	.	+	0	2- ✓
7	.	+	0	2- ✓
8	.	0	+	1-
9	.	0	+	1-
10	.	0	+	1-
11	.	0	+	1-
12	.	0	+	1-
13	.	0	+	1-
14	.	0	+	1-
15	.	0	+	1-
16	.	0	+	1-
17	.	0	+	1-
18	.	0	+	1-
19	.	0	+	1-
20	.	0	+	1-
21	.	0	+	1-

18 1-
3 2-

hysatz	2/10/54	2/75	750	hmm	6/20/54
3	19/1	1/0	-	2-	} failed to minimize
6	17/1	0/0	-	2-	
7		2/2	90/1 (20%)	2-	

11/16/50

W2252 = 1895 Hfr hp^s

(-) handpicked with HFT) to obtain Hfr gal-

2341 {

1. Several (-) colonies with N16 HFT gal-
isolated and purified - HFT λ tests indicate gal-
cross with W902

Results indicate HFT cross - almost solid
smear (with dilute cells also)
no (+) is probably 100,000 colonies

(2) Streaked against λ - not seen.
S18 - not lysogenic = L_{p1}^R hp^s
 $\lambda 2$ - sensitivity

(3) "Lysate" prepared - very viscous

0.1 ml / 2281 = no effect, neither phage nor λ
" 518 = " " (?) 14/9 = not λ / control area

11/25/50 Repeat with NA-4 lysates 1+2

W 2345 =

From lysate 1 - one gal- isolated - (tested against N16, NA-1 gal-
From lysate 2 - 5 gal- isolated - " " " " gal-)

NA-1 } lysates
NA-2 }
sketchy
confirmed

also nonlysogenic

11/30/53

Transductions for ^{segregant} analysis

A W1765 gal⁻ (made by transduction) W2373

	λ	cont'd %	phage $\frac{1}{2}$
1.	811	1	30
2.	1210	0	56
3.	902 (by 100)	1	11
4.	K-12	1	33

W 750

1.	K12	1	46
----	-----	---	----

W1210

1.	K-12	4	21 (33)
2.	750	2	65
3.	811	5	56

12/5/53

See also next page
 T18 } is against 2070 - to see if both 1, 2- is in these apparent doubles.
 T19 }
 (-2) lysate not very active

245B

T18 - $\frac{5}{11}$
 T19 - $\frac{7}{2}$

1. Analysis of transductions of 2070 by T19-2

	segregants / N1	/N16	Focus
7.	+	0	2-
8.	+	0	2-
9.	0	0	---
4.	+	0	2-
5.	+	0	2-
6.	+	0	2-
7.	+	0	2-
8.	+	0	2-
9.	+	0	2-
10.	+	0	2-
11.	+	0	2-
12.	+	0	2-
13.	+	0	2-
14.	+	0	2-
15.	+	0	2-
16.	+	0	2-

2. Analysis of segregants from 2 hand. T18/2070

	N16	N1	d %	Attenu
7.	+	0	R	1-
8.	+	0	"	1-
9.	0	+	"	2-
4.	0	+	"	2-
5.	+	0	"	1-
6.	+	0	"	1-
7.	+	0	"	1-
8.	contaminated (+)			

12/2/53

21756750 A-

A: T18⁺ transduced by machine N16, ~~N1~~, but not by single λ 's

Proposed Conclusions: ① 18 pop (gold) picked and streaked. Many unpaired colonies - all appear to be segregating
 $\left. \begin{matrix} \text{loc}^+ & 2^- & 1^- \\ & 2^- & 1^+ \\ & 2^+ & 1^- \end{matrix} \right\} 3_n$ on loc should be loc slow if gal^- present - which is the case
 can't read gal segregating

② One of $+$ a gal streaked out on loc, gal - segregating on gal - also on loc, (+) appears to be segregating loc slow

	on loc	Segregating	by band
loc 2-1-	s	no	wt.
loc 2+1-	s	no	1-
loc 2-1+	+	no	2-
loc 2 1/2 1/2 +	+	seg loc slow	2-, wt.
loc 2 1/2 1/2 -	-	no	1-, wt.

see next page 246A for details

B. 2345 / 2251 λ to make HFI, HFr

0.1 wt 2251 λ = 56, control γ_2 - 0

One second streaking only 4/24 $+$ - greater stability of trans. than expected
 The 4 restreaked and segregants examined

12/19/53 T18 continued.

Transl. Array

1. The loc flow -

#	Flow	Zone	NI	N16	hours
1	slow	(-)	0	+	1-
2	slow	"	0	+	1-
3	slow	"	0	0	durk-
4	slow	"	0	0	"
5	"	"	0	+	1-
6	"	"	0	0	entire-
7	"	"	0	0	"
8	"	"	0	+	1-

- An abundance (12/14) there are populated quite heavily

41-
4 (-)

Low chemical not good indicator

2. The loc +

#	Flow	N16	hours
1	(?)	+	1-
2	0	0	0
3	0	0	0
4	0	+	1-
5	0	0	0
6	0	+	2-
7	0	0	1-
8	0	+	2-
9	0	0	2-
10	0	+	2-
11	0	0	0
12	0	0	0
13	0	0	0
14	0	0	2-
15	0	+	1-
16	0	0	0
17	0	0	0
18	0	+	?
19	0	+	1-
20	0	0	2-
21	0	0	0
22	0	0	0
23	0	+	1-
24	0	0	0

24 loc all picked from here and good tested

Can find heavy flow and locate - suggest this is 2-

Ignore
Transl. Array
according to data
no. probably #14 + #15
On 2 observations the following marked @ were found stable - why stable?

#	N16	hours	Notes
1	+	1-	(+)
2	0	0	
3	0	0	
4	+	1-	
5	0	0	
6	+	2-	
7	0	1-	
8	+	2-	
9	0	2-	
10	+	2-	
11	0	0	
12	0	0	
13	0	0	
14	+	2-	
15	+	1-	Rev. on this
16	0	0	
17	0	0	
18	+	?	
19	+	1-	
20	0	2-	
21	0	0	
22	0	0	
23	+	1-	
24	0	0	

lines remaining in double - ?

save stable only

save stable only

1st 12 seq. (-)

3/5/54 lysate made of 246A-15

1/78	solid spot	indicates HFT gel 2
1/120	0	
1/811	solid spot	

(unit level) what to think about this

Correlation of duck... above with 1st... HFT gel 2... but that this is a... HFT gel 2

13 --
6 1-
5 2-

6/18/54
246A-15 + check discarded
246A-15 on check

12/6/53

See pg 215 -
Sample size = 22
6/12 KAT

S K
4- +
2+ -
↓
S R
4* + → (247A) R
2- - 2- +

W/1210 transduced by K12

Transduct	Transduced / 518	Segregant / N16	NI	Segregant / R _x	Milelet	
1.	lyp	0	+	✓	2-	
2.	...	0	+	↓	2-	
3.	...	0	+	↓	2-	
4.	...	0	+	↓	2-	
5.	...	0	+	↓	2-	
6.	...	0	+	↓	2-	
7.	pure salt	→				
8.	pure salt	→				
9.	...	0	+	✓	2-	
10.	...	0	+	↓	2-	
11.	...	0	+	↓	2-	
12.	...	0	+	↓	2-	
13.	weat	0	+	↓	2-	
14.	...	0	+	↓	2-	
15.	pure salt	→				
16.	pure salt	→				
17.	...	0	+	✓	2-	
18.	weat fine salt	→				
19.	pure salt	→				
20.	salt + salt	→				
21.	pure salt	→				
22.	...	0	+	✓	2-	
23.	...	0	+	↓	2-	

6/11/54 Stock discarded
See ↓

19/53 +

15/15 salt

Cross x 902
247A-4
247A-13
247A-22
247A-33
6/20/54
out
5/13/54
in stocks

lysozym	4/20/54	5/13/54	6/11/54
4	1/20.1	2/0.5	2/75
13	0/20.1	0.70/0.5	
22	0/20.1	0.20/0.5	
23	0/20.1	0.40/0.5	

these two lysates give large numbers of small papillae } possibly not sterile? probably Hfr?

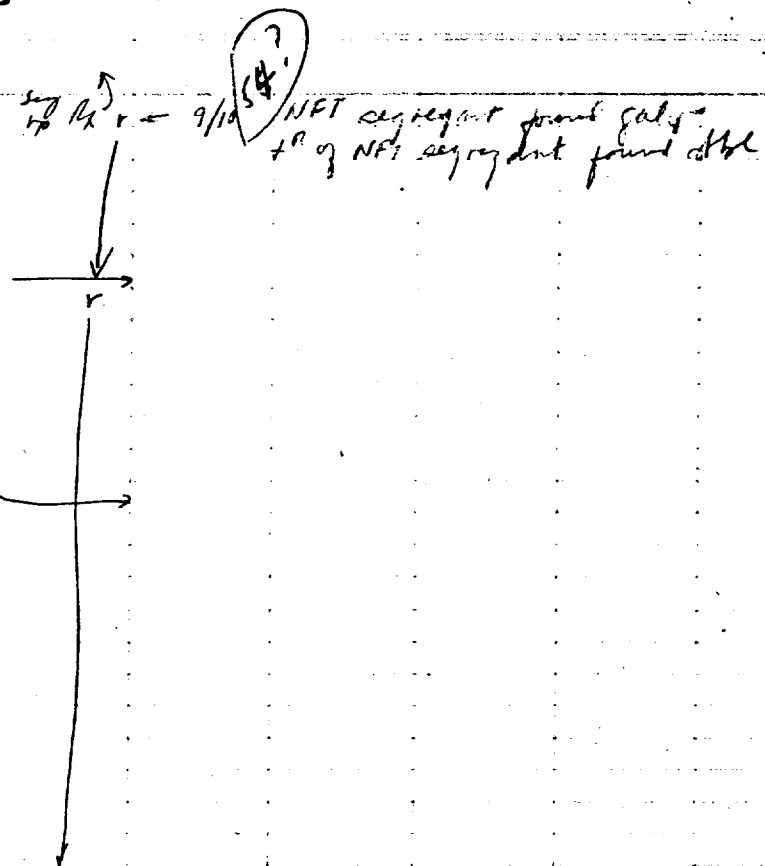
12/6/53

W 1210 t 811

See 245 for origin

245B

Transd.	Transd. / ST	seg ↑	MT +	ST 0
1	lys	0	+	0
2	"	0	+	+
3	"	0	+	+
4	"	0	+	+
5	"	0	+	+
6	"	communicated with gult		+
7	"	0	+	+
8	"	0	+	+
9	"	0	+	+
10	"	0	+	+
11	"	0	+	+
12	"	0	+	+
13	"	communicated with gult		+
14	"	0	+	+
15	"	0	+	+
16	"	0	+	+
17	lys	0	+	+
18	"	0	+	+
19	"	0	+	+
20	"	0	+	+
21	"	0	+	+
22	"	0	+	+
23	"	0	+	+
24	"	0	+	+
all +		22 gult - 1 gult		



Transd x 1436, 90v

- 1
- 8
- 14
- 18

6/18/54
stocks discarded

7/9/54
Lysals

-1 / 0%

solid green = HFT gult

(ALSO as the result F+ x Fx)

12/16/53

W1710 E 750

See 245

(247C)

Transd. #	Transd. Sp Rx / SR	Seq N16 M3 R1A	Seq N1 M3 R1A	Seq Rx Sp Rx	Notes
1	lyu	0	+	v	2'
2	"	0	+	v	2'
3	"	contaminated with gold			2'
4	"	0	+	v	2'
5	"	0	+	v	2'
6	"	0	+	v	2'
7	"	0	+	v	2'
8	"	0	+	v	2'
9	"	0	+	v	2'
10	"	0	+	v	2'
11	"	0	+	v	2'
12	"	0	+	v	2'
13	"	0	+	v	2'
14	"	0	+	v	2'
15	"	0	+	v	2'
16	"	0	+	v	2'
17	"	0	+	v	2'
18	"	0	+	v	2'
19	"	0	+	v	2'
20	"	0	+	v	2'
21	"	+	0	v	1'
22	"	+	0	v	1'
23	"	contaminated with gold			1'
24	"	+	0	v	1'
all +"		19 gold 2-			
		2 gold 1-			

6/24/54
 Phase on
 at
 1
 3
 24
 5/13/55
 " "
 "

12/5/50

4750 transferred by K121

Trans. #	Trans. / 528	N1	N16	Abilities
1	Up.	0	+	1-
2	"	pure galt	→	
3	"	"	→	
4	"	"	→	
5	"	contaminated with galt	→	
6	"	0	+	1-
7	"	0	+	1-
8	"	0	+	1-
9	"	0	+	1-
10	"	0	+	1-
11	"	0	+	1-
12	"	0	+	1-
13	"	0	+	1-
14	"	0	+	1-
15	"	pure galt	→	
16	"	0	+	1-
17	"	0	+	1-
18	"	0	+	1-
19	"	0	+	1-
20	"	contaminated with galt	→	
21	"	pure galt	→	
22	"	0	+	1-
23	"	0	+	1-
24	"	0	+	1-
17/24 +4				16 1-

hyath	2175	750
1	1/0.5	0%
7	2/0.5	0%
9	2/0.5	0%
11	5/0.6	0%

6/20/54
 ← only culture surviving in stock of the first

12/8/53

W2373

249 A

W1765 gal, - (byes transd. with NAY byate) transduced by 841 - See pg 245-

Transd.	From 4 2	Stab exp-2	Seq. exp-2	SIF
1	S X am by	stable	N1	SIF
2	R X by	"		
3	R X by	"		
4	R X by	"		
5	R X by	"		
6	R X by	unstable		
7	S X am by	stable		
8	R X by	"		
9	R X by	"		
10	S X am by	"		
11	R X by	"		
12	R X by	"		
13	R X by	"		
14	R X by	"		
15	R X by	"		
16	R X by	"		
17	R X by	"		
18	R X by	"		
19	stopped	"		
20	R 5 am by	"		
21	R 5 II	"		
22	S X am by	"		
23	S "	unstable		

+ (?) ~~up~~ A² = gal, -

aha?

Lpt = 11
Lpt = 5
1/23+
Lpt

check
shades discarded

5 s
17 +
0 r

0.60
59141.0
354
560

23	22
24	16
12	12
23	13
82	63
23	23
18	72

41
53

12/8/53 W2373

W1765 gal- (by c) transd. NA9) transd. by 1210

Transd #	Transd by	Transd. stability	NI	NI	NI	NI
1	lys	S	-	+	1 ^R	1 ⁻
2	H	S	-	+	1 ^R	1 ⁻
3	H	S	-	+		
4	H	S	-	+		
5	non-lys	S	-	+		
6	lys	S	-	+		
7	non-lys	S	-	+		
8	lys	S	-	+	1 ^R	1 ⁻
9	H	S	-	+	(+)	
10	non-lys	S	+	-	1 ^R	2 ⁻
11	H	S	-	+		
12	lys	S	-	+	1 ^R	1 ⁻
13	lys	S	-	+		
14	non-lys	S	-	+		
15	lys	S	-	+		
16	H	S	-	+		
17	H	S	-	+		
18	H	S	-	+		
19	non-lys	S	x	-	1 ^R	1 ⁻
20	lys	S	-	+	1 ^R	1 ⁻
21	H	S	-	+	1 ^R	1 ⁻
22	H	S	-	+		
23	H	S	-	+	1 ^R	1 ⁻
24	H	S	-	+		

what price?
 encumbrance =

an l_p^R transduction segregating the allelic allele and l_p^R simultaneously.

12/24/53
 Publish an error in recording

S may refer to segregating and not stable.

see next page as well.

$l_p^+ = 18$
 $l_p^R = 6-3 l_p^S$
 $8/24 + "$
 $8 l_p^+$

6 1⁻
 1 2⁻

Line 249B
 10
 13
 20
 21

6/18/54
 stocks discard

12/10/53 12373

1765 gal - transferred by 902

Trand #	Trand. L/lot	Trand. S/lot	Stability	Sag	gent	gas	analysis	S18
1	-L/S	R	S					
2	-L/S	R	S					
3	-nm/L	R	S					
4	-nm/L	S	S					
5	-L/S	R	S					
6	-L/S	R	S					
7	-nm/L	S	S					
8	-nm/L	S	S					
9	-L/S	R	S					
10	-L/S	R	S					
11	-nm/L	R	S					
12	-L/S	R	S					

7 Lp
2 Lp
3 Lp

6/18/54

(Stocks discarded)

There is the only
equipment shown

12/10/53

W2373

Sample 23

2490

W1675 Sol - Transduced by K12

Trans. #	1485	Trans. Sub	W16	W1	W16	W1
1	lys	r	s			
2	lys	r	s			
3	lys	r	s			
4	non-lys	s	u	-	+	GR 1- ✓ lys
5	lys	r	s			
6	non-lys	s	u	-	curtam. E (+)	
7	lys	r	s			
8	u	-	+	GR 1-
9	s			
10	s			
11	s			
12	u	-	+	GR 1-
13	s			
14	u	-	+	GR 1-
15	u	-	+	GR 1-
16	non-lys	s	s	-	+	GR 1-
17	lys	r	s			
18	"	"	s			
19	"	"	s			
20	"	"	u	-	+	GR 1-
21	"	"	u	-	+	GR 1-
22	"	"	u	-	+	GR 1-
23	"	"	s			
W16			10/23 + u			

20 Lp⁺ 8 Lp⁺ 9 Lp⁺
 3 Lp⁺ 2 Lp⁺

lys 4, 8, 12, 14 } none have any activity on 1st pass

6/18/54 } stocks discarded

EXPT. 642

250
A

12/12/53 EML - cross 1210 x 2234 Gal²⁻ x Gal⁴⁻ Lp⁺ Lp^v

1. The small colony "prototypes" (probably require B)

#	M16	S18	Lp Rx	locus
1.	+	0	s weak	4
2.	+	0	s	4
3.	+	0	s	4
4.	+	0	s weak	4
5.	+	0	s weak?	4
6.	0	+	R	2 -
7.	+	0	s	4
8.	+	0	s weak	4
9.	+	0	s	4
10.	0	+	s	2 -
11.	+	0	s weak	4
12.	+	0	s	4
13.	+	0	s	4
14.	+	0	s	4
15.	+	0	s	4
16.	+	0	s weak	4
17.	0	+	s weak	2 -
18.	+	0	s weak	4
19.	+	0	s	4
20.	+	0	s	4
21.	+	0	s	4
22.	+	0 (?)	s	4
23.	+	0	s	4
24.	+	0	s	4
25.	+	0	s	4
26.	0	0 (414)	s	?
27.	0	+	s	2 -
28.	0	0	s	?
29.	+	0	s	4

23 = 4⁻ all Lp^v
 4 = 2⁻ 3 Lp⁺
 2 = 2⁻ 4 Lp^v
 (1 Lp^v)

F⁺ Gal²⁻ Gal⁴⁺ Lp⁺ x F⁻ Gal²⁺ Gal⁴⁻ Lp^s

Cont. of 250A.

2. The Large colony photo people.

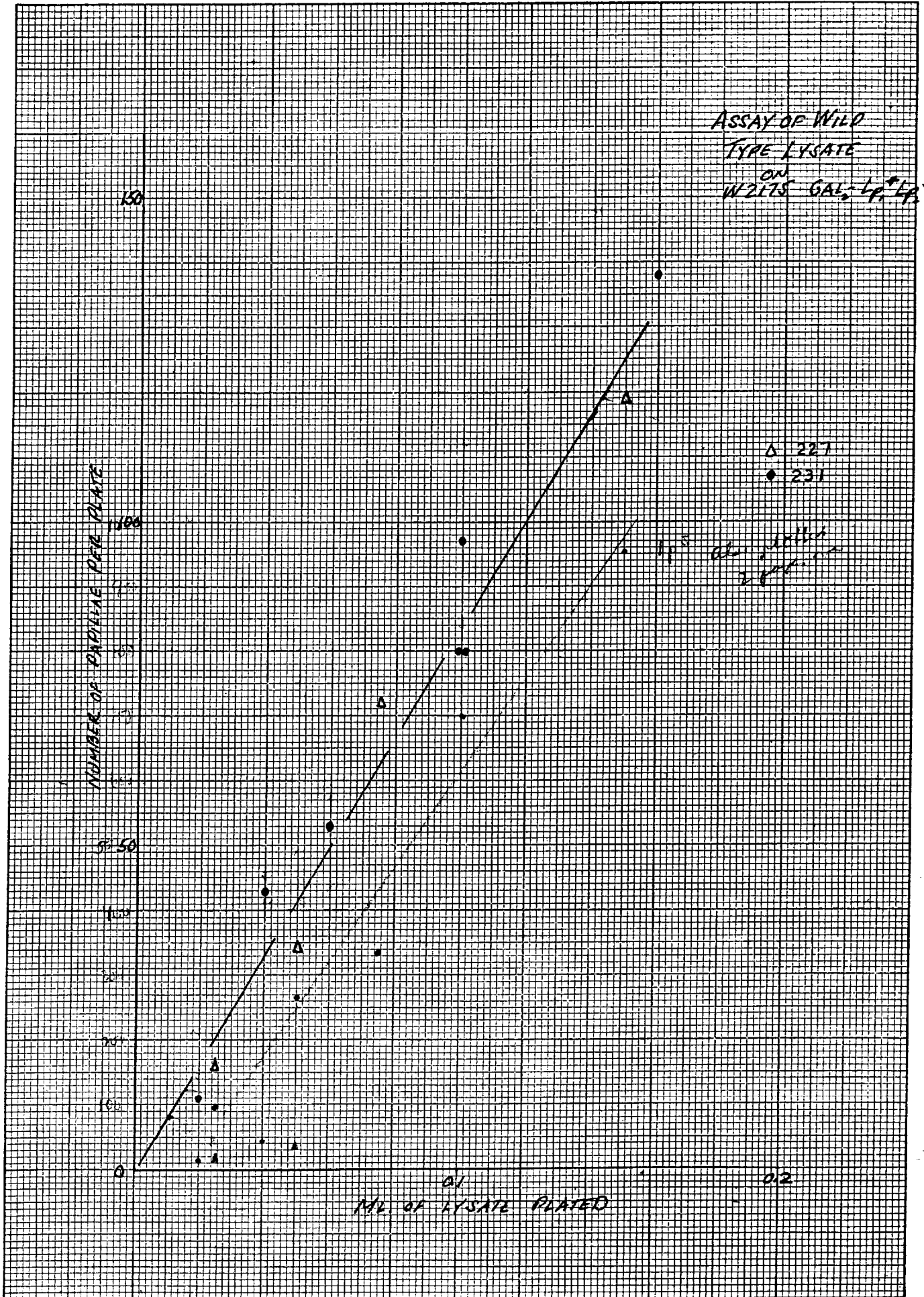
	N16	S18	Lp	Locus
1.	0 ✓	0 ✓	S	---
2.	0	+	r	2-
3.	+	0	S	4
4.	0	+	r	2-
5.	+	0	S	4
6.	+	0	S	+
7.	+	0	S	4
8.	0	+	r	2-
9.	0	+	r	2-
10.	+	0	S	4
11.	+	0	S	4
12.	0	+	r	2-
13.	+	0	S	+
14.	+	0	S	4
15.	+	0	S	+
16.	0	+	r	4-
17.	0	+	r	2-
18.	+	0	S	4
19.	+	0	S	4
20.	+	0	S	4 ✓
21.	0	+	r	2-
22.	0	+	r	2-
23.	+	0	S	4
24.	+	0	S	4
25.	0	+	r	2-
26.	+	0	S	4
27.	+	0	S	4
28.	+	0	S	4
29.	0	+	r	2-
30.	+	0	S	4
31.	+	0	S	4

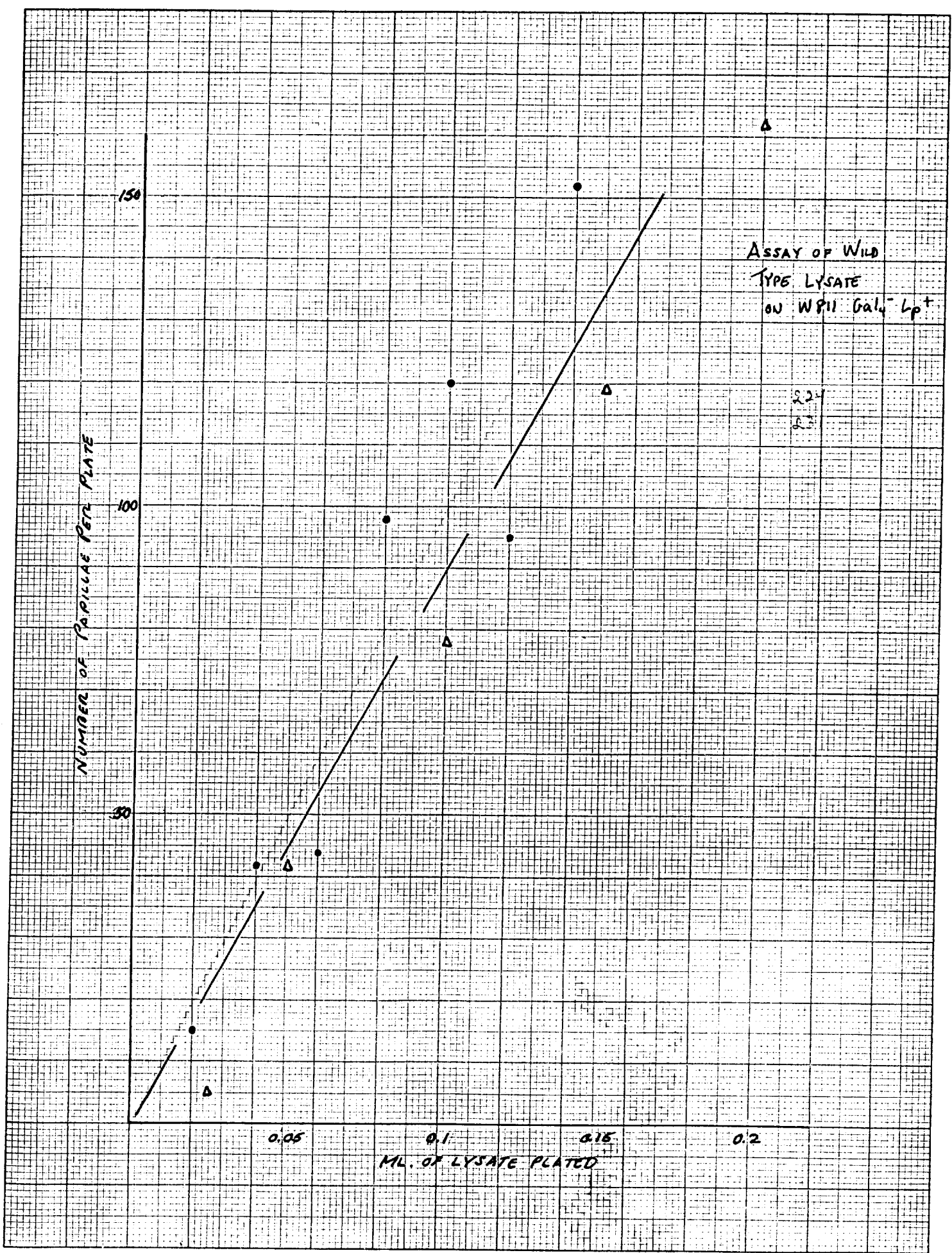
4- Lp^v X 2- Lp^t

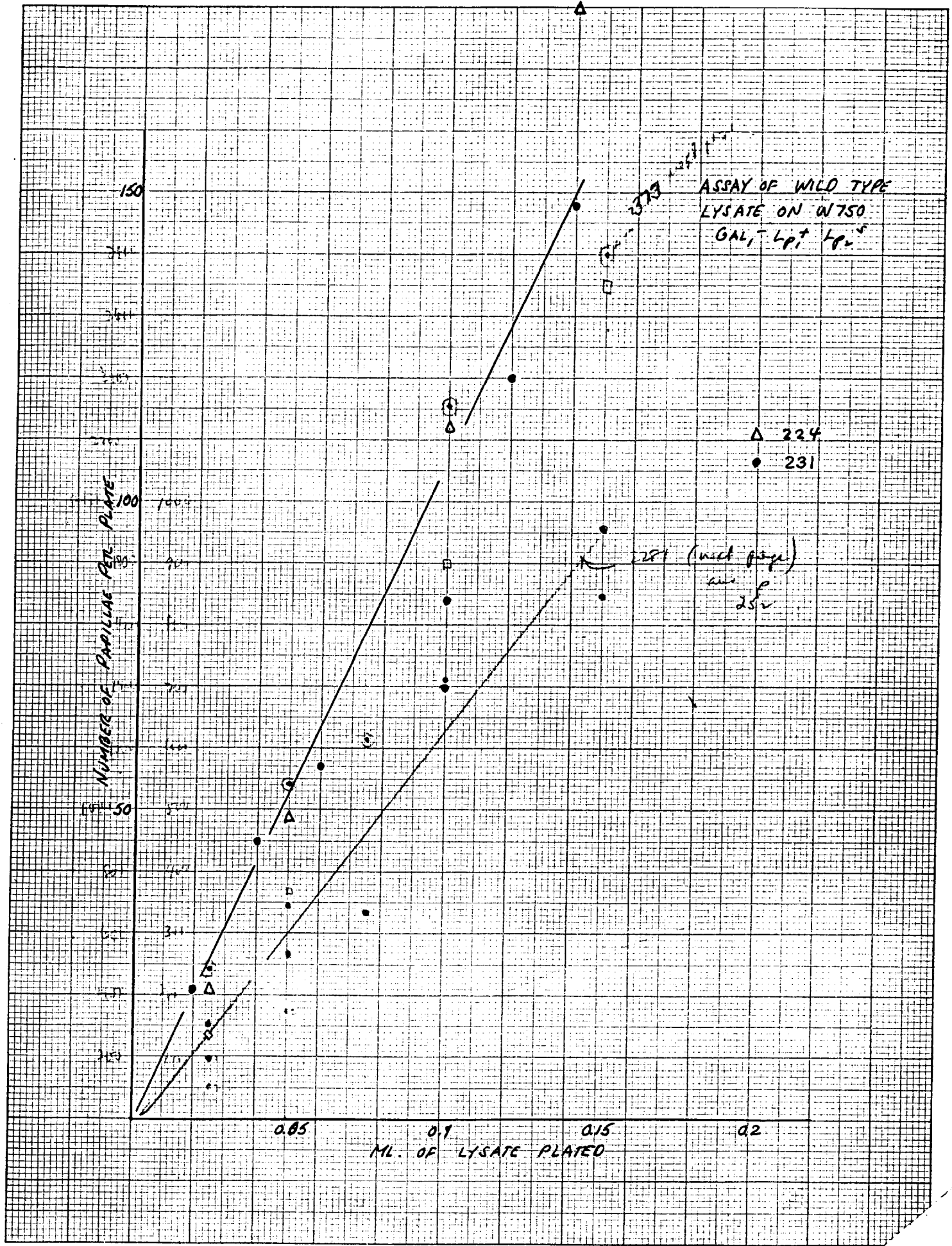
Total of 250A + 250B

43	4-	Lp ^v
13	2-	Lp ^t
1	2-	Lp ^v
3	2-4-	Lp ^s

20 = 4- all Lp^s
 10 = 2- all Lp^t
 1 = 2-4- Lp^s







Assay

K121 5/20/53

$3.6 \times 10^{10} \lambda/ml$

LpK

ML	227	231
0.025	16	0.02 11
0.05	34	0.04 43
0.075	72	0.06 53
0.100	—	0.1 97
0.15	119	0.16 138
0.20	—	—
0.3	255	—

224	231
0.025 - 21	0.02 - 21
0.05 - 49	0.04 - 45
0.1 - 112	0.06 - 57
0.15 - 183	0.08 - —
—	0.10 - 84
—	0.12 - 120
—	0.14 - 148

200C

224	231
0.025 5	0.02 - 15
0.05 42	0.04 - 42
0.1 77	0.06 - 40
0.15 119	0.08 - 51
0.2 162	0.1 - 120
—	0.12 - 95
—	0.14 - 152

2281
LpS
K121 12/28/53
 $1.7 \times 10^{10} \lambda/ml$

0	0
0.025 - 96	151
0.05 - 266	345
0.075 - 334	—
0.1 - 700	706
0.15 - 955	842

2377
2094
LpS
K121 12/28/53
 $1.7 \times 10^{10} \lambda/ml$

0.025 - 483
0.05 - 1079
0.075 - 1223
0.1 - 2314
0.15 - 2799

578
K121 12/28/53
 $1.7 \times 10^{10} \lambda/ml$

0	0
0.025 276	
0.05 738	
0.1 1778	
0.15 2689	

1924
K121 12/28/53
 $1.7 \times 10^{10} \lambda/ml$

0	0
0.025 14853	
0.05 3772	
0.1 143	
0.15 249	

1.7
1.5
8.5
2.5

12/17/53

2345 - Hfr gal⁻ Use as a allele tester by growth with prototrophic gal⁺ - for sexual hous and then plating out on B gal

1. Preliminary encouragement.

Tube	Final test culture	Plated on B gal after 6 hours	no. of (+) spots
1. 2175 (= gal ⁻)	0.1 ul	12	
2. 2345 (= gal ⁻) Hfr	0.1 ul	1	
3. 2175 + 2345	0.1 incub	166	

} suggest method may work

12/20/53

2. Cultures apparently need purification in some places

purify	Parent 2	0	Gal ⁻ Hfr Parent 2341	Gal ⁻ 2345	Gal ⁻ 2341	Gal ⁻ 2345	Gal ⁻ 2341	Gal ⁻ 2345	Gal ⁻ 2341	Gal ⁻ 2345
✓	2341	47	—	—	—	—	—	—	—	—
	2345	3	—	—	—	—	—	—	—	—
	2343	3	(+) > 500 ✓	(-) 3	—	—	—	—	—	—
	753	26	(+) > 1000 ✓	(-) 14	—	—	—	—	—	—
✓	1210	9	51 107? ✓	(+) > 500	—	—	—	—	—	—
	2175	14	(-) 17 ✓	(+) > 1120	—	—	—	—	—	—
✓	2281	1	(-) 41 ✓	(17) 21	—	—	—	—	—	—
	518	11	(+) > 10 ⁴ ✓	(+) > 250 ✓	—	—	—	—	—	—

12/29/53

Are lysates "decaying" at room temperature? Probability to repeat massive deletion 518 and the inability to obtain expected ^{210.5} ~~count~~

no band.
 3 518
 2 1765 gal⁻
 7 K12 518

of transduction of λ p⁵ or compared to λ p⁺ says perhaps this would also argue that transduction of λ p⁺ is affected by particles different from those affecting λ p⁵ deletion

NEW lysates of N16, K12 mode - designated 12/25 approximate titer

1. K12 - $167 \times 10 \times 10^7 = 1.67 \times 10^{10}$ against 2281
2. N16 - $> 50 \times 10 \times 10^7 = > 5 \times 10^9$ " "

N1765 gal⁻ - made by (-) band. cultured into stock both on

W2373

12/31/53

Array against λ^5 - use of K12 λ 12/28/53

1. 2373 - λ added - L_p^5 best conc. - no. cells in suspension

λ added	pop/plate	Δ	
0	3	0	
0.025	486	483	
0.05	1082	1079	
0.075	1226	1223	
0.1	2314	2311	
0.15	2799	2796	

$= \frac{1378 \text{ 1104}}{2} = 1243 \times 10^7 = 1.2 \times 10^{10}$

added / plate = 1.2×10^9

2. 2287 - λ added - L_p^5 - no. cells in suspension = $2145 \times 10^7 = 2.1 \times 10^{10}$

λ added	pop/plate		
0	0		
0.025	96	- 1st of row	
0.05	266		
0.075	377		
0.1	700		
0.15	955		

Suspension appears mixed - large and small colonies

picked and added - tabs against 5th - all reg. λ^+

2.1×10^9 added

11/9/53 Purified 7211 - small colony from cell array

λ added	pop/plate	
0	0	
0.025	151	
0.05	345	
0.1	642	
0.15	706	

$221 \left\{ \begin{array}{l} 211 \\ 200 \end{array} \right\} \times 10^7 = 2.1 \times 10^{10}$

2.1×10^9 added

1/4/54

cells from aerated 10x conc. cultures.

Arrays - K12 A 12/28/53

1. 881

no add	40	Δ	
0.025	94	0	
0.05	138	54	✓
0.1	247	98	
0.15	319	207	
		279	

2. 1974

no add	27	Δ	
0.025	80	0	
0.05	105	53	
0.1	170	72	
0.15	276	143	
		249	

3. 2371

no add	34	Δ	
0.025	217	0	
0.05	423	183	
0.1	945	389	
0.15	1388	911	
		1364	

4. 518

no add	55	Δ	
0.025	331	0	
0.05	793	276	
0.1	1833	738	✓
0.15	2744	1778	
		2689	

1/11/54

518 Transduction of λ , with HFT gal⁻ lysate of 1/9/54

① One ml lysate added to one ml of cell suspension - incubated at 37C for 10 min
 then exposed cell titer =

708
 889
 794
 889
 $3280/4 = 820 \times 10 \times 100 \times 100 \times 50 = 410 \times 10^7 = 4.1 \times 10^9/ml$

1/12/54
 centrifuged and resuspended in
 1.0 ml broth. aliquots and
 plated.

λ exposed cell titer

761
 651
 670
 719
 $2807/4 = 700 \times 10 \times 100 \times 100 \times 50 = 3.5 \times 10^9/ml$

② Distribution of λ types -

	gal(-)	gal(+)	gal - mutation	Total
1. Control (both exposed)	3280	0	0	3280
2. λ exposed	2801	31	54 38	2886

31 (-) picked and streaked / 518 - all um lys.
 / λ - all sensitive

30(+) picked

#	518	λ	Sensitization?	518	λ	Sensitization?	518	λ	Sensitization?	Summary
1. Lpt ⁺	lys	s	yes	um lys	s	no	21. Lpt ⁺	a	"	4
2. Lpt ⁺	"	"	"	12 Lpt ⁺ lys	s	yes	22. "	s	"	23 Lpt ⁺
3. Lpt ⁺	"	"	"	13 Lpt ⁺ "	"	"	23. Lpt ⁺	s	"	3 Lpt ⁺
4. Lpt ⁺	"	"	"	14 Lpt ⁺ "	"	"	24. Lpt ⁺	"	"	"
5. Lpt ⁺	"	"	"	15 Lpt ⁺ "	"	"	25. " Lpt ⁺	"	"	"
6. Lpt ⁺	um. lys.	"	"	16 Lpt ⁺ um. lys.	s	"	26. " Lpt ⁺	"	"	"
7. Lpt ⁺	lys	"	"	17 Lpt ⁺ lys	"	"	27. um. lys	some s. present	"	"
8. um. lys.	s	"	"	18 Lpt ⁺ "	"	"	28. um. lys	"	"	"
9. "	"	"	no int. 81	19 Lpt ⁺ "	"	"	29. Lpt ⁺ Lpt ⁺	"	"	"
10. "	"	"	a (-)	20 Lpt ⁺ "	"	"	30. Lpt ⁺ Lpt ⁺	"	"	"
11. "	"	"	"	"	"	"	31. um. lys	"	"	"

254B Lytic λ - Plaque (#3) of lytic d. prep. previously reported machine -
 um. lys. no control spotted done.

- 1. 750
- 2. 2781
- 3. 2373
- 4. 518
- 5. 811
- 6. 4/4 tested stable - 2 appear slow (+)
- 7. 39 - all stable (!)
- 8. 9 - all discarded as slow and slow growing
- 9. 4/4

Streaks inside

2/4/54
 Stacks of these discarded
 excepting #s 6, 16, 27
 Stacks made of these

254-5 = W2866
 254-16 = W2867

1/13/54

Crosses - check on *Lp gal aggregatus*

A 7. 1765 X 750 on S gal
39 protoplasts - all (-)

B 2. 2281 X 2035
25 (-)
329 (+)

C 3. 902 X 1655
not counted - many small - appear about 50-50

1/22/53

246A-15 Reversions continued

Reversion	Segregating?	Segregating Characters	Proposed Genotype
1	yes	(--)	$\left. \begin{array}{c} \frac{1 \quad 2}{- \quad -} \\ \hline + \quad + \end{array} \right\}$
2	"	(--)	
3	"	(--)	
4	"	(--)	
5	"	(--)	
6	"	(--)	
7	? <u>no</u>	stable +	$\left. \begin{array}{c} \frac{1 \quad 2}{+ \quad -} \\ \hline - \quad + \end{array} \right\}$
8	? <u>yes</u>	mixed +	
9	? <u>no</u>	stable +	
10	yes	(--)	
11	"	2-	
12	? <u>no</u>	stable +	
13	yes <u>no</u>	"	
14	"	(--)	
15	"	(--)	
16	"	(--)	
17	"	(--)	

a 1-6, 10

2/4/54

811E902 (New lysate of 902 - 3 bottles)

for the purpose of isolating HFT gal₂

1. mixed 1/2 = 19
2. 0.1 902λ = 89

2/8/54

750E902 gal₂ (regents sitting around about a month) suspected of HFT gal₂ - Origins now cloudy - Originally 3 but one lost
Both in need of purification

#1 Test for HFT	#2 Test for HFT
257-1 → Col. 1	257-3 → 1
2. NFT	2. "
3. HFT	3. "
4. "	4. NFT
5. "(?)	5. HFT
6. "	6. "
7. "	7. "
257-2 → 8. "	8. NFT
9. NFT	9. HFT
10. HFT	10. NFT

7/10/54

NFT cultures examined
257-1 = gal₂, +R stbc
257-3 = gal₂, +R stbc

W2869

518 transduced with N16λ 1-9-54 for the purpose of establishing that the transduced colonies contain both λ⁺, λ⁺ cells and gal₂ cells - (regulation for both?)

Fresh unselected cell. 518, 0.2 ml cells + 0.4 ml λ incubated at 37° for 10'

Plate	(+)	(-)	(-) mibbled	total
1.	1*	28	2	31
2.	1**	32	0	33
3.	0	39	2	41
4.	0	43	1	44
	2	142	4.5	149/142

* a colony such ● } both streaked - 10 (+) and 10 (-) colonies picked + 1/15/54

1.	λ (+) R	λ (-) R	λ (+) R	λ (-) R	λ (+) R	λ (-) R
2.	λ (+) R	λ (-) R	λ (+) R	λ (-) R	λ (+) R	λ (-) R
3.	λ (+) R	λ (-) R	λ (+) R	λ (-) R	λ (+) R	λ (-) R
4.	λ (+) R	λ (-) R	λ (+) R	λ (-) R	λ (+) R	λ (-) R
5.	λ (+) R	λ (-) R	λ (+) R	λ (-) R	λ (+) R	λ (-) R
6.	λ (+) R	λ (-) R	λ (+) R	λ (-) R	λ (+) R	λ (-) R
7.	λ (+) R	λ (-) R	λ (+) R	λ (-) R	λ (+) R	λ (-) R
8.	λ (+) R	λ (-) R	λ (+) R	λ (-) R	λ (+) R	λ (-) R
9.	λ (+) R	λ (-) R	λ (+) R	λ (-) R	λ (+) R	λ (-) R
10.	λ (+) R	λ (-) R	λ (+) R	λ (-) R	λ (+) R	λ (-) R

see p. 262 for consideration of regent from the

W2868

2/13/54

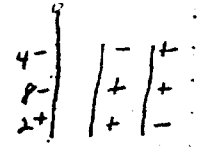
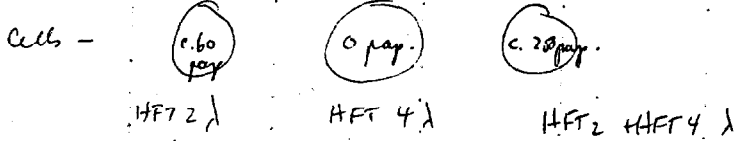
258

W 2350 - double ductin

parent culture streaked out and 10 colonies examined to eliminate possibility of including reversions at one locus in the inoculum - - - - - + and therefore transducible by one of the 2 (-) lysates.

1. All 10 colonies not transducible by HFT 2, 4
2. Lysates of overite both placed on B gal

2350 = 4-8-



Are the pap on HFT 2 due to reversions at one of loci being transduced? If so they should segregate 4- segregants predominantly and by (-) Pick up 12 and see -

258-A

Of the 12 - 6 found unstable Segregant. 1. 0 2. 0 3. 0 4. 0 5. 0 6. 0

HFT 2 HFT 4

0 0 0 0 0 0

+ + + + + +

+ + + + + +

0 0 0 0 0 0

+ + + + + +

0 0 0 0 0 0

+ + + + + +

0 0 0 0 0 0

+ + + + + +

3. Examine HFT 2 + HFT 4 for double ductin

2/18/54 double ductin 258-B pap

	HFT 2	HFT 4	loci			
1.	0	0	(-)			
2.	0	0	(-)			
3.	0	+	2-			
4.	0	0	(-)			
5.	0	+	2-			
6.	0	+	2-			
7.	0	+	2-			
8.	0	0	(-)			
9.	0	+	2-	9	0 (+) pap	
10.	+	0	4-	10	46 (+)	4% stable
11.	+	0	4-	11	40	4% stable
12.	0	0	(-)			
13.	0	+	2-	13	22	4% stable
14.	0	+	2-	14	9 (+) pap	4% stable
15.	0	0	(-)			
16.	+	0	4-	16	11 (+) pap	5% stable
17.	0	0	(-)			
18.	0	+	2-	18	10	10% stable
19.	0	+	2-	19	12	12% stable

control & minute papillae

9 2-
7 (-)
3 4-

recloning 9 (2-4-8-)
7 (2-4-8-)
3 (4-)

3/5/54 Lysate made of #16. Lysate spotted
 750 / 1200 / 800
 0 / 0 / 0

2/13/54

W2350 lysate on 2070 - After about 4 days c. 10 small pop. a
 lysate half - 1.5 larger pop. (reversus?) and on
 control 1/2 about 5 larger pop. - m smaller -

24 picked to gal for purification - all (-)

2/22/54 Rpt HFT duction with N16

N16 lysate 1/9/54 - 0.5 ml
 N16 titer = 2×10^9
 cell titer = $892 \times 10^5 = 8.9 \times 10^7$
 $\frac{2 \times 10^9}{8.9 \times 10^7} = c. 200$ multiplicity 0.5 ml N16 + 0.5 ml cells

EML
 Analysis
 EML 667

control plate c. 900/plate - all (-)

exposed plates: 1/4 of (1) = 213 = 882 per plate - indicates little killing

1/4 of (2), 23, 35, 24 = $\frac{103}{4} = 26$
 $870 \frac{0.03}{2610} = 3.0\% (+)$

St HFT duction with N16 1/9/54

Control culture 1/10 - some sediment - sample in
 0.5 ml N16 added - absorb. c 3 min centrifuge - Rpt 3 times except no
 centrifugation after in 3rd - dil $10^2, 10^2, 25/10 \rightarrow$ plate

See above

Control plates:

37C 371, 341, 319 = $1031/3 = 344 \times 10^6 \times 2 = 688 \times 10^6 = 6.8 \times 10^8$ $\frac{36}{344} = 10.4\%$
 30C 360, 358, 361 = $1109/3 = 369 \times 10^6 \times 2 = 738 \times 10^6 = 7.3 \times 10^8$ $\frac{39}{369} = 10.5\%$

N16 Phage Plates

37C $\frac{63}{763}, \frac{22}{309}, \frac{23}{207} = \frac{108}{1279} = 8.4\% (+)$ 13 = 426
 30C $\frac{31}{175}, \frac{45}{197}, \frac{41}{194} = \frac{117}{566}$ 12 = 189

Esten's
 Page
 667

N16 Assay -

λ 10^7 dil = 106
 $\frac{130}{236} = 120 \times 10^7 = 1.2 \times 10^9$

Taking of (-) in plate - 10⁶ and (+) present
 in any (-) colony?
 30 gal - in "

7/8 found mixed by 10^7 + 10^5 = self plaque

found.

10^7 dil = plaques = 240
 population = 104
 why 2-fold diff? 0.2 ml pipette?

Control plate for sp. done on assumption = 44 ratio $\frac{60}{240} = \frac{1}{4}$

2/18/54

Looking for Recombination between tp^+ by selecting for Φ recombinants when two diff. gal- are tp^R

1. This is second attempt - previously no tp^+ found in (+) recombinants between 2341 gal₁- tp^R x 19204 gal₂- tp^R

2. In this case 2341 gal₂- tp^R x W1 (= 1924 t 902 \rightarrow neg gal₂- tp^R) out of 33 gal(+), in a Bgal mixed culture of these (probably includes sp. reversis) none found tp^+ . Not expected? since in 1st test tp^R x tp^R 1924 failed to give tp^+ .

What goes here? 2341 = $\frac{\text{Gal}_2 - tp^R}{\text{Gal}_2 - tp^+}$
Where did Gal+ come from?

518 t N16 - again for higher fraction (+)

0.4 ml N16 1/9/54 + 0.1 ml of 518 culture c. 10^8 cells/ml

cell array of above	$10^2, 10^4, 10^5$	3 plates	1 = 600 W (+)
duplicate plate	" " $\frac{1}{5}$	3 plates	1 = 163 x 5 815 W (+)

Failure why

2/19/53

246A-15 gal⁺ R-1 } ~~Missed by my self to find a study probably~~
 -11 } ~~probably R-1 w~~

246A 75 - labelled 11 reversions examined

Reversion	single seg	HFT	Structure	Label
1	0	0	$\frac{++}{--}$	246A-15 + R-1 A
2	+	0	$\frac{++}{--}$	- a common " " -1 B
3	0	0	$\frac{++}{--}$	
4	0	0	a-1	
5	0	0	a-1	
6	0	0	a-1	

Because gal⁺ chlo. seg. reverts from structure labelled 246A-15 + R-1 to see if conforms to original definition of 246A-15 + R-1

261 C	3/5/54	lysate of 261 C	reversion	single seg	HFT	Structure	Label
			1	0/7	(-)	1 (gal ⁺)	
			2	shade			} according to this it will be 261
			3	7/7	(--)		
			4	7/7	(--)		
	750	solid spot					
	120	" "					
	811	" "					

Lysates of 261A + 261B made

Ratio of (+) to minus colonies in	Explanation of the (-)	
	(+)	(-)
261A	82	44
261B	103	63

Acting lysate	261A	261B
	CHP	CHP
750	HFT	HFT
2175	HFT	HFT
578	HFT	HFT

1976 2042 Acc. by ... 270 ... NFF
 copy of NFF ... 1-2 ... since NFF ...
 ... 1-2+ ... 1-2-

262

2/28/54

257C-2 5FFt N16 → +^a hp^R - Examination of the segregants

Segregant	from this	NFF ²	S18 NFF ⁴	hp ^R		2 + 1
1	nonlys	0	0	S	7	+
2	"	0	+	S	mt	-
3	"	0	+	R	+	-
4	"	+	0	S	+	-
5	"	0	0	S	+	-
6	"	0	+	S	mt	-
7	"	0	+	S	mt	-
8	"	+	0	S	+	-
9	"	0	+	R	V	-
10	"	0	+	R	V	-
11	"	0	0	S	+	-
12	"	0	0	S	+	-
13	"	0	0	R	V	-
14	"	+	0	S	+	-
15	"	0	+	R	V	-
16	"	0	+	R	V	-
17	"	0	+	R	V	-

results suggest
 that the ...
 1-2-
 1-2-
 On this ...
 # 2, 6, 7 in the
 table ...
 1-2-
 In which case
 the ...
 1-2+4-
 1-2+4-

Not certain that
 these segregants came from
 separate ...

The ...
 results ...
 are not (+)

been observed before

2/20/54 ^{qui's} the original (-) that is dup'd for some portion of rain gal region

246A-15 - Steaked out and (-) shows precep and tested against Both H₂O gal₁ - and gal₂ - (-) and gal₂ - found - no gal₁ -

discuss with lab

The gal₁ - By the present test were plated and ⁽⁴⁾revised stained and streaked on B gal

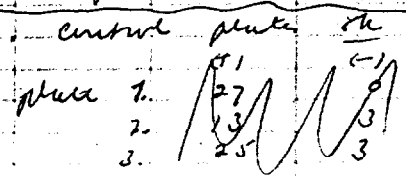
gal ₁ -	Streaks	Result
1	unst.	+
2	"	+
3	stab.	+
4	unstab.	+
5	"	+
6	stab.	+
7	unstab.	+
8	"	+
9	stab.	+
10	"	+
11	unstab.	+
12	stab	+

6/18/53 These stocks discarded -

(5 stable 7 unstab.)

246A-15 ^{photograph stupid!} cross with 1503 gal₁ - EMS gal
 246A-15 $\frac{2-}{-}$ → arrested = F -
 1503 gal₁ $\frac{+}{-}$ → F +

10/26/56
~~these are not~~
 where did these come from? we in phototype?



2/22/54

1503 - (-) direction to obtain gal⁻ in Het, stocks?

1. 2 gal ₂ - obtained	one Lp ⁻	W2424
	one Lp ⁺	W2429
2. 1 gal ₁ - obtained	one Lp ⁺	W2430

Also These are }
 2431 = 7478-1
 2432 = 1673 gal₂⁻
 2433 = 718
 2434 = 719

Crosses of gal⁻ - Control stocks in B gal = pure

1. (X) 750 - c. 30 (small) / plate x 11 = 330
checked

2. (X) 1210 - 9, 8, 13, 12, 14, 12, 6, 6, 8 = 100
checked

3. (X) 811 - 3/17, 3/14, 1/15, 1/28, 1/17, 1/15, 1/16, 1/23, 1/29, 1/16 = 190
checked

(+) gradability of 18 checked
to see if diploids - 7 found (+)

5 of which may be diploid.
4 were

Analysis of these diploids by examination of P₂ progeny.

#2 9 (-) progeny	tested against HFT 1 ⁻ , HFT 4 ⁻	- all Lp ⁺ and gal ₄ ⁻
#3 10 (-) progeny	" " " "	- all Lp ⁺ and gal ₄ ⁻
#4 8 (-) progeny	" " " "	- all Lp ⁺ and gal ₄ ⁻
#5 12 (-) progeny	" " " "	- all Lp ⁺ and gal ₄ ⁻



W2360 - a Lp⁺ form of W2350 gal₂⁻ gal₄⁻
 6 colonies taken from old bottle streaked out from each
 all Lp⁺ and not screened by HFT 2 or HFT 4 in
 two days

Stock made.

3/4/54

Examination of Transducer Complex (+^u) for HFT nature of at all
 Stocks have been maintained in slink for considerable time (6 mo?)

Designation	Temp spotted			characteristic of pp. used in made by spec		
	750	710 advent	811	(-)	(+) mean	
Stock Slant	solid spnt	solid spnt	solid spnt	19	53	5
518EK12	solid spnt	solid spnt	solid spnt	13	79	8
750EK12-1	solid spnt	solid spnt	solid spnt	26	110	8
750E902-1	o	o	o	14	c.50	4
250E912-2	o	o	o	4	c.50	pur plate 1-2
217SE700-1	+++	+++	+++	+	2K1000	?
217SE80-1	+++	+	+++	18	83	?
518E700-1	+++	+	+++	14	32	5
750EK12-2	+++	+ ck	+++	27	62	6
750E902-2	+++		+++			
2						

not sterile
 not sterile
 both cancelled
 217SE800-1

HFT Stocks

<u>Designation</u>	<u>Page</u>	<u>Genotype</u>	<u>Comment</u>	<u>Autotrophic Markers</u>
2342 (N16)	192B	gal ₂ -		none
2346 (N44)	230	gal ₁ -		M-
(D1)	153	gal ₂ -	from 8921 - x 518	M-
(S18)	202	gal ₄ -	probably NFT now	M-
(H14)	242	gal ₂ -		none
(#19)	242	gal ₂ -		none
246A-15	246A	gal ₂ -		none
2000-16H	202	gal ₂	516 has an NFT & it	M-
257-2	257	gal ₂ -	has an NFT & it	M-
257-4	257	gal ₁ -	M-
247B-1	247	gal ₄ -		M-
261 A	261	+ ^u		none
261 B	261	+ ^u		none
261 C	261	+ ^u		none
21752758 + ^u		+ ^u	7501 - x 2175	none
578E1412	263	+ ^u		M-
750E1412-1	263	+ ^u		M-
750E1402-1	263	+ ^u		M-

3/21/53

Tests of some lysates for HFT and allele - See also page 265

Lysate	750	1210	811	Comment	Cell Characteristics Mating w/ lys	Stability of lysate	Yellow canyon #
in also → 2342 (3/14/54)	+++	0	+++	Known gal ⁻ HFT	-	64 1/2 + ⁹ ⊕	1
Pen- 56th. 2342 (3/17/54)	+++	0	+++	Known gal ⁻ HFT	-		2
2175E811-1	++	small	+++		mixed		3
750E612-2	+++	+	+++	belongs to + HFT	mixed		4
two / HFT → 750E902-2	+++	+	+++	" " "	mixed		5
diff. cultures mislabelled here 2478-1 pop 2	1 pop	2 pop	0	gal ⁻ HFT lost?			6
2175E811-1	0	0	2	NFT?	mixed		7
57E750-1	+++	+++	+++	belongs to + HFT	mixed		8
750E902-2	1 pop	0	2	NFT?	mixed		9
258-16	5 pop	1 pop small	0	gal ⁻ unstable transition			10
246A-15	+++	0	+++	gal ⁻ giving + ⁹ reverse			11
261C	+++	++	+++	segregates (---) principally			12
57E812	+++	c. 100 pop	+++	belongs to + HFT	mixed (A) (-)		13
750E812-1	1 pop	+	+++	" " "	" "		14
750E902-1	18 pop	2 pop	c. 30	NFT?	mixed	5 4/5 + ⁴ ⊕	15
234L							

+++ = solid clump
++ = some descent. pop

one of these is 2175E811 other 2175E751

* Gal⁻ + Reverse stability - no obs., no found +⁹ / no. of mutants

4/10/54
all tested streak in Pen. all stable 0.5 ml sample

16 also tested streak
12

3/22/54

Transductions with 2342 (=U16) in NSB

① WSTF	No (+)	No. (-)	NOE (control)	NOE (pap)	cellular
lysate	8	190*	9	1	compd. in 10 ml, 10 ⁷ , 1-50 → 20 ml
Bactn	0	679*	0	2	"

$$\frac{8}{190} = \frac{140}{8.00} \frac{0.057}{700} = 5.7\% (+) \text{ transd. on } l_p^S$$

* difference in numbers probably due in part to loss in centrifuge tube decantation

2. W811

lysate	1	426 *	0	5	above
Bactn	0	448 *	0	1	"

$$1 \frac{0.0022}{1.000} \frac{852}{1480} = 0.23\% (+) \text{ transd.}$$

* two colony types - large and small.

3/25/54 Experiment with TCN - Transmissions of "injectate" in crosses.

(Hfr $l_p^+ l_p^2$ treated) crossed to 2342 \otimes $F^- l_p^- l_p^R$ TCB = Gal₂

Experiment couldn't succeed in first place - gal.

Absorption of 2342 in (2341) *shyformycin* treated

1 ml phage, 1 ml cell suspension

Before	$10^7 = 273, 214 = 2.44 \times 10^9$	$\frac{1}{2} = 1.22 \times 10^9$	No adsorption
After	$5 \times 10^7 = 20, 26 = 1.15 \times 10^9$	(surface plating on B(1))	

$\frac{1.2}{1.2} = 0.0$

on 2252

Before - above] $\frac{7}{12} = 12 \frac{0.58}{7.1} \frac{5.0}{100} = 92\% \text{ adsorpt.}$
After	

3/30/54 Repeat examination of adsorption of λ cells.

W 2341 and W 2342 HFT (Assays by pour tubes)

1. Pre adsorpt. titer = $247, 167 \times 10^7 = 2.07 \times 10^9 = 2.1 \times 10^9$ (compare with above)

2. 5^E cells (0.3 ml) + 0.3 ml 2342 (NSB prep 1/9/54) - micrb. 37 C 10' $\frac{9.4 \text{ ml } l_p^+ \text{ added}}{\times \text{ centrifuging } 5 \text{ min}}$
 dil $10^2, 10^4 \rightarrow 0.1 = 444, 4 P_2$
 titer = $463 \times 10 \times 33 \times 10^4 = 3.3 \times 10^6 = 1.53 \times 10^9$ 20% adsorpt. $\frac{65}{21} = 71\%$ 29% adsorpt.

3/28/54

SIFC 750-1 (Look up origin) Nature of segregant.

Seq.	/HFT-1	/HFT-4	Locus
1	o	+	1-
2	o	+	"
3	o	+	"
4	o	+	"
5	o	+	"
6	o	+	"
7	o	+	"
8	o	-	"
9	o	+	"
10	o	+	"
11	o	+	"
12	o	+	"
13	o	+	"
14	o	+	"

what happened to 4-?

all 1st

lysis made of here to end of HFT-Hfr & has any additional properties

2341 *hydrodromus* (for T₄) using his culture which was found $\frac{1}{5}$

7. K12 lambda

1. small = 4
2. 0.1 K12 (90%) = $3 \times 58 = 174 = 1856$

Use in crosses of added segment as a test of elim. and package of segment in $\frac{1}{5}$ crosses.

Just done beyond

2341 (abm) 2. 902 lambda - (indate HFT Hfr)

1. small = 2
2. 0.1 902 = 42

4/7/54

In Relation to SIFC 750-1 Above. Transduction of 81. with HFT 1- 2346
 Examination of stability - Do the $\frac{1-}{1-}$ combinations +? Does 1- and 4- segregate
 from such a complex? Are $\frac{1-}{1-}$ observed (+) in 4, 1- transductions
 (+) after elim. of (-) allele?

1. 8 peculiar examined 0/8 stable.

	/HFT 1-	/HFT 4-	HFT/1-	HFT/4-
A1.	-	+	3.	+
2.	-	+	2.	+
3.	-	+	3.	+

269-3

3/28/54

Examination of HFT stocks

1. Previously noted (in last week) that cultures of N16 (= 2342) and NA-4 (= 2346) were segregating NFT lines (opt. plate test) (mixed. on plate possible - loopful added to indicator, plate unadmitted). On each case of 10 colonies 8 were HFT - From each a NFT and HFT selected and placed on slants and stored in refrigerator to prevent segregation. A portion of each plated on Royal and reversed in ethanol - (See page 267)

2. 241-14 (a gas₂ - HFT) tested against 1210, 750 - 10 colonies

against 750	^{Colonies} ⊗ ⊗ ⊗ ⊗ ⊗ ⊗ ⊗ ⊗	8/10 HFT. Stocks made of an HFT, NFT.
against 1210	⊙ ⊙ ⊙ ⊙ ⊙ ⊙ ⊙ ⊙	

3. 241-19 (a gas₂ - HFT) as 2 above

against 750	⊙ ⊙ ⊙ ⊙ ⊙ ⊙ ⊙ ⊙
against 1210	⊙ ⊙ ⊙ ⊙ ⊙ ⊙ ⊙ ⊙

Examination of NFT HFT stocks above - also 2342 2346 - See page 267 also

Culture	As Reverse	Stability	Transmission anal. of				Killed
			1/4 HFT	1/1 HFT	1/2 HFT	4 HFT	
241-14 HFT	19 (brown)	12/12 + ⁿ	(+)	+	0	(+)	2-
" NFT	12	12/12 stable	+	+	0	+	2-
241-19 HFT	18	12/12 + ⁿ	(+)	+	0	(-)	2-
" NFT	0	—	+	0	0	13/13 +	1-2
2342 HFT	62	12/12 + ⁿ	+	0	0	+	1-2
2342 NFT	0	—	+	0	0	+	1-2
2346 HFT	15	4/5 + ⁿ	+	0	0	+	1-2
2346 NFT	8	0/8 stable	+	0	+	+	1-

4/5/54

Examination of HFT by state

1. 7505K12-2 - Assay on different indicators - 0.1 ml of overnight unassociated cult. used.

dil	0.1 ml No. pop.	Assay cult	dil	0.1 ml No. pop.	Assay cult.	dil.	0.1 ml No. pop.	Assay cult
na	0	750	na	0	2438(1-2)	-	3	12/10
10 ²	+++	"	10 ²	++	"	10 ¹	+	"
10 ⁴	565	"	10 ⁴	2/4	"	10 ⁴	23	"
10 ⁶	12	"	10 ⁶	4	"	10 ⁶	1	"
<u>Titer</u>		1.2 x 10 ⁸					2 x 10 ⁶	

2. 7505902-2 4/8/54 (lysate that worked)
 Filtered on U2 (sp. 1st) 10⁷ = 717 = 7.2 x 10⁹ - lysate tested stock. p. 67

Assay No. cells	750	12/10	2433	Crustidulin presumed
0.1 ml of dil	0	3	0	1-2+
10 ²	+++	1200-1600	183	2+ 1+
10 ⁴	++	13	1	
10 ⁶	12	?	0	
<u>Titer</u>	1.2 x 10 ⁸	1.0 x 10 ⁶	1.83 x 10 ⁵	

suggests

4/10/54

U2 - sp. 10⁵? from 21755811 -> (-) seq 10⁵ sp. 4 - Only
 useful marker for elim. possibility of contamination is
 p. 67. This culture grows on EMS gel. Presumably
 this is a valid occurrence of 10⁵ - No other gels - 10⁵
 p. 67. about at time of isolation.

4/5/54

In search of triple (-) 1-2-4-

7. W2350 ^{separated} ~~isolated~~ ^{to} HFT 1- (2346) - Segregants examined

Seg #	HFT 1	HFT 2	HFT 4
1	+	0	+
2	+	0 ↓	+
3	0	+	+(weak)
4	0	+	+(weak)
5	+	0	+
6	+	0	+
7	+	0	+
8	+	0	+
9	+	0	+
10	+	0 ↓	+

2-
1+
4-
X-
-
+

8 2-
+ 1-

In these cases
a putative population used
similar to the 1% of 2350 against
HFT 2 - Significance?

8. T18 = W2432 ^{separated} ~~isolated~~ ^{to} HFT 4- (2470-1) - Segregants exam.

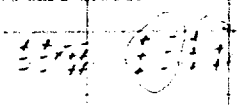
Seg #	HFT 1	HFT 2	HFT 4
1	0	0	+
2	0	0	+
3	0	0	+
4	0	0	+
5	0	0	+
6	0	0	+
7	0	0	+
8	0	0	+
9	0	0	+
10	0	0	+

272-2

→ minute population in non-uv in control

} all 1-2-

2-
1-
4+



4/7/54 - and several days earlier

Recovery of Lp_2 from Lp_1 cultures

W902

1. Inoculated = 2346 HFT 1 - Δ - Reaction suggest family large
2. Reaction of Lp_2 present.
3. Checked for Lp_2 - seg + observed.
3. Just for seg anal (c.b.) all pl₁
4. One tested for nutrition -

D(0)	no growth
+ T2	growth 2 days
+ LB	no growth
+ TB ₁	no growth
+ TB ₂	growth 24 hrs.
5. Examination of imm segs and

} nutrition sh. tested on
 make sh. 13. Mal = met+

W2584

W945

1. Inoculated = 2346 HFT 1 - Δ - React. also suggest family large no Lp_2 present.
2. checked 13 fac - +^a observed.
3. 6 seg tested / 2346 / 2342 - all seg₂
4. Examination of the cultures for sugar reactions.
 Mal + Xyl - Mtl -
 3 additional colonies also found Mal +

W2580

4/9/54

In search of phage mutants.

Platings on E. coli - all platings gave no plaque

With the following cultures.

Assays on NSA (pumps)	10^8 dil	10^5 dil
2281	274	10
1485	27	10
578	281	6

4/10/54

Repeated at lower dilutions

Sp. out. Lp_2 →	U2	10^3	10^5
578	0	0	0

4/14/54

Is the complex $\frac{1+4^+}{1-4^+}$ phen. (+)? All (+) appear to be secondary - ~~very clearly one (+)~~

SIF # 2746 λ - Single exposure to phage - unacrated fish, sea cat. per cent.

①

Treat	SIF cell	No. phage (+)	No. (+)	No. phage (-)	Total
1. broth	10^6	0	0	67	67
2. 2376 HFT 1	"	9	2 (papillate)	41	52

② Examine (-) a treated plate to see what they are segregating

- Single (-) appearing colony (up to 2 days - then giving a papulating appearance thus \odot , and turning darker in the following day.
- Streaked out and giving

- (+) some of which appear to be segregating (-)
- (-) non papulating (?)
- parental type (-) which papillate

3. 10 (-) colonies picked and streaked against HFT 1 + HFT 4

#	HFT 1	HFT 4	lp Rx	Comment	Allele
1.	+	0	5	} lp to segregating also	4-
2.	+	0	5		4- (?)
3.	mixed (+) and (-)	0	5		4- (?)
4.	+	0	5		-
5.	mixed (+) and (-)	0	5		-
6.	"	"	5		-
7.	"	"	5		-
8.	0	+	5		-
9.	intermed (+) turning dark after 3 days	"	5		-
10.	"	"	5		-

Only Segregants are 4- lp

lp to λ remaining segregating 1-

parental
 dark exsiccum
 from
 parental
 exsiccum
 parental
 parental

4/28/54

902 Mal- → Mal+ = Ap_2^R → Ap_2^S ?

10 well isolated colonies (c) on Mal, tested and found Ap_2^R checked on Mal for reversion exam. (1 / streak) and a gal coated with HFT 1-

1. Exam papillae on gal for instability, indicating Ap_2^S + transduction.
2. Exam Mal+ for Ap_2 reaction.

Original Colony	No. reversions B mal	Ap_2 Reaction of Mal+	No. pap. B gal + HFT 1- (6 picked)	Stability of Gal pap.	Mal + Ap_2 sample characteristics	Transd. of Ap_2 from HFT 1- HFT 2-
1	+	Ap_2^S	31	7/6 stabl	Mal- Ap_2^R	0 0
2	0	—	16	4/6 stabl	" "	0 0
3	+	Ap_2^S	18	0/6 stabl	" "	0 0
4	+ internal (+)	faint reaction	12	5/6 stabl	" "	0 0
5	+	Ap_2^S	14	3/6 stabl	" "	0 0
6	0	—	10	3/6 stabl	" "	0 0
7	+ internal (+)	faint reaction	16	3/6 stabl	" "	0 0
8	+	Ap_2^S	26	1/6 stabl	" "	0 0
9	+	Ap_2^S	12	1/6 stabl	" "	0 0
10	+	Ap_2^S	21	4/6 stabl.	" "	0 0

6 Ap_2^S
2 internal S.

All run transducible

Σ. Mal+ reversions are Ap_2^S or intermediate sensitive

Selection here may not be for the occurrence of Ap_2^S under conditions of extreme purity of cell population but may be for host range mutants in the HFT phage population.

③ Lysozyme made of #1 (unstable, and seq(-))

	0.1 ml /	results
1.	750	solidomer
2.	578	" "
3.		

5/6/54 W2331 mal- tested for transd. with N16 4/9/54 (HFT) not found transd.

5/4/54

J.C. Single cell est.

1. 2342 (1/9/54) + 578 cells
 2.4×10^9
0.5 ml + 0.5 ml

2. Control

0.5 ml Pen +

0.5 ml ul →

Adsorption
10 min. 37°C

Centrifuge
10 min.

resuspended
0.5 ml Pen.

Total Cells No. (+)
 $\frac{213,185}{2.44} = 2.0 \times 10^8$ 16 (18)

del $10^2 \cdot 10^4 \cdot 10^6$ 425 (16) = 3.8% (4)
 dil 10^6

$\frac{401,851}{2} = 4.15 \times 10^8$ 0

24 hour count

It reports no titer present in his sample of cells studied in the microtiter.

5/6/54

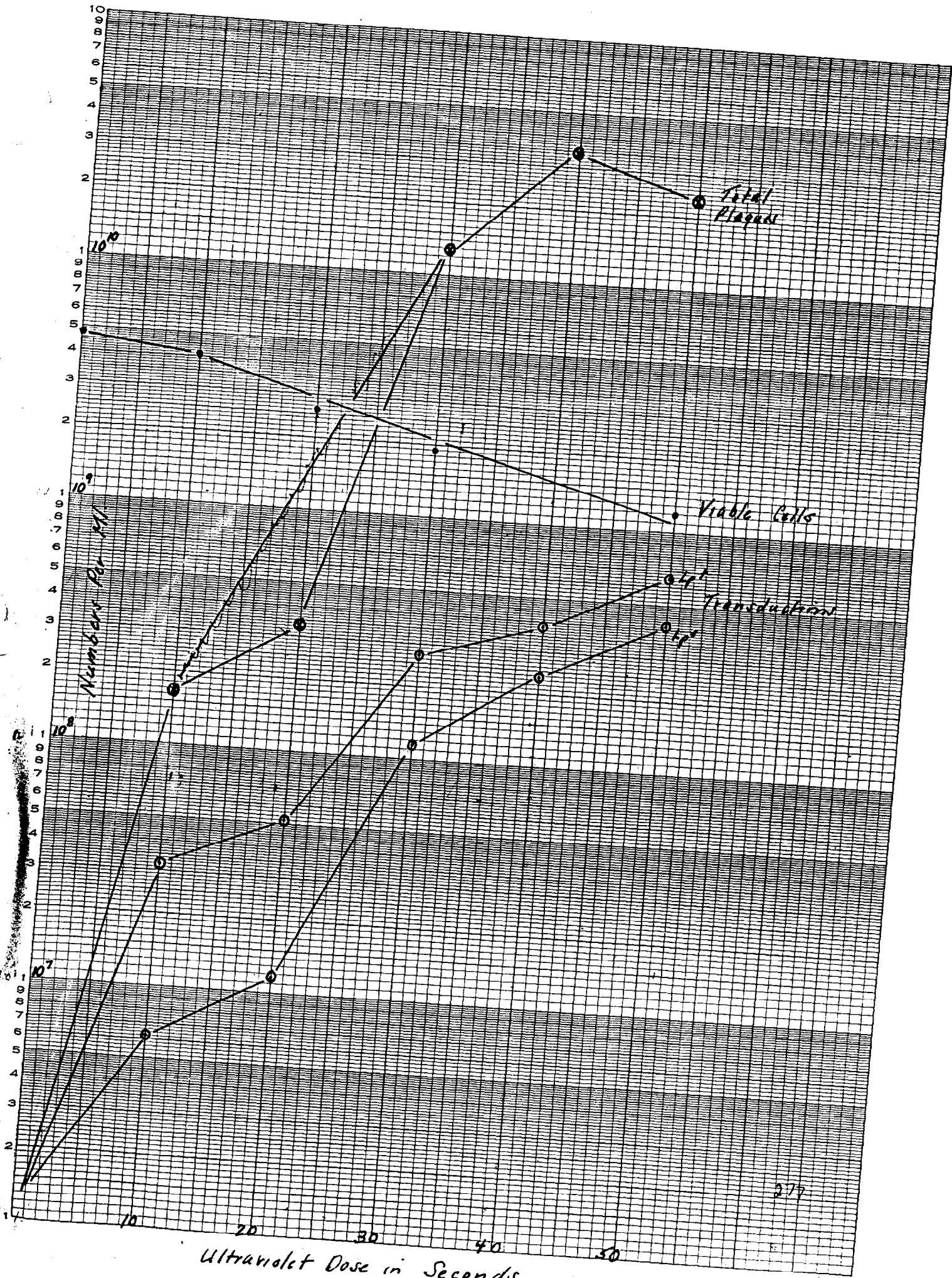
This day 2252x Hft gal⁻ (2477134) entered in stock book as W2487 presumably made and tested c. Mar. 24, 1954 - Culture to TCD at this time also.

5/6/54 For the purpose of studying the transmission of the plaque in crosses. No sugg. by St. Prinst. 2252 (= Hft gal⁻ gal⁺) to $\frac{++}{-+}$ and

then cross to $\frac{++}{-+} \otimes \frac{+-}{-0}$

1. 2252 treated Hft gal⁻ (W116) colonies obtained for "phage control" experiment - some found streaked out.

2. or second streaking, all + still



Numbers Per ml

Total Plaques

Viable Cells

Transductions

Ultraviolet Dose in Seconds

5/26/54

2342. Test on survival to see if inability to get lysates & high titers is caused by application of wrong a.v. dose.

1. Acetated cultures, nearly saturated - centrifuged and respl. in saline

2. Inoc.

	Single plate Cover 6	N/N ₀	Plaque (area) 1-100 dia	Transd./plate x 10 ⁸	Plaque/10 ¹⁰ cells
0	477	1.0	0	50	0
10	417	0.88	103 x 16.16 x 8 x 10 ⁵	112	62 x 10 ³
20	263	0.55	c. 2 x 10 ⁶	164	114
30	194	0.41	120 x 11 x 1.5 x 10 ⁶	1200	115
40	(256) -	-	3456 x 10 ⁵ x 10 ²	2368	2318
50	117	0.25	-	4160	4110

0.1 ml
add back
to
10 ml
Pen
inoculated

ditto
add. in
NBS for
cell assay

1-100

Time	Viable Cells Present	Inoculum Cells	Transd. produced	Plaque produced
0	4.77 x 10 ⁹	0	0	0
10	4.17 x 10 ⁹	0.6 x 10 ⁹	6.2 x 10 ⁶	1.65 x 10 ⁸
20	2.63 x 10 ⁹	2.14 x 10 ⁹	1.14 x 10 ⁷	2.3 x 10 ⁸
30	1.94 x 10 ⁹	2.83 x 10 ⁹	1.15 x 10 ⁸	1.24 x 10 ¹⁰
40	1.59 x 10 ⁹	3.18 x 10 ⁹	2.32 x 10 ⁸	2.46 x 10 ¹⁰
50	1.17 x 10 ⁹	3.60 x 10 ⁹	7.1 x 10 ⁸	2.4 x 10 ¹⁰

Transd. assay
on 8H

no a.v.
Inoc. dose

17

Time	Calculation	Plaque Transd.	Ratio
10'	16 x 200 x 10 ⁴ = 3.2 x 10 ⁷	16.5/3.2	5.2
20'	30 x 163 x 10 ⁴ = 5.2 x 10 ⁷	33/5.2	6.4
30'	169 x 16 x 10 ⁵ = 2.7 x 10 ⁸	103/2.7	38
40'	379 x 10 ⁶ = 3.8 x 10 ⁸	340/3.8	90
50'	654 x 10 ⁶ = 6.5 x 10 ⁸	240/6.5	37.

6/1/54 Transduction of λ_{p}^R with HFT lysates - An indication of host range change in λ ?

1. 1412 ($\lambda_{p}^S \lambda_{p}^R$ Gal⁻) transid \bar{c} 2342 λ
 no add 40
 23% λ 456
 2. 30 papillae picked - after 2 streakings, all stable on Gal
 - #1, #2 3. 24 Gal. papillae " after # 2 unstable - ~~not~~ non- λ_{p}^R in STP both -
 - #3 4. 24 " papillae " " 3 streakings 1 " - ~~not~~ Mal(-), non- λ_{p}^R on STP
 - #4, #5 5. 24 " " " " 2 " - non- λ_{p}^R on STP
- also 23 stable Gal⁺, non- λ_{p}^R on STP

6. Examine Mal + Reversion of above and see if sens.

Mal	Reversion	λ_{p}^R	Gal	Mal ⁺ Revert	λ_{p}^R of Mal ⁺	Revert. of λ_{p}^R	Gal ⁺ of 3
+	+	L	Sen	full	full sens.	2 non- λ_{p}^R	+
+	+	L	Sen	"	"	3 non- λ_{p}^R	-
+	+	R	alg	"	"	1 " "	+
+	+	R	alg	intermed	intermed sens.	" "	+
+	+	R	alg	"	"	" "	+

Suggesto that trans. accomplished by λ_{p}^R forming phage

278

Examine to see if reversion of λ_{p}^R occurring - Tated against λ , STP

Traced. STP	2	3	4	5
1. 5 non- λ_{p}^R	+	+	+	+
2. 5 "	+	+	+	+
3. 5 "	+	+	+	+
4. 5 "	+	+	+	+
5. 5 "	+	+	+	+

Interdict. some cultures look for further phage

Stacks made as 278-1, 2, 3, 4

7/7/54 Lysate made #2 gal⁺ Mal⁺ λ_{p}^R - no phages found before irradiation
 #1 no mod. stack indicates mixed for gal - lysate showed no evidence of phage or trans. in STP

10/2/54 - In looking for whether selection for (Mal- λ_{p}^R) intermediate ^{allele} from admixing λ selected by HFT cell above. Stacks found to be reversion of Mal and λ_{p}^R . Two July λ_{p}^R alg found from #3 #4. (3% revert.) Both found stable such that it is likely more λ_{p}^R are int. differ.

80 λ for hybrid lambda prep. Labeled 6/1/54

1. Titer = $209 \times 10^8 = 2.1 \times 10^{10}$

STP \bar{c} 2342 - Attempts to get higher fraction of transducing

	(-)	(+)	total	cell count	
1. out cell	216	0	216	$\frac{216}{3} \times 10^5 = 7.2 \times 10^6$	
add to					
1. out lysate	206	10	215	$\frac{215}{3} \times 10^5 = 7.1 \times 10^6$	c. 5% (+)

See. 5 min centrifuge present resuspend in 1 ml saline

WS877
 2878
 2879
 2880

6/17/54

279-1 14/2t 234z plate - the sporulation (+) - Revert^A and examine by ⁶MoCh. ^{if picked}

279-2 14/2t 234z plate - 12 picked from by site picking of plate - to see return of ⁶MoCh of the stable gal (+) - by site picking of plate:

1.	2.	3.	4.	5.	6.	7.	8.	9.	10.	11.	12.	13.	14.
1. All 12 mal-gal + were found to be ⁶ MoCh	1. um lps	R ³	s wk	+									
2. DeMalt ^P (of varying degree of +ness)	2. "	R	s	sq									
3. "	3. "	R	s	sq									
4. "	4. "	s wt. confirmed	s wk	sq	rev. ⁶ MoCh	sq + R-							
5. "	5. "	s	s	+									
6. "	6. "	s	s	+									
7. "	7. "	R	s	sq									
8. "	8. "	R	s wk	+	rev. ⁶ MoCh	stbl, + R-							
9. "	9. "	s	s	(-)									
10. "	10. "	s wt. confirmed	s wk	+	rev. ⁶ MoCh	stbl, s-							
11. "	11. "	s	s	+									
12. "	12. "	s	s	+									
13. "	13. "	R	s wk	+	rev. ⁶ MoCh	stbl, R-							
Σ													

Σ 14 spout (+) ⁶MoCh

4 spout + ⁶MoCh
3 " + ⁶MoCh
4 " + ⁶MoCh
Refer to 278

General Summary

sterile
full malt - distribution above not obtained

transd. appears to be gal + ⁶MoCh
gal + ⁶MoCh → sq gal - ⁶MoCh
gal + ⁶MoCh → sq gal - ⁶MoCh

279-3 2580tK12 FT X 1321 3 plates
Control plate ~ 2580tK12 FT for distribution of (+), (-) gal 109F
5K-

279-3 1. Three plates gave "micrographs" early (1st day) - all ⁶MoCh
After 2 days - c. 2000 / plate - 1 apparent (+) instead of (+) on streaking in B gal - Squiggle of gal - noted after 2
6 sequents of this (+) tested against HFT 2, 4 - all transduced by 4, ∴ Gal- from 2580tK12 FT

W 2896

279-4 2580tK12 F X 518 3 plates (discarded one, similar to other because of minimum)

1.	2.	3.
300	241	541
53	46	99
353	287	

30 (+) picked from this cross to find (-) allele - of the 30, 24 were apparently stable (+), of the remaining 6, 25 tested microscopically against HFT 2, HFT 4 - all transd. by HFT 4 ∴ Gal- from 2580tK12 F-

W 2897

6/21/54 1342 purified of 10 colonies examined 7 were HFT on 84 - new stock prepared

For reporting above transmission experiments

1321 transd. by K12 →

1321 tK12 F-

W 2898

Stock - 279-5

6/24/54

280-1 Are the bands of L_p^+ really of L_p^R/L_p^+ nature? Examined on L_p^+ transd. W 25705K12

- ① The Gal (-) seg (20)
- ② The Gal +^u (20)

-(280-1) the Gal- (In General these were from test) all found to be L_p^+ and L_p^R

(280-2) the Gal +^u is found to be L_p^+ and L_p^R - 2 tests were too poor to judge

280.3 Lytic λ - 1485 - 0.5 ml of this lysate into Pan - sterile after 2 days.

① 1485 grown and NSB overnight - then aerated culture in NSB started until nearly full density - centrifuged, resuspended in 2.0 ml SLD lambda (p2.78) tube = 2.1×10^{10} Centrifuge resuspended in 0.20 ml NSB aerated. After 4 hours partially cleared. Centrifuged and checked

② Assay:

$10^7 = 757 - \text{tube} = 7.6 \times 10^7$

③ Transduction with

in P(0) 0.1 ml this lys + 0.1 ml 891 overnight broth cells. - no colonies after 48 hours - No M ductin

④ Coelocution

	Expt/Control	Stability ✓	
1. W 750	2/0		After 2 shakings - 2 unseg stbl (+) directly 9 stbl (+) after 2 shakings
2. W 1210	2/1	2 very faint (+)	
3. W 84	10/14	1, a very faint (+)	

Further evidence that lytic λ does not transduce

averaged 2.242 deleted to find included by in scheme
 had. all o.i.m.e + 10 pm

287-1

Time	Productive Cell Assay	HFT Infective Centers *	Assay Tinned. Centers (=41)	plaque $\frac{1-100}{100}$ $\frac{50}{25}$	Plaque $\frac{1-100}{100}$ $\frac{50}{25}$	A $\frac{100}{100}$	HFT/NFT	Assay of Colony of Infection
0	622	3 0	45 0	97 72	6 0	0	0	$\frac{3}{25} = 0.12$ <small>all data here is others</small>
10	381	193 140	51 2	47 22	11 5	500	2.6	$\frac{10}{25} = 0.4$
20	170	372 364	87 38	38 13	27 41	4100	11	$\frac{9}{25} = 0.36$
30	63	517 516	136 87	18 3	68 62	6200	12	$\frac{9}{25} = 0.36$
40	20	449 446	128 79	25 0	59 53	5300	12	$\frac{4}{20} = 0.20$
60	1	259 256	98 49	26 1	35 29	2900	11	
70	1	230 227	80 31	16 0	24 18	1800	9	

* Some of these infective centers appear to have papillae in them. There also appears to be papillae cont. infective centers about them.

↑ these values cannot be trusted since they represent some growth effect after induction of genome

Time	Total N/N ₀	P/N ₀	T/N ₀
0	1.0	—	— x 3.5
10	0.61	0.31	0.0032 0.11
20	0.27	0.59	0.061 0.21
30	0.10	0.83	0.14 0.48
40	0.032	0.72	0.13 0.45
60	0.0016	0.41	0.27 0.27
70	—	0.36	0.050 0.17

Estimate of fraction HFT

$$\frac{3 + 10 + 9 + 9 + 5}{25} = \frac{36}{25} = \frac{7.2}{25} = 0.29 \text{ HFT}$$

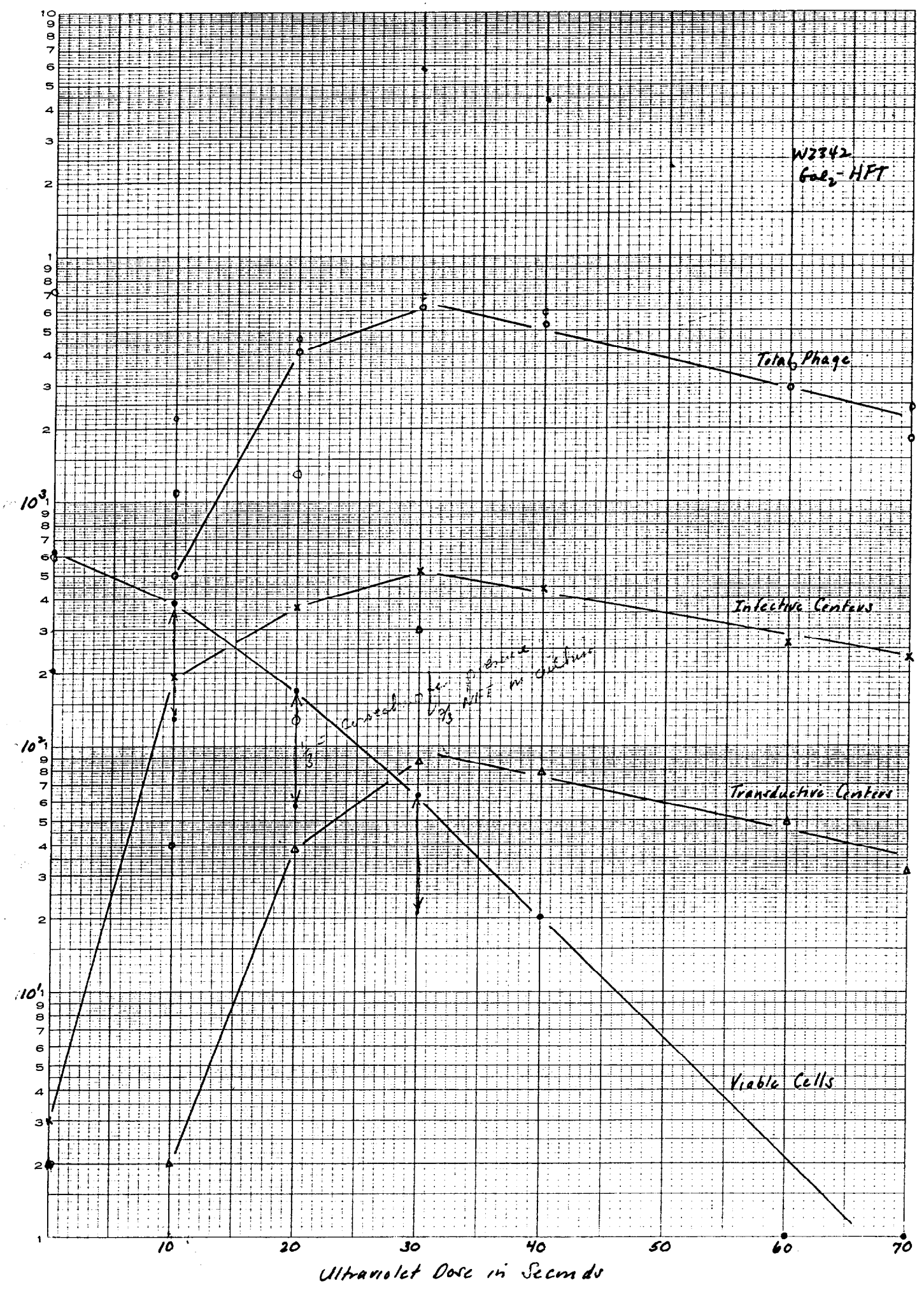
Corrected for LFT

$N_0 = 622 \times 0.29 = 180$

Time	N/N ₀	F/N ₀	P/N ₀
0	1.0	0	
10	0.61	55	
20	0.27	103	
30	0.10	150	
40	0.032	124	
60	0.0016	74	
70	—	66	

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282-1 In the 1st data so far suggests lysates of 1st carry more often than not the (+) of the segment. Really so?

1. 217E750-1 streaked out and 5 colonies picked - cultures made and lysates of these prepared.

Culture	W cell - Assay/750	no cell - Assay/1210	Plaque/SIF	Comment	Ratio ^{pl} /fr	Comment 2
1	1124/448	49 (small)	533x10 ⁶ x 2.1x10 ⁸		2132/497 = 4.3	
Lysate culture 2	460	29 (small)	307x4x10 ⁶ x 2x10 ⁸	(omit 2x as many minutes plaq)	1228/486 = 2.5	Higher titer in Gal ₂ than 2 suggest the fragment is 2- or Gal ₂ ⁺
3	336	36 (")	134x4x10 ⁶ x 5x10 ⁸		544/372 = 1.5	
4	281	57 (")	217x4x10 ⁶ x 2.7x10 ⁷		868/337 = 2.6	
Lysate culture 5	382	34 (")	191x4x10 ⁶ x 2x10 ⁷		764/416 = 1.8	

due to lower
avg. titer
with 1210?
Possibly a
W2175

2. Single colony isolated from each and tested for allele - all were gal⁻

also for
deactivation cultured

W2852 | W2851

282-2 Inbred. of W2851 with HFT 1⁺ → lysate = 282-2
408 (-), 2 (+)

① W2851 cells exposed to HFT 1⁺ (NA-4 prep?) plated out. Many partially lysed colonies after 3 days, 6 colonies showing some evidence of reversion. Other 2 possible inbreds, both predominately (+) in streaks, also appear to be segregating. Original colonies were not (+), but pink in color, also of large size as if host selection advantage in Gal.

② Search for 1⁺ 4⁺
② 1 discarded culture #2 - lysate prepared - streak of culture before down ... 50-50 (+/-) started from (-) colony - assay of lysate (spores) on 575, 1210, 750 - all HFT resistant
③ Single (-) pop. colony (the original) streaked out → single (+) colonies streaked out: 24 in number
④ 6/14 stable (+)

Remember now (-) came from pure (+)

	HFT 1 ⁺	HFT 4 ⁺	Result	Comment
1.	+	+	resist.	Gal ⁻ 4 ⁺
2.	+	+	resist.	Gal ⁻ 4 ⁺
3.	+	+	resist.	Gal ⁻ 4 ⁺
4.	+	+	resist.	Gal ⁻ 4 ⁺
5.	+	+	resist.	Gal ⁻ 4 ⁺
6.	+	+	resist.	Gal ⁻ 4 ⁺
7.	+	+	resist.	Gal ⁻ 4 ⁺
8.	+	+	resist.	Gal ⁻ 4 ⁺
9.	+	+	resist.	Gal ⁻ 4 ⁺
10.	+	+	resist.	Gal ⁻ 4 ⁺
11.	+	+	resist.	Gal ⁻ 4 ⁺
12.	+	+	resist.	Gal ⁻ 4 ⁺
13.	+	+	resist.	Gal ⁻ 4 ⁺
14.	+	+	resist.	Gal ⁻ 4 ⁺

W2851
283-1 →
Discarded 283-2 →
5/13/55

HFT 1 ⁺	HFT 4 ⁺	1 ⁺	Comment
0	+	resist.	Gal ⁻ ①
+	0	"	Gal ⁻ ②
+	0	"	Gal ⁻ ③
+	0	"	Gal ⁻ ④
+	0	"	Gal ⁻ ⑤
+	0	"	Gal ⁻ ⑥
+	0	"	Gal ⁻ ⑦
+	0	"	Gal ⁻ ⑧
+	0	"	Gal ⁻ ⑨
+	0	"	Gal ⁻ ⑩
+	0	"	Gal ⁻ ⑪
+	0	"	Gal ⁻ ⑫
+	0	"	Gal ⁻ ⑬
+	0	"	Gal ⁻ ⑭
+	0	"	Gal ⁻ ⑮
+	0	"	Gal ⁻ ⑯
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+	0	"	Gal ⁻ ㊼
+	0	"	Gal ⁻ ㊽
+	0	"	Gal ⁻ ㊾
+	0	"	Gal ⁻ ㊿

2. parental
Gal⁻ 4
Gal⁻ 6
Gal⁻ 5
Gal⁻ 2
Chromosomal
1
IP
all tested
5/13/55
These are not derived from original
NFT

7/11/54

283

283-1 } from previous page - are presumably Out. - Gen. -
283-2 }

- 283-1 Transl. \bar{c} HFT \bar{c} to get triple \bar{c} .
- (*) Reaction weak - (*) not strong in cross bound - segregation $\frac{1+4-2}{10402}$ is not full + ?
 - (*) Stealing out - suggestion that (*) may be of two classes - one full (*), other unlearned.

A 283-1 cross A = \otimes 2234
B cross B = \otimes 902 gal,

(A) 1st cross not prod - many 6 kg. - no (*) observed

(B) $\frac{1}{8} = 83$, = $664 \times 10 \text{ plates} = 6640$ (-), no (*) observed.

- c 283-1 Further test - transd. 283-1 \bar{c} HFT get + 1 and obtain (+) - this should then segregate phenotypic (-) which revert rapidly - where genetic constitution is $\frac{1+4}{1-4}$
- (*) Transd. \bar{c} STK12 (#73) p. 267 - del 10 = solid media, del 10 = cross, del 10⁶ = 20 (loop full)
 - (*) 24 (+) seg checked out - all appear to be stable (-) - allele = are found 1-4-
 - (*) rest checked 12 (-) seg are stable (-)

283-3 Combination of 283-1 - #2 lysate. The small plaque? a δ mutant. Attempt to obtain lysogenic culture of - Plaques picked and streaked out, colonies b. but

1. 1st 24 colonies found non lys. / 518
2. 2nd / 518

7/16/54

- A Repeat examination of +4 - after check on lysate cell population

Sample ID	Condition	Streaks	1750 CFU	1750 CFU	1750 CFU	1750 CFU
1. 518E902-1	6(1) cl.	sterile	14	14	20	15
2. 518E902-1	reg, mostly +	sterile	0	0	17	13
3. 518E902-1	"	not sterile	9	9	9	4
4. 750E902-1	"	sterile	2	2	9	4
5. 750E902-2	"	"	5	5	11	6
6. 217E902-1	"	"	3	3	11	6
7. 217E902-1	"	"	9	9	15	10
8. 217E902-1	"	not sterile				

dilution for growth

Repeat assay 10⁶ dilution

Sample ID	Condition	CFU	CFU	CFU	CFU	CFU	CFU
1. 518E902	5	0	0	0	0	0	0
2. 518E902-1	0	0	0	0	0	0	0
3. 518E902-1	0	0	0	0	0	0	0
4. 750E902-1	1	0	0	0	0	0	0
5. 750E902-2	0	0	0	0	0	0	0
6. 217E902-1	0	0	0	0	0	0	0
7. 217E902-1	24 (C)	0	0	0	0	0	0
8. 217E902-1	0	0	0	0	0	0	0

no. all	no. all	no. all	no. all	no. all
13	75	111	74	active on 1, 2, 4
18	5	45	8	
10	0	39	2	
24	0	25	0	
32	18	50	13	active on 2, 4
10	0	35	0	
160	147	170	133	active on 1, 2, 4
12	0	30	0	

Assay 250 (1) 10⁶ = 501, 5 x 10⁷
 Control = 0
 (5) 10⁶ = 24 = 2.4 x 10⁶
 (7) 10⁶ = 433 = 4.3 x 10⁷

Presumptive activity, diluted 250 culture, which that assay expect - other experiment about this time suggest this is true activity

284-B

1. 2487 x 2593 - Gary-HF x Geo-Het - to secure dry load? not given (4)
2. Culture given by gelman c 3 hours - plated out 475 loc
 10² dil - c 10³
 10⁴ dil - c 10⁵
 10⁶ dil - c 10⁷⁻⁸
3. Streaks with of prep. c 10⁵ - slightly purple in appearance, large 14 in number
 1. (-) (-)
 2. (-) one no growth
 3. (-) pop. ch. no growth
 4. + ches. + (free)
 5. + " " + (underneath)
 6. + " " + (underneath)
 7. + " " no growth
 8. no growth " "
 9. + also (-) +, also (-)
 10. + " " one (-) col.
 11. (-) minute col.
 12. (-) (-)
 13. (-) (-)
 14. (-) (-)

7/18/54

2799 = Gal, - tp^s - Repeat of 578t HFT, using Gal, cells instead of Gal⁻

1. Actively growing culture of 2799 in Pan. c. 10^8 cells per 0.5 ml cells + 0.5 ml NA-4 prep. 2. Admb 10', dia 10', 10', 10'. Control cells both treated

(a) Control cell. - one of three plates - 155 = $1.55 \times 10^2 \times 10 \times 10 \times 10^4$ 1.6×10^8 no (+), lysed, or phase spread colonies observed, nor any weak (+) colonies

(b) exptl. cell. - one of three plates - 115
 on 38 partially lysed colonies observed
 no transph. 2. colonies papulating in appearance, slightly pink toward similar to appearance of gal⁻ band. = HFT 1 = the 2 colonies in streaking out gave many (+), and parental (-), and possibly un-parental (-). Some of the (+) appear to be segregating

285-1 streaks
 285-2

Picking (a) to examine (+) seg.

285-1	(+) colony	stability	Seg. test	Miles	285-2 HFT	(-) Seg. test	Notes	2nd batch (-) seg
1.	u	s. stabl	0 - +	-	u	parental		picked from gal ⁻ segregates
2.	u	u. unstbl	0 + +	1-	s	-		
3.	u	s	-	-	s	-		9 in all
4.	u	u	+ var 0	4-	u	parental		6 were parental 3 were gal ⁻ tp^s
5.	u	u	0 +	1-	u	parental		
6.	u	u	+ 0	4-	u	parental		
7.	u	u	0 +	1-	u	0 + tp^s		
8.	u	u	0 +	1-	s	-		
9.	u	s	-	-	u	parental		
10.	u	s	-	-	u	"		
11.	u	u	reverting to + also trans. by 4?	parental?	u	"	Many of parental show evidence of lysogenicity in that this is an tp^s transducing segregating (-) seg that are tp^s	
12.	u	u	0 +	1-	u	"		
13.	u	s	-	-	s	-		
14.	u	s	-	-	u	parental		
15.	u	s	-	-	s	-		
16.	u	u	0 +	1-	s	-		
17.	u	s	-	-	u	parental		
18.	u	s	-	-	u	"		
19.	u	s	-	-	s	-		
20.	u	u	0 +	1-	u	parental		
21.	u	s	-	-	u	"		
22.	u	u	0 +	1-	s	-		
23.	u	u	0 +	1-	u	parental		
24.	u	u	0 +	1-	u	"		
25.	u	u	0 +	1-	u	"		
26.	u	u	0 +	1-	u	"		
27.	u	u	0 +	1-	u	"		
28.	u	u	0 +	1-	u	"		
29.	u	u	0 +	1-	u	"		
30.	u	u	0 +	1-	u	"		
31.	u	u	0 +	1-	u	"		
32.	u	u	0 +	1-	u	"		
33.	u	u	0 +	1-	u	"		
34.	u	u	0 +	1-	u	"		
35.	u	u	0 +	1-	u	"		
36.	u	u	0 +	1-	u	"		
37.	u	u	0 +	1-	u	"		
38.	u	u	0 +	1-	u	"		
39.	u	u	0 +	1-	u	"		
40.	u	u	0 +	1-	u	"		
41.	u	u	0 +	1-	u	"		
42.	u	u	0 +	1-	u	"		
43.	u	u	0 +	1-	u	"		
44.	u	u	0 +	1-	u	"		
45.	u	u	0 +	1-	u	"		
46.	u	u	0 +	1-	u	"		
47.	u	u	0 +	1-	u	"		
48.	u	u	0 +	1-	u	"		
49.	u	u	0 +	1-	u	"		
50.	u	u	0 +	1-	u	"		
51.	u	u	0 +	1-	u	"		
52.	u	u	0 +	1-	u	"		
53.	u	u	0 +	1-	u	"		
54.	u	u	0 +	1-	u	"		
55.	u	u	0 +	1-	u	"		
56.	u	u	0 +	1-	u	"		
57.	u	u	0 +	1-	u	"		
58.	u	u	0 +	1-	u	"		
59.	u	u	0 +	1-	u	"		
60.	u	u	0 +	1-	u	"		
61.	u	u	0 +	1-	u	"		
62.	u	u	0 +	1-	u	"		
63.	u	u	0 +	1-	u	"		
64.	u	u	0 +	1-	u	"		
65.	u	u	0 +	1-	u	"		
66.	u	u	0 +	1-	u	"		
67.	u	u	0 +	1-	u	"		
68.	u	u	0 +	1-	u	"		
69.	u	u	0 +	1-	u	"		
70.	u	u	0 +	1-	u	"		
71.	u	u	0 +	1-	u	"		
72.	u	u	0 +	1-	u	"		
73.	u	u	0 +	1-	u	"		
74.	u	u	0 +	1-	u	"		
75.	u	u	0 +	1-	u	"		
76.	u	u	0 +	1-	u	"		
77.	u	u	0 +	1-	u	"		
78.	u	u	0 +	1-	u	"		
79.	u	u	0 +	1-	u	"		
80.	u	u	0 +	1-	u	"		
81.	u	u	0 +	1-	u	"		
82.	u	u	0 +	1-	u	"		
83.	u	u	0 +	1-	u	"		
84.	u	u	0 +	1-	u	"		
85.	u	u	0 +	1-	u	"		
86.	u	u	0 +	1-	u	"		
87.	u	u	0 +	1-	u	"		
88.	u	u	0 +	1-	u	"		
89.	u	u	0 +	1-	u	"		
90.	u	u	0 +	1-	u	"		
91.	u	u	0 +	1-	u	"		
92.	u	u	0 +	1-	u	"		
93.	u	u	0 +	1-	u	"		
94.	u	u	0 +	1-	u	"		
95.	u	u	0 +	1-	u	"		
96.	u	u	0 +	1-	u	"		
97.	u	u	0 +	1-	u	"		
98.	u	u	0 +	1-	u	"		
99.	u	u	0 +	1-	u	"		
100.	u	u	0 +	1-	u	"		

From the above information it appears that these (-) seg. are not derived from separate crosses

7/22/54

Linearity of G_{al}^- in G_{al}^- and G_{al}^- ?

3/17/54
902A

α	Δ	Δ	Δ	Δ
0.025	62	60	357	325
0.05	81	79	415	389
0.075	206	204	725	699
0.100	243	241	1392	1366
0.125	157	155	1734	1708
0.150	130	128	1526	1540

From the above 12 plaques picked - to observe if trans. of tp^+ are by tp^+ particles
 examine bands for lysogenicity and also try to confirm the tp^+ character
 of reduced "lysogenicity" with in G_{al}^- variants. Examine say the one of two you
 are tp^+ or tp^- . Also type one of each to see behavior of + with. Make
 check.

286-1 = 750 E902 | of the 12 picked - after 2 shk. 1/12 mutant (+)
 286-2 = 811 E902 | " 6/12 " (+)

1. In both cases - no clear distinction as regards lysogenicity response
 between transduction and segregants. No cases of tp^+ segregants observed.
 One has assumed that tp^+ particles are transducing tp^+ cells, perhaps the
 heterotypic segregants should be tp^+ . However previous evidence
 indicates that such segregants need not be tp^+ , since lysates were
 made of them. see 811 E902

286-1 titer against 518 10^7 MS = 1 plaque = c. 1×10^7

10 days	10^2	0.1 ml of	10^6	tblr
750 (cont. 10)	177	10^4	0	2.8×10^6
2175 (cont. 11)	63	11	13	6.3×10^4

750 E902

286-2 titer against 518 $\frac{1}{3} \times 10^7 = 73 = 7.3 \times 10^8$

10 days	10^2	0.1 ml of	10^6	tblr
750 (cont. 10)	44	251	2	2.5×10^7
811 (cont. 11)	276	13	14	2.8×10^5

811 E902

||| assay culture

Populace / Plate

*Assay of Gal₂ - 1
in Gal₁ - and
only*

Gal₂ W84

*Gal₁
W84
10X
W84*

Gal₁ W750

1500

1000

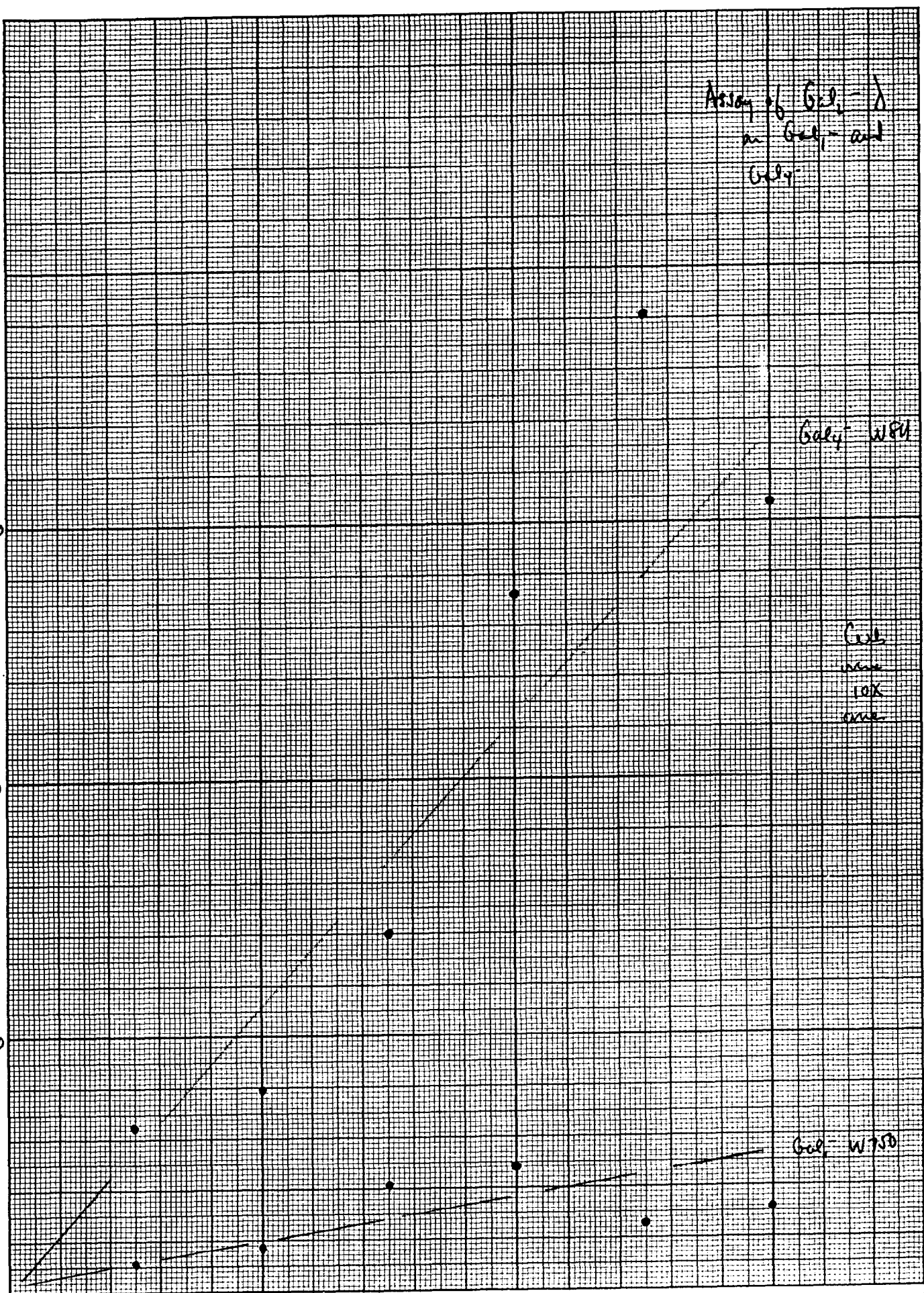
500

0.05

0.1

0.15

ml of lysate plated



9/9/54

2341 Its constitution. Isolated on ~~the~~ ^{Hfr} gel - lp^R - TCN isolated lp^S against gms . Is this an indication of lp^R/lp^S ?

- ① Streaked out and single wt tested.
 - $9 lp^R$
 - $1 lp^S$
- ② One lp^R and ten lp^S grown up and plates B gel to obtain reversions.
- ③ Reversion (obtained from ~~single~~ ^{separate} ~~crosses~~ ^{crosses})
 - ④ lp^R - 8/8 appear to be neg \pm on streaking
 - ⑤ lp^S - 6/6 " " " " stable on streaking

\therefore 2341 is $\frac{2- lp^S}{2- lp^R}$ coming from one following Rx

$$\frac{2- lp^R}{2- lp^R} \times \frac{+ lp^S}{0} \rightarrow \frac{+S}{-R} \rightarrow \frac{-S}{-R}$$

9/10/54 7 ml of 2341 added in stock
 Labeled as 2341R

9/12/54 lysate of 2341 lp^R (made with ant lp^+)
 analyzed against SR
 no plaques
 no transductions

9/13/54

19246KR - Recheck and even \bar{c}^{1436} to see if fragment is $\pm R$ by recovery of $\frac{-S}{+R}$

① possible trans. retained - all non lyogenic 1518
 ② crossed \bar{c}^{1436} - both cultures not aerated. B gel
 3 plates -

$\frac{1}{8}$ of plate = $\frac{106(S)}{39(S)}$

The (-) 23 picked ^{streaked out one} and tested (2 may be ~~80%~~ ^{80%}) - 21 lp^S 2 lp^R

The (+) 24 picked streaked out 2⁺, (-) neg (stab) = 24 (3 appear extra mixed in recovery) ^{streak} mixed
 21 were SR

stock plate of the
 saved for further

9/20/54

2341 - 2344

2344 is a V_1^R derivative of 2341. Since 2341 is $\frac{2-L^S}{L^C}$ what is 2344 with regard to V_1 ?

① taking colonies of 2344 streaked out

17 colonies tested / 11 - all L^S . Suggestion that only L^S seg. of 2341 can be made V_1^R since according to Cavalli suggestion crossing over between fragments and chromosomal genes results in duplicate for V_1 and L^S ?

② Plate 2341 / 17 see if any $L^S V_1^R$ exist.

③ 21 V_1^R colonies picked, purified once and tested against 10 were successful and phage reaction script
7 Lambda Sam
20 L^S

Both repeat with reported Gal as to judge recombination

SP11 - Making HFT by adding fragment?
(5/8 (9/10/54) -

The run with 588 compared with HFT 4 - 48 tested found NFT against 2175

84 (9/20/54)	24	Colonies tested against 2175	- all NFT
single	18	"	- " NFT
run	2	"	- probably HFT - <u>no</u>
9/30/54	20	"	1210 - all NFT

9/18/54

S18

To make HFT blocks of other gal

289-1

u2870

289-1 →

① 98 against EM's 2238 d - of 20 ^{more} py picked (very ^{more} py in exposed portion of plate) found segregating

10 segregants tested are transduced by HFT 4, must be false?
 ^{lytic phase} - tested for transd. / 2297, 811, 0.1 ml of dil 10⁶

② 2070 d - in + a found of 24 tested

84	25	73	33	file
2297	223	313	251	4.8 x 10 ⁶
				9.0 x 10 ⁶

9/22/54

2350 - stock to see contaminated to reverse in

10 colonies tested against 2 of 4 HFT

H make false - HFT by segregation from double dueteri

① after 16 hours in transduction

② " 72 " transduction by HFT 2 - of 4+ reversions?

This behavior is consistent with previous.

9/23/54

Recheck on probability that revision spots give more stable reproduction than wild type as typified by K-12

1. The hands down:

		lysate	Control	0.1 lysate	Stability
290-2	750	750gt-1	0	384	21/24 +u
291-1		2175gt-1	0	80	22/23 +u
292-2	2175	750gt-1	3	540	21/24 +u
293-4		2175gt-1	3	93	23/24 +u

W's
out
stock
out 5/12/54

Storks 290-178

1210 - hand. E 2070A to strain $\frac{2-6t}{2-6-}$
 lysate Control 0.1ml
 290-5 1. 1210 2070A 1 17 9/12 +u

Single day clone of chel- lysate prepared
 7 seg from seg hand. tested against HFT 2 - see wt hand. F 2)

1010 E 8H for Carachi Ept
 292-A Control 3
 811k 62

saved 5/13/51

[2, 3 cultures made (3/11/51)] 291A1, 291A2, 291A3

1. 291A-1 ① 8 day tested by uv. irradiation on 8H - all NFT
 ② 8 spots on 8 day to see if reversions stable ④ c. 8 found stable
2. 291A-2 ① 12 control tubes 0.4 HFT - none HFT ⑤ c. 5 reversions found stable
 ② 10 day as 291A-1 NFT, spotted for reversions ⑥ c. 8 found stable
3. 291A-3 ① 12 additional tested via HFT spot - none HFT ⑦ c. 4 reversions found stable
 ② 8 day as 291A-1 NFT, spotted for reversions ⑧ c. 8 found stable
 ③ 4 reversions found stable

HFT strains 2-
 202-16 } look found stable
 241-14
 241-19

Effect of λ-2 Resistance on adsorption

811 vs. 1439

1. Ept. 0.5 ml K12 λ (7/27/54, Prep 1) + 0.5 ml 10x conc. cell from non-adsorb. cell.
2. λ prep tube 1578

Adsorb. cell titres
 10^7 dil. = $\frac{113,66}{66,16} = 41 \times 10^7 = 4.1 \times 10^8$ (with memory plaque/ml)

8H 10^7 dil. = $\frac{927,793}{860} = 860 \times 10^7 = 8.6 \times 10^9$ cells/ml
 1439 10^7 dil. = $\frac{961,967}{964} = 964 \times 10^7 = 9.6 \times 10^9$ cells/ml

4. Adsorb. tubes contain:
 811 4.3×10^9 cells + 2.1×10^8 λ
 1439 4.8×10^9 cells + 2.1×10^8 λ

5. Assay after 15 min adsorpt. 37C, 10' centrifugation
 811 10^7 dil. = 0.1 = $\frac{0.05}{0.5} \times 10^8$ λ
 1439 10^7 dil. = 85.56 = 70×10^5 λ
 est. $\frac{5}{270}$ remain.

w2888 (292.3 really?)

292

10/10/54

① Reexamination of 257C-6 $\frac{4-2+5}{+ - 72}$, its segregants to study crossing over - 29 colonies picked and streaked out from original streaking of stock. Single (+) seg selected.

	HET 2	HET 4	4	Genotype	STEL	2-	4-	4	Genotype
292-1	1. 0	+	R	2-R	16	0	+	R	2-R
	2. +	0	S	4-S	17	0	+	R	2-R
	3. 0	+	S	2-S	5(4)18	0	+	S	2-S
	4. 0	+	R	2-R	19	0	+	R	2-R
292-3	5. 0	+	R	2-R	20	0	+	R	2-R
	6. +	0	S	4-S	5(1)21	0	+	R	2-R
292-2	7. 0	+	S	2-S	5(2)22	0	+	partial S	2-seg? 292-22
	8. 0	+	R	2-R					
	9. 0	+	S	2-S	(1)				
	10. 0	+	R	2-R					
	11. 0	+	R	2-R					
	12. 0	+	R	2-R					
	13. 0	+	R	2-R					
	14. 0	+	R	2-R					
	15. 0	+	R	2-R					

Editype 4-S = 2 $\frac{1}{2}$ +^{one} S61
 Allotype 2-R = 15 $\frac{1}{2}$ +^{one} S61
 Allotype 4- = 4 $\frac{3}{2}$ +^{one} S61
 Allotype seg? 2- = 1 ? not done

CONTINUED 292A

See also 29F

② Transduction of 4^r - 1924 x gal²- \rightarrow gal²- 4^r . Culture is W1 from Egypt W. Is this culture heterogenic in any way? Plated out on B gal and revertants examined. 12 picked.

- 6 obviously slow (+)
- 6 fast (+). Of these 1 appears to be seg.

292-3

3 The seg clone of 2 above - 3 col examined for *Stamboda* reaction

- seg +
 - pure +
 - pure (-)
- 2279 All non lysogenic

Attempt to isolate $\frac{2-4^r}{2-4^r}$ and see character of phage

③ 1503 4^r - Inoculation of gal 4 - After c. 100 colonies examined from single colony experiment (cells + HET 4 plated out) - failure to find (-) spotting on B(0) of cells and HET - After about 30 streakings out - 2 (-) observed also a segregating (+), perhaps $\frac{+}{-}$ stock of (+) made $\frac{+}{-}$ (292-1) discarded

Fast of the gal ~~from 292-1~~ 292-1
 transduced by HET 1, 2, 4 \rightarrow seg (-) in spontaneous gal²-
 116 11/153 292-7

292-1, 22

292-1 - The original part of this in association with the trans. test was pure (-), no segregation of population, (+)

- on streaking out for colony test, all of 4×10^8 segregating - streak found mixed (+), (-). Pure (-) colonies tested against 1 and 1 (+)

- ① The (+) - weakly sensitive, (segregating probably 4×10^8)
- ② 1 of 10 (-) colonies sensitive 4×10^8 # 6 (+), #7 from beginning
- ③ ~~no above (-)~~ spotted ~~against~~ on Gal to check reversin stable
 - { 6 reversin
 - retained: 2
 - Normally unstable
 - Others may be
 - Streaks incubated
 - to long

292-22 - The case obviously segregating 4×10^8

- ① 10 (-) colonies tested 7 out of 10 4×10^8 #s 1, 2, 5, 6, 7, 9, 10
 - { reversin found
 - seq = #8, #3, #4
 - hem. seq = #1, 5, 6, 9, 10
 - no reversin obtained
 - in others (2, 7)
- ② The above (-) spotted to check reversin stability.

2734

ETL released a better growing isolate for this and labeled 2734

Apparently two cultures of 292A-1503 retained and from the trans. fold of 292 original = W2733

1503 In search of Gal^r Het 1503

9-20/84 From the spotting of HET 4 on 1503 a (-) obtained not trans. E HET 1, 2, 4, apparently sensitive to 1 (less so than usually found)

Cross with 2279 - Gal^r - Gal^r to see 1. un-Gal^r, 2. diploid obtained

- ① cultures on B Gal. after 3 days both show occurrence of reversin (papillae)
- ② The cross of Gal.

	(+)	(-)
1.	12	121
2.	296	57
3.	7	81
4.	17	127
5.	4	56
6.	23	160
	69	592

Actual no (+) may be higher since small colonies appear to be untrans. in appearance and may be immature (+), wanted here a (-).

$$\frac{592}{514} = 11.4\% (+)$$

12 (+) obtained and found stable for Gal

10/17/54

Search for HFT 4 (After many unsuccessful attempts)

2175 ^{SSU} - trans. plate = 40/10 (old lysate)

of 15 segregants tested (transd. saved also to see if only contain trans. gene due to HFT, as distinct from a new segregation event ~~found~~ from any transduction - two found to be HFT in 2175 stocks mated and also checked for purification (results could also be certain E (+))

Stocks: 293-11 = transd., 293-11S = HFT 4 }
 293-13 = " , 293-13S = " " }

also found 293-12 - lysate prepared 7 12, 13 (from lysate once cultures were vacated 56 hours before induction),
 - uninduced lysate streaked against 2175 ^{lyse 12} ¹³
 8H ⁺⁺ ⁰ ^{??}

SEE 305

11/2/54

293-1, 2, 3 all transductions from which HFT seg were obtained
 Behavior of transductions in giving HFT or NFT - Separate segs from disp. (1)

Seg. packet sprinkled 1210 SA and used 15 sec.	293-1		293-2		293-3		293-11		293-12		293-13		280/100	total
	seq. #	SS	SS	SS	SS	SS	SS	SS	SS	SS	SS	SS		
2.	0	(30)	1	6	0	(18)	0	2	(50)	0	0	(18)	0	0
3.	2	5	0	9	0	(13)	0	7	(55)	20	0	0	0	0
4.	0	(18)	1	(17)	0	(23)	0	3	(53)	20	0	0	0	0
5.	0	(14)	0	(55)	0	11	0	3	0	(12)	0	0	0	0
6.	0	(24)	3	3	0	4	0	3	(18)	2	0	(C. 70)	+	0
7.	0	(12)	0	(55)	0	(30)	0	6	(55)	0	0	(C. 60)	0	0
8.	0	(10)	1	(12)	0	6	0	3	(8)	6	0	(C. 60)	0	0
9.	0	(10)	1	1	0	2	0	1	(55)	2	0	(C. 20)	0	0
10.	0	(10)	0	(10)	0	(10)	0	0	(55)	18	0	0	+	0
11.	0	(10)	0	7	0	(30)	0	0	0	1	0	0	0	0
12.	0	(10)	0	7	0	(30)	0	0	0	1	0	0	0	0
		1 HFT 2		2 HFT 2		1 HFT 2		7 HFT 4		3? HFT 4		3		0

Request only	1/100	B/100	1/100	1/100	1/100	1/100	1/100
1.	(55)	1	0	0	SS	25	0
2.	2	5	3	1	SS	11	0
3.	0	4	2	2	SS	6	0
4.	0	6	3	2	SS	C. 40	0
5.	0	(55)	0	1	-	3	1
6.	1	4	1	1	-	0	1
7.	1	(55)	0	2	SS	C. 30	0
8.	0	4	0	0	-	2	3
9.	1	4	2	(55)	-	0	2
10.	0	3	4	0	SS	C. 50	0
11.	0	3	1	0	0	1	1
12.	0	3	0	0	SS	12	0

293-12A substituted for W02431 as HFT 4 stock
 E. coli picked and streaked out and found HFT on 1210 and used & lysated

12/16/54

11/7/54

Recombination between tp^s to give λ

- Attempted by growing tp^s cultures together - tp^s cultures known to have differences, 2281^{non} different to lysogeny, 2373 gives stable transductions more frequently, etc. All tp^s believed to be separate mutational events or cases where the tp^s had been in contact with transducing phage and had emerged as tp^s again
- Proced. - cultures given in all combinations in peracry. Starting from 0.1 ml of overnight culture. Cultures centrifuged after 24 hours and loopful of supernatant spotted on 2281

3. Set up

	A	B	C	D	E	F
	W2281 ^{FT}	W2373 ^{FT}	W578 ^{FT}	W2344 ^{HFT}	W1673 ^{FT}	W2280 ^{FT}
	Gal ⁻	Gal ⁻	Gal ⁻	Gal ⁻	Gal ⁺	Gal ⁻
A +		+	+	+	+	+
B +			+	+	+	+
C +				+	+	+
D +					+	+
E +						+
F +						

+ = culture or combination

- Results
 - Spotting on 2281 showed no obvious evidence of λ plaques
 - Culture transferred and given to naturalist - loopful spotted on 578 and indicated no plaques found on indicated plates

11/24/54

At Luca Corradi's suggestion transduction of HFT or FT [\in HFT]

- W1321 (SFT⁻)
 - untreated
 - 0.3 ml ~~of overnight culture~~ of overnight culture ~~10x~~ } incubated 10' at 37, dil to 10 ml both added
 - as 6. only using 269-1 by date = 2342 + K12 HFT HFT with the same conditions.
- a, b, c incubated at 37C overnight, centrifuged 100g in saline.
- a, b, c mixed with W945 S. gal.
a, b, c also spotted on Sgal separately, 945 as well, as controls, no further
- After 3 days in suggestion of protoplasts - indication of no F⁻ detection or HFT detection

1412 Attempts to transduce ϵ λ -2

1. Intermed. malt⁺ of (Mal-lyc^a) have interaction with λ -2 such that transductions might be obtained?
2. 8 Mal+R obtained with varying degrees of +-ness
1 = extreme (+) 8 = (-)
3. λ -2 prepared by gamma on 1405. 0.1 ml plated 3 fold

Row	Phenotype	Control area	Lysate area
1	(++)	8	19
2	(++)	6	18
3	+	5 (small blue)	19
4	±	10 (small blue)	29
5	±	10	29
6	±	8	16
7	(-)	14	25
8	(-)	20 *	26

* note the streptomycin of this plate and plate does evidence of lysin by the phage.

W2872

1210 transductions of, wells 1-4 lysate. ADDITIONAL PCW - 297

1. lysate 283-1 (appears to be a good lysate, cleared promptly at 2 hours, viscous)
2. control lysate 81 after 24 hours no signif. diff. lysate >> control.

The transd. these (-) came from

Row	295-1	295-2	295-3	295-4	295-5	295-12
1	+	-	+	+	-	+
2	-	+	-	-	+	-
3	+	-	+	+	-	+
4	+	-	+	+	-	+
5	-	+	-	-	+	-
6	+	-	+	+	-	+
7	+	-	+	+	-	+
8	+	-	+	+	-	+
9	+	-	+	+	-	+
10	+	-	+	+	-	+
11	+	-	+	+	-	+
12	-	+	-	-	+	-
13	+	-	+	+	-	+
14	+	-	+	+	-	+
15	+	-	+	+	-	+
16	+	-	+	+	-	+
17	+	-	+	+	-	+
18	+	-	+	+	-	+
19	+	-	+	+	-	+
20	+	-	+	+	-	+
21	+	-	+	+	-	+
22	-	+	-	-	+	-
23	+	-	+	+	-	+

W2853

E: 15 sides 3 also

Row	HET 1	HET 2	HET 4
1	+	-	+
2	+	-	+
3	+	-	+
4	+	-	+
5	+	-	+
6	+	-	+
7	+	-	+
8	+	-	+
9	+	-	+
10	+	-	+
11	+	-	+
12	+	-	+
13	+	-	+
14	+	-	+
15	+	-	+
16	+	-	+
17	+	-	+
18	+	-	+
19	+	-	+
20	+	-	+
21	+	-	+
22	+	-	+
23	+	-	+
24	+	-	+
25	+	-	+
26	+	-	+
27	+	-	+
28	+	-	+
29	+	-	+
30	+	-	+

W2853

possibly HET or control. No indicate contamination

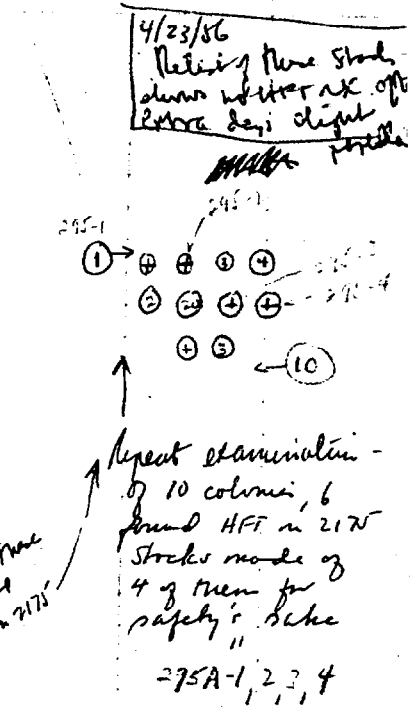
* These machines for HFT
 required a day
 extra for fuel development
 suggesting that the
 is not fuel (+)

295A
 17424
 14422

F = fuel
 N = week
 1210 + 283-1 Continued.

Qty	HFT1-	AFL-	AFT 9-
31 59	+	F	+
32 59	-	+	+
33 59	+	F	+
34 59	+	F	+
35 59	Cont	+	+
36 59	+	F	+
37 59	+	F	+
38 59	Cont	+	+
39 59	+	F	+
40 59	+	F	+
41 59	blue	+	+
42 60	+	F	+
43 61	blue	+	+
44 62	blue	+	+

39	+	-	+
30	+	-	+
31	+	-	+
32	+	-	+
33	+	-	+
34	-	+	+
35	+	-	+
36	+	-	+
37	+	-	+
38	+	-	+
39	+	-	+
40	+	-	+
41	+	-	+
42	+	-	+
43	-	+	+
44	+	-	+
45	+	-	+
46	+	-	+
47	+	-	+
48	+	-	+
49	+	-	+
50	+	-	+
51	-	+	+
52	+	-	+
53	+	-	+
54	+	-	+
55	+	-	+
56	+	-	+
57	+	-	+
58	+	-	+
59	+	-	+
60	+	-	+
61	+	-	+
62	+	-	+
63	+	-	+
64	+	-	+
65	+	-	+
66	+	-	+
67	+	-	+
68	+	-	+
69	+	-	+
70	+	-	+
71	+	-	+
72	+	-	+
73	+	-	+
74	+	-	+
75	+	-	+
76	+	-	+
77	+	-	+
78	-	+	+



52 idis - + 6 = 58 idis
 8 allos + 1 = 9 allos
 2 amphi = 2 amphi

(45) 63	+	-	+
(46) 64	+	-	+
(47) 65	+	-	+
(48) 66	+	-	+
(49) 67	-	+	+
(50) 68	+	-	+
(51) 69	+	-	+

Batch II

1.	+	-	+
2.	+	-	+
3.	+	-	+
4.	+	-	+
5.	+	-	+
6.	+	-	+
7.	+	-	+
8.	+	-	+
9.	+	-	+
10.	+	-	+
11.	+	-	+
12.	-	+	+
13.	+	-	+
14.	+	-	+
15.	+	-	+
16.	+	-	+
17.	+	-	+
18.	+	-	+
19.	-	+	+
20.	+	-	+
21.	+	-	+
22.	+	-	+
23.	+	-	+
24.	+	-	+
25.	+	-	+
26.	+	-	+
27.	+	-	+
28.	+	-	+

49 idis
 4 allos
 1 amphi

79 idis
 5 allos
 1 amphi

found HFT
 one of these
 found HFT in 2175

All listed

	1	2	4
1.	-	+	-
2.	-	+	-
3.	-	+	-
4.	-	+	-
5.	-	+	-
6.	-	+	-
7.	-	+	-
8.	-	+	-
9.	-	+	-
10.	-	+	-

1412 Repeat transduction with HFT 2 to observe the nature of transductions

1. 70 odd $\frac{1}{2}$ = 41
HFT 2 (2342 phage)¹¹ = 317
2. 24 ^{Batch I} populus picked and streaked 3 fold (2) - all stable, all non lysogenic / STB, are λ^+
(2) revert dual test λ^+
3. 22 ^{Batch II} populus picked and streaked 8 fold (2) - cell stable, all non lysogenic / STB
4. 24 ^{Batch III} populus picked (2) - all
5. 24 populus picked (2) 1st found, all non lysogenic / STB (296-1)
5. 24 populus picked (2) 3rd found, all non lysogenic / STB (296-2, 3, 4)

W2881
W2882
W2883
W2883

Expt 1 experiment

HFT 2	HFT 4
3	0
8	0
0	0

6. tests of 296-1, 3, 4 - ~~these~~ fold - seq obtained from each and brushed against HFT 2, 4 after 2 days

	HFT 2	HFT 4
1	4 pop	0
3	1 pop	0
4	3 pop	2 pop
2		

Since this HFT 2 is the host to make the original selection these results do not indicate that selection has been for a mutant of the cell λ

7. 296-1, 3, 4 fold seq used as indicators in plating with conc. (> 10⁹) particles of λ in the hope of selecting host range plaque mutants. None found on either 1, 3, 3, 4

Re number	Batch I #	Appearance of fold on Hal	Reversion observed	Reversion % o type	Gal transd. λ^+	Mal + e λ^+ / STB	revert λ^+
	1	pink	+	free +	revertant	S	..
	2	blue	+	free +	"	S	..
296-5	3	pink	+	free +	"	S	..
296-6	4	blue	+	free +	"	R	..
296-7	5	pink	+	free +	"	S	..
	6	pink	+	intermed (+)	"	S (weak)	..
	7	pink	+	free +	"	S	..
	8	"	+	-	"	-	..
	8	"	+	free +	"	S	..
	9	"	+	free +	"	S	..
	10	"	+	intermed (+)	"	S (weak)	..
	11	"	+	free +	"	S	..
	12	"	+	free +	"	S	..
	13	"	+	revert (+)	"	S (weak)	..
	15	"	-	-	"	"	..
	14	"	+	free +	"	S	..
	15	"	+	free +	"	S	..
	18	blue	-	-	"	"	..
	19	blue	-	-	"	"	..
	16	pink	+	free +	"	S	..
	21	pink	-	-	"	"	..
	17	pink	+	free +	"	S	..
	18	pink	+	free +	"	S	..
	19	pink	+	revert	"	S	..

W2884
W2885
W2886

K-12 lysate 11/12/54 - hand tube about 150/ml^{0.1}
Agar 1210. Papillae slow in development. 3 days
required and small. Other 'decidua' with 1210
were also slow at this time. 1210 bad medium?
See Paou

1210 handlined with 283-1 - same ¹²¹⁰ culture as ↑

7. no add 3
2. 283-1 0.1ml 71

SEE 297A

K12 lysate above. Test of layer plating method.

Assays. 0.1 ml lysate + 2.5 ml Bgal (0.6% agar) poured in Bgal plate

817	① no add	38	
	② 0.1 ml	294	256
1210	① no add	4	
	② 0.1 ml	44 very small	40
750	① no add	4	
	② 0.1 ml	34 (quite a few small)	32
2289	① no add	7 c. 400 small plaques)	
	② 0.1 ml	1/2 = 162, 1296	

Continue checked culture?

11/30/54

For J.L. tests of some segregants from P₁ hybrids?

① 3-14x Gal - against HFT 1, 2, 4 - cures not fol - no suggestion of heritability

② lysates of cures TCN II-2 and TCN VII-94, also lysate of 583.

lysate	1700	2125	PA	2307	3-14x
control	0	2	0	0	not
583	0	49	9	0	not
II-2	1	34 *	1 *	0	fol -
VII-94	c.120	c.150	c.120	c.100	

+ a little chloroform transferred to plate causing inhib. of growth

Attempted heritability of elimination of Arabidopsis and also direct attempt at (Mal-Lp²).

1. 2347 = 583 mal - ara - Lp² Lp⁵ SR ← obtain heritable part for 5th marker (this results in disease of HFT)
 on Basal control 0.1 ml HFT 2 } no heritability

2. 2307 = 583 mal + Lp⁵ ara - 5th
 on Basal control 0.1 ml HFT 2 } 4 days

2307 against HFT 1, 2, 4 - Trans. were by 2, 4, less by 1 after 3 days

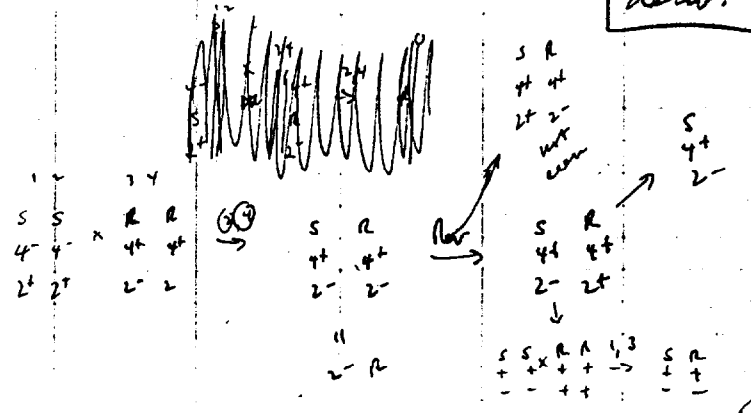
257C-6 - Combination. see also 292

seg	from HFT 2	from HFT 4	A. Recomb.	Seg from HFT 2	Seg from HFT 4	Page 292	Line
1	-	+	R	2-5	2-5	2	3
2	-	+	S	-	-	6	16
3	-	+	S	-	-	2	4
4	-	+	R	2-2-5	2-R	1	?
5	-	+	R	3	-		
6	-	+	R	4	2-5		
7	-	+	R	5	2-5		
8	-	-	S	(Re)	-		
9	-	+	R	7	2-R		
10	+	-	S	two include			

298
298-8

On the basis of the reversion study the 2-R were derived from

W2890 also got derived.



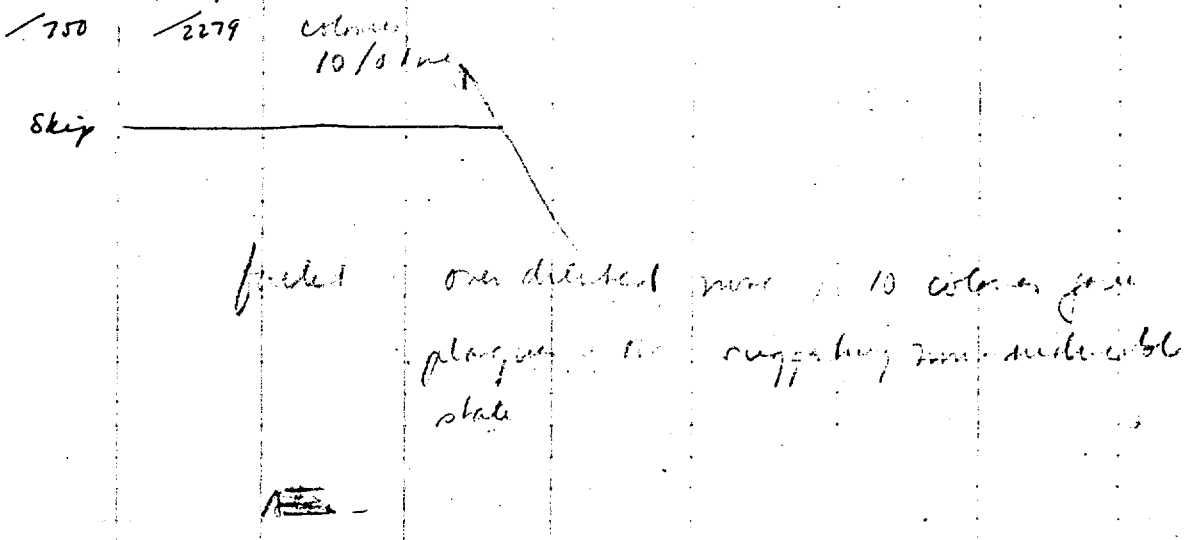
12/7/54

One step = HFT 241-14

7. 241-14 streaked out single colonies picked to both and tested for HFT by read. at 750. Broths saved. After determ. which were HFT, 2 ~~broths~~ were added to 10 ml saline, incubated 15 min at 37C. Fairly turbid = $<10^7$ / ml. Irradiated 45 sec to obtain approx. 10^{-2} survival. Dilute $0.1 + 10$ sal. = $<10^5$ ^{trials} dil. $0.1 + 10$ sal. = $<10^3$ $1.0 + 10^{100} = 10^2$

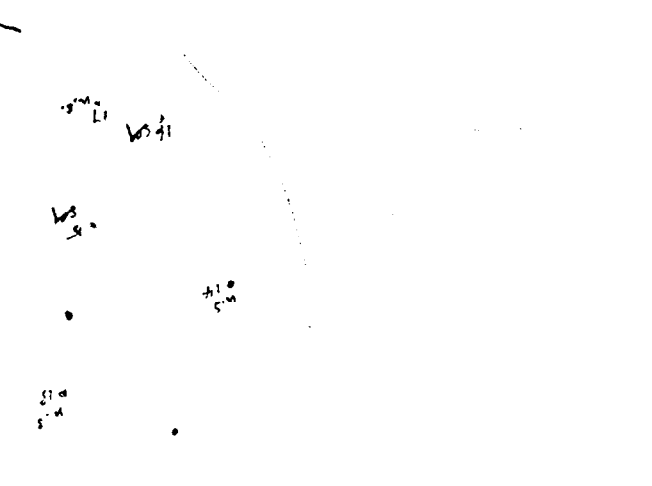
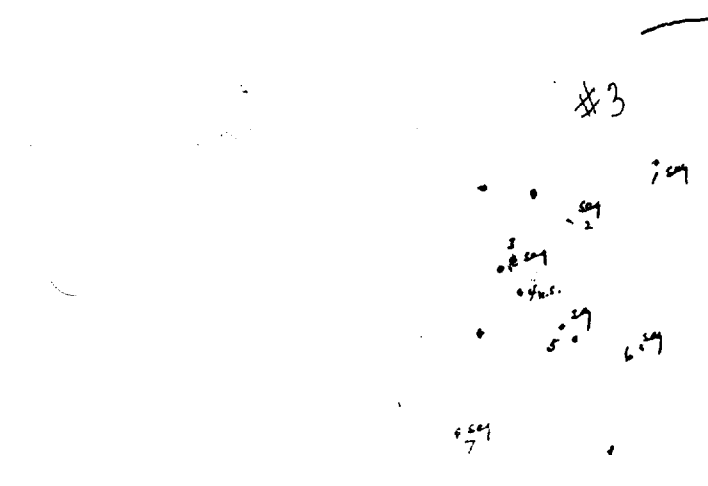
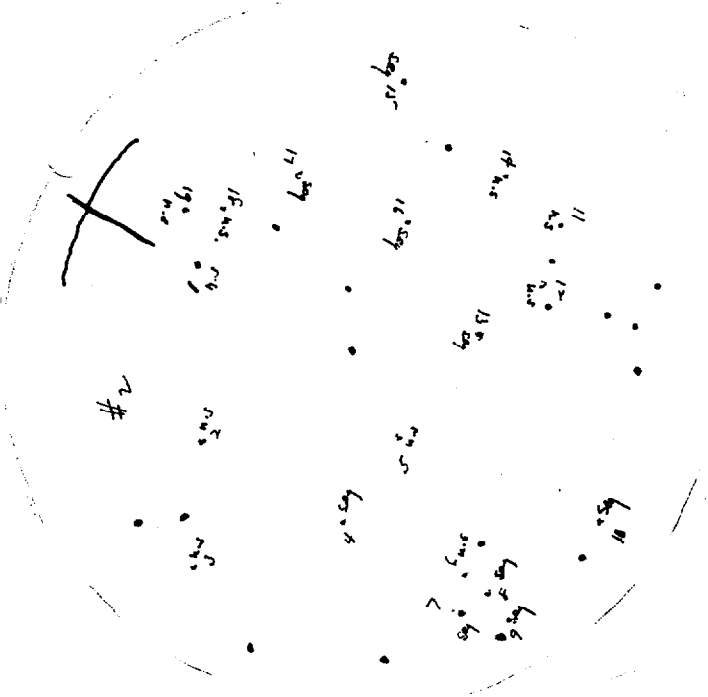
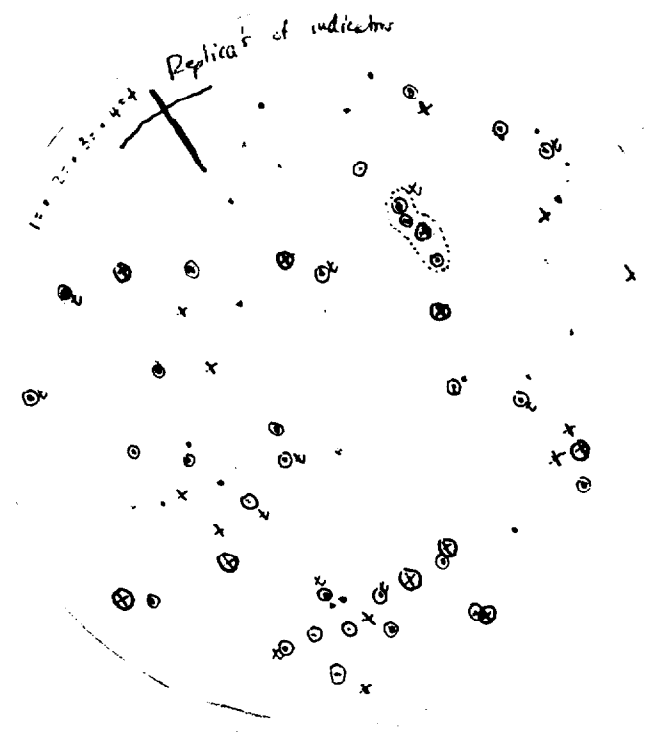
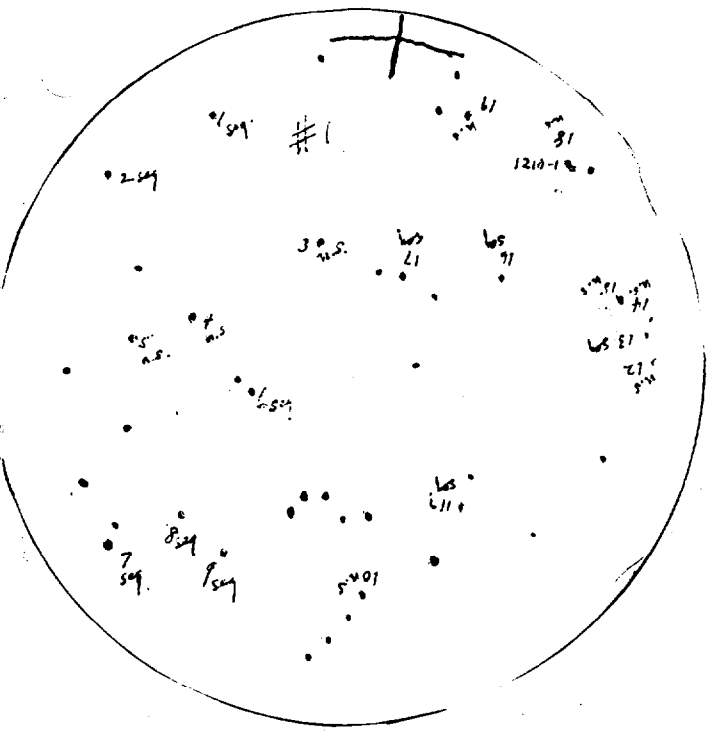
Assay against 750, 2279

Time
10:24 0
5
10:34 10
10:39 15
10:44 20
10:49 25
10:54 30
10:59 35



~~10:04 40~~
~~10:14 45~~
~~11:14 50~~
~~11:19 55~~
~~11:24 60~~
~~11:29 65~~
12:29 125

2. 241-14 - attempt to make 6×10^8 .
 Streaked out on λ -2 on B mal.
 1. Mostly Mal + seen -
 2. 2 mal - col picked and streaked out
 3. purest mal - streaked on B. gal - all gal (-)
 4. Repeat
 Supernat. broth of culture used to make this = 2×10^5 hand? 0×10^6 plaques
 (B) Repeat assay
 5. on next run several mal - obtained - tested for HFT all HFT except 1
 6. Stock made called (241-14 Mal)
 1. lysate 12/18/54 53×10^7 plaques / 2279 = 5.3×10^8
 hand. assay $13 \times 10^7 = 1.3 \times 10^8$
3. 202-16H - / 100 241-14 above. plated against λ -2, mal - obtained. Tested 2/7 col as HFT against 750 - For use in one step growth culture



12/11/54

Are the NPT cells giving ~~plate~~ transducing phage closely related?

1. W1250 - spread on B gal for overnight culture.
2. Incubated about 12 hours ~~over~~
3. replicated to 4 prewarmed B gal
4. after 30 min. 4 replicas inoculated 20" - 50 cm sterile loops, incubated 10"
5. Phaded plates replicated to B gal plates spread with W750 cells as indicator of transduction. W750 cells = 1/2 dilution, 0.1 ml of overnight culture, spread 1/2 hour before replication
6. Result of replica comparison - attached figures.
7. All colonies on indicator plates picked, streaked out twice to see which are segregating

12/12/04

Observation on 292-3 = 4+2-5 (from 510 by 2-X)
against 295A-4 = presumed HFT 1-4- do all of initial complex
to phen. (+).

1. c. 10³-10⁴ cells exposed to lysate of 295A-4 (later unknown) 1.0 ml, etc.
15 minutes, 0.1 ml plated on 8. gal. Both exposures used as
control

1. Control = 3x10 ³ (+) col.	After 3 days
2. control	c. 12 1/2 on column region c. 12 reversions
3. 295A-4 = 201-301 col	c. 14
4.	c. 12 representing ?
	c. 8

2. Presumably lysate not active either through low titer or HFT
nature - Repeat = good lysate.

Observation on 2307 = 583 mult by 2³ fold - by +

W2874

302

1. 2307 diluted to about 10³ cells/ml
Exposure.
a. Growth - Control
b. 1.0 ml 2342
c. 1.0 ml NA4 prep

2. Exposure

1. Growth	c. 400 (-) col.	2 days	c. 5 reversions	3 days	
2. " "	" (-) col.	"	"	"	
3. HFT 1	c. 500 col (-)	0	0	c. 2000 - ab. mult. colonies populating	
4. " "	"				
5. HFT 2	"				

For further
investigation
see p 309.

302-1A = 59
302-1

Colonies picked
and checked - Seq. for agent HFT 2 all had,
or 7- Spiked and used on 8/1, apparently
all HFT - Do lysates and see

c. 1 pop. - ab. at the end of 2 days all
colonies regularly populating and reacting to (+)
colonies almost sterile appearance
because of (+) growth. At the end of 3
days c. 70% of colonies (+)

4. Exposure. 1-7+ phenotype also
2308 = 1/2 2307

302A

1. Sept. interrupted for about a week - 2308/HFT (2)
3000 populating plate with colonies of about 1000 cells - Controls about 56
per the same number

2. Pick the presumed (+) and all of 1-7+ phenotype can be described
1. the phen. is mod. slow shaker mt, 24 clonal cell picked and checked
out to show definite change to (-), 22 mixed streaks obtained, 2 pure (-)
shaker

Some of
the (+) appear
to be the
parent
(± +)

Single (+) picked from each of the 22 shakerings
1st (+) checking 7/22 shows "slight" reversion, i.e. few (-) col.
3rd (+) " 3" apparently stable, "slightly" neg., one col.
" 1/5 neg., stable (+)

SEE PAGE 307

12/12/54

Observations on t_p^s transductions and stable transduction

$\left\{ \begin{array}{l} 292-3 = \text{ST gal}_2 - \text{Array of H-12 d on 0.1uc} = \\ 298-F = \text{ST gal}_2 - \text{gal}_4 - \\ \text{ST also} \end{array} \right.$

285-2. Strained out on $B(d)$ after 3 days, 20 colonies present

① Pure (\pm) were 8 in number. Presumably derived in part directly from $\frac{+}{+}$

combination	λ_{RK}	mate	Resistant to tetracycline
1. + - -	S	1-	✓
2. + - -	S	1-	✓
3. + - -	S	1-	✓
4. + - -	S	1-	✓
5. + - -	S	1-	✓
6. + - -	S	1-	✓
7. + - -	S	1-	✓
8. - + -	R	1-	✓

cistrons mixed up and partially lost, in attempting to recover, tested against λ , $\lambda_{14}(\pm)$ and gal_2 , λ^s 3/3 (\pm) $\lambda^s(?)$

apparently 4 by λ^s lost somewhere during this procedure. Correct!

Same one as in 1st 7 STP

Gal (\pm)	but segregating?	segregant	Seg. t_p
1. R	yes	1-	S
2. S	no	—	S
3. R	yes	1-	S
4. R	yes	1-	S
5. R	yes	parental?	S
6. F	yes	1-	S
7. R	yes	—	S
8. R	yes	1-	S
9. R	yes	parental	R
10. R	yes	1-	—
11. S	no	—	—
12. S	no	—	—

continued

298-8 - Gal⁺ Gal⁻ Lys⁺ from 257C-6

From the purpose of settling two problems, possibly?

1. Lys⁺ transduction?
2. Stable transduction how many?
3. Size of the fragment.

1. 298-8
 200 add. 4 days (Lys⁺ trans) 0? (perhaps 50 very small)
 K12(11/12/54)

A

a. This is odd since in 297, this lysate approximated at 150/0.1 ml
 found 250/844
 40/1210
 342/750
 1300/2099
 this lysate tested against 238
 1/8 = c. 2000 = ltr. 2000
 control = 0

304A-

b. the same prep, inhibited by something?

① 6 patches	5 found to contain (H) in first streak	② 3 found segregating (one of these appears to be stable)	2	4	1/4	1/4
1	-	-	-	-	-	R
2	-	-	-	-	-	R
3	-	-	-	-	-	S
4	+	-	-	-	-	R
5	+	-	-	-	-	R

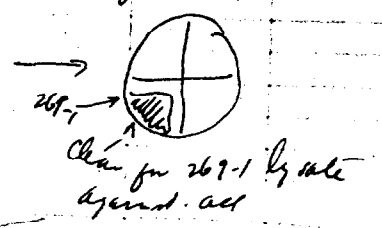
N2892
 1/2 R 2+4+ / Stocks:
 1/2 S - - / 304A-1

B

269-1 - because of EML's finding of unusual phase in lysate of this.

1. prep of 269-1 obtained and handpicked and 10⁵ dil. of 267B-1
 un. odd = 10
 0.1 267B-1 lys = 53

Material against	269-1 ^{lys}	269-1 ^{lys}	269-1 ^{lys}	269-1 ^{lys}
on Bgal 1. 2297	dark, choct.	lys, solid film of contact, pink in water	supernat. of 269-1 contains 0	even control or
on Bgal 2. 750	c. 20 col remaining	no lys, contain 2+	no lys, contain 2+	de
on B(i) 3. 269-1	c. 4 dist. col.	no change	no change	str
4. 1412	c. 10000 col.	no lys, contain 0	no change	



1/9/54 Additional lysate made on in 1150, another in Pan, also using actinic glass tubes -
 Pan
 ↓
 2279 1/2
 811 1/2
 1177 1/2
 1/2 ← usual Lys⁺

Presumably the new phase is not in the prep or the new lysate. In addition, prep. appears not sterile

12/29/54

2341 x wild type HFT to Ashani $\left. \begin{matrix} 0 & \frac{5}{2} & 2^- \\ \frac{R}{2} & 2^- & \\ + & 2^+ & \end{matrix} \right\}$ and see segregants

- 2341 fresh culture P₁ overnight at end/line incubated in water bath, no air, for 3 hours, diluted to c. 10⁸ cells/ml
- 750 tK12 lysate = 5.8 x 10⁸ plaques/ml (see pg 284)
- 0.1 ml cell susp + 0.8 ml lysate incubated at 37C. for 10' 1.8 x 10⁸ trans/ml
- control (broth) exposed cells = $\frac{1.1 \times 10^6 \text{ cells}}{5.8 \times 10^8 \lambda}$ = $\frac{1100 \times 10^3}{5.8 \times 10^8}$ plates about the same no. of colonies, no (+) on any of them, no evidence of lytic activity.

5. sept. plates:
- 1/4 = 216, probably not significantly different in no. from control, evidence of lytic activity
no. (+) = 8
 - no. approx. same as control
no. (+) = 4
 - no approx. half of control
no. (+) = 6

6. purified twice		single seg		single seg					
stbl	λ	2279	λ	2279	stbl	λ	2279	λ	2279
1.	u	R	R	nl	13.	u	R	R	lyt
2.	u	R	R	nl	14.	u	R	R	nl
3.	u	R	R	nl	15.	u	R	R	lyt
4.	u	R	R	lyt	16.	u	R	R	lyt
5.	u	R	R	nl	17.	u	R	R	nl
6.	u	R	R	nl	18.	u	R	R	nl
7.	u	R	R	nl					
8.	u	R	R	nl					
9.	u	R	R	nl					
10.	u	R	R	nl					
11.	u	R	R	lyt					
12.	u	R	R	lyt					

found not segregating - It should be noted that segregants are derived from the same clone as the (+), not from the (-) shown

283-1 Cont. 304.
lysate #12 17/18/54 m,

no cell = 0
0.1 ml K12 = 175

	L ₁ ^S	L ₁ ^S	L ₁ ^S
292-3	27	9	6
170di 0.1	520	65	263
wt	493	56	257
0.1 ml	4930	560	2570

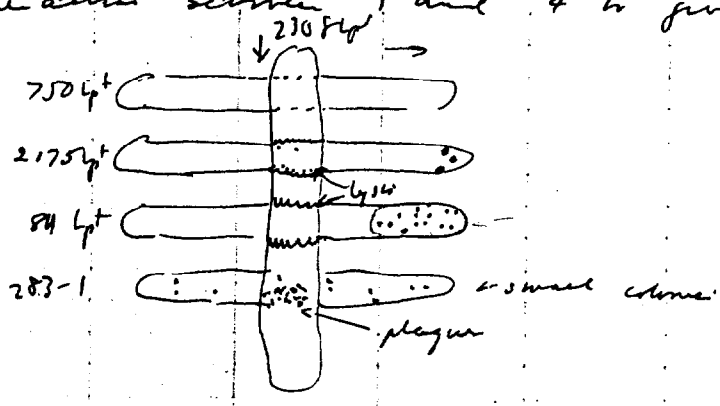
On to → 307

200 07. 304
/2279 c1300
/750 342
/1200 40
81 256

B

230 f

previous test of some 7 segments against 811 to see if they were HFT, suggested they were, but a lysate made from one of the segments was not. Do there a phenotype into which between 7 and 4 to give (+)?



202-16H Mal - One step - created culture from overnight pen. $10^9 \rightarrow 10^7 \rightarrow 10^5 \rightarrow 10^3$ moderate $\rightarrow 0.1 + 10$ ml pen incubator

Time	0.1 ml / 2279	0.1 ml / 750	0.1 ml plated	0.1 + 10 ml Pen	0.1 / 2279	0.1 / 750	0.1 / Plated
0	3	5 (1)	0		0 (3+)	24	20
15	4	5 (1)	-				
38	4	5 0	-				
45	4	5 1	-				
60	1	5 (19?)	-				
75	5 (8+)	7 1.75	-				
210	58 (trud)	14 10	Prod. = 35 pac.				

750 no. prod. = 3

3 of these were also marked as plaque.

Ant size	c. 16	c. 3	c. / 2279	/ 750	/ Plated
	plaque	trud. prod	0 (2)	30	59

10x	0	P prod.	T	C
obs	2:45 60	15 (6) 8	6	9
Med:	2:35 80	21 (4) 16	6	-
35"	2:55 100	33 (4) 8	10	-
	3:15 120	165 0 2	13	-
		145 (4) 6	13	-

Ant size c. 10 +
in the plates marked for plaque from top to bottom 3, 5, 2, 2, 0

1+7-
1-7+
← pinkie effect streaks

307-1A, 1C
1- x 7-
307

307A Continued.

Truss	Sagittatus	Sagittatus	Sagittatus	σ	all of these spotted	Re ✓	Genotype
1	"slightly"	-	+	1, 1, 7-	m	-	1-
2	"	parental	di-heterozygote	-	2125	survived	-
3	"	-	+	1, 1, 7-	m	+ - +	1-
4	"as usual"	parental	di-het.	-	5 gal	-	-
5	"	+	+	7-	and	+ -	7-
6	"	+	+	7-	wad.	+ -	7-
7	"	+	+	1, 1, 7-	none	- -	double
8	"	-	+	1, 1, 7-	found	- +	1-
9	"	-	+	1, 1, 7-	HFT	- -	double
10	"	-	+	1, 1, 7-		- +	1-
11	"	+	+	7-		+ -	7-
12	"	+	+	7-		+ -	7-
13	"	-	+	1, 1, 7-		- -	double
14	"	+	+	7-		+ -	7-
15	"	parental di-heterozygote	-	-		- -	-
16	"	+	+	7-		+ -	7-
17	"	+	+	7-		+ -	7-
18	"as usual" → stable (1)	-	-	-		- -	7 7-
19	"slightly" → stable (1)	-	-	-		- -	4 1-
20	"	-	-	-		- -	3 1-7-
21	"	-	-	-		- -	3 parents

307-1 →

all take 7- ave.
(primary)
page after 4 days.

~~7 7-
7 1, 1, 7-~~

4 stable 2

Continued from 305 - the 2341 x (+) $\frac{1-}{1-}$

W2854
= 307-1

- ① # 4, 5, 11, 12 re-treated for normal taking from "bush"
- ② single colonies (+) picked and tested (1), 2279, streaks out 2 gal

also for
derivatives
W2855

Stems primary	2279	△	Sagittatus
→ 4	lysoy	R	+
5	windy	Sens	no
11	lysoy	R	+
12	windy	R	+

307-4
-11
streak out 4, 11, all by present.

Stocks 1/4/55

307A

- K12
- 58-161
- W67
- W518
- W578 part #1
- W583
- W588
- W677

84 & HFT 7 (309-1) to obtain 4-7-
py. removed from a cross bush

1. 7 sep. sep. hand. 1/12 picked
nick sep. from each 3 7-
4 4-

2. Reheated to obtain 7 more sep.
6 obtained 3 7-
3 4-

74- (67)

← stock of this made 11-7-55
307A-1

W2792

1/9/54
ST?

Promin effect. between Gal₁ and Gal₆ (2070)

1. MAT pup - mixed with cells of 2070 with λ in tubes, incubated for 15 min at room temp., diluted and plated. Broth used as control \rightarrow c. 10⁷ and 10⁸

	(-) col	orange col. after 2 days
broth	45	0
	82	0
	<u>140</u>	<u>0</u>
CHFTI ⁺ plate	277	0
	73	1
	104	2
	<u>90</u>	<u>0</u>
	267	3

Stocks of the diheterozygotes 308-1, 308-2, 308-3
Stock del. +
discovered Sept 1, 1954

2. Segregants from sep. (+)

no.	orig. pup	orig. pup	tested adults
1.	o pup	o pup	populatus parental
2.	+	o	6-
3.	-	+	1-
4.	-	+	1- \leftarrow 308-4
5.	o pup	o pup	populatus parental
6.	+	o	6-
7.	+	o	6-
8.	o pup	o pup	populatus parental
9.	+	o	6-
10.	+	o	6-
11.	+	o	6-
12.	o	o	pup. parental
13.	o	o	1-6- same pup and strength
14.	o	o	1-6- \leftarrow 308-5
15.	+	o	6-
16.	o	o	pup. parental
17.	o	o	1-6-
18.	o	+	6-
19.	o	o	pup. parental
20.	5 stable Gal+	2	6 parental 8 6- 2 1- 3 1-6- how big pupae

W 2056

308-5

W 2057
also pupa derivat

+ -
- +
+ +
- -
+ +
+ -
+ +

1/9/54

Continuation of 302 -

302-1, 2, 3, 4, 5 - In search of HFT gals -

1. Since gal⁻ cultures were spotted on either 2 or 5 and irradiated give "apparent" false positive reactions for HFT, experiment was done on 5 gal using prototrophin 2175 gal⁻.
2. Above cultures streaked to individual (+) picked from each and streaked out to obtain 6 separate segregational events. The (-) obtained spotted on W2175 on 5 gal and irradiated. After 3 days result were

302-1	302-2	302-3	302-4	302-5	derived from 2307
no HFT	in HFT	in HFT	no HFT	1/6 appan HFT	309-1

1/26/54

To test h.h. Couette's hypothesis about crossing over between fragment and chromosome.

7. 309-1 plated on B⁺ gal, antibiotic to obtain reversion and to test stability

1. no reversions obtained B⁺ gal, several on B gal. B arab. result appears identical mixed culture (how can you see such colony arise). Streaking out of the growth on the plate suggests segregation for gal⁺ going on. Poss. of random reversion?
2. 2307 streaked against HFT 1, 2, 4, 7 on B arab.
3. 309-1 gal⁺ should be checked (after purification) on B arab. - Apparently 309-1 is ara⁺ as 2307 has been on v streak of 309-1 rather showing different colony type, do not show any (-)
4. Gal⁺ of 309-1 are unstable 1/6

Presumably

2-1-4+
2+1-4-

309B-1

48 x 1

2350 x 1 - See 272 For order of loci - Gatter additional information
24 x 1 - on segregants.

Seg	1-	2-	3-	4-	5-	6-	7-	8-	9-	10-	11-	12-	13-	14-	15-	16-	17-	18-	19-
1.	mixed with + ?	105 words del. of NA/prop	0	4B	7+ 5th/47	11-	12-	14-	Alleles										
2.	+	-	+	+	2-	21	+	+	2-										
3.	mixed with + ?	Uncolored			22	22	-	-	12-4-										
4.	+	-	*	+	2-	23	-	*	12-4-										
5.	+	-	*	+	2-	24	+	*	2-										
6.	-	-	*	-	12-4-	25	-	-	12-4- (New spring)										
7.	-	-	*	+	12-	26	-	+	1-										
8.	+	+	*	-	4-	27	+	-	2-										
9.	+	-	*	+	2-	28	-	*	12-4-										
10.	+	-	*	+	2-	29	-	+	1-										
11.	+	-	*	+	2-	30	+	-	2-										
12.	-	+	*	+	1-	31	-	+	1-										
13.	-	+	*	+	1-	32	-	+	1-										
14.	+	-	*	+	2-	33	-	+	1-										
15.	+	-	*	+	2-	34	-	+	1-										
16.	+	-	*	+	2-4-	35	+	-	2-										
17.	+	-	*	+	2-4-	36	+	-	2-										
18.	+	-	*	+	2-	37	+	-	4-										
19.	+	-	*	+	2-		+	-	4-										

W2859 also 1/2 den

309B-11
triple (-)
plus 31%

Classes	2-	17	27
(ideo)	2-	17	27
(allo)	1-	8	10
(ath)	14	0	0
Ampli	124	6	6
	12-	1	1
	4-	1	1
	35	45	

Previously
also 2-
also 1-

1/9/54

2433 x-4 For order of loci - see 272 and also other comb. on 309, but my

Seq	no add 2470-1	0	17	2/17	Recombination observed, 1 loc, 14 tailed	Comment
1	/1	/2	/4	+?	heavily populating after 3 days	
2	-	-	-	+?		
3	curtan e	(+)				
4	-	-	-	+ √	as ①	
5	-	-	-	+	as ①	
6	-	-	-	+	no pop.	
7	-	-	-	+	as ①	
8	-	-	-	+	as ①	
9	-	-	-	+	no pop.	
10	-	-	-	+ √	as ①	
11	-	-	-	+	as ①	
12	-	-	-	+	no pop.	
13	-	-	-	+ X	as ①	
14	-	-	-	+	no population	

A

Rpt to

also regarding on 2nd day
that some of the populating allele forms are h^s or h^m but
of something. h^s from
usually when thinned growth
restored on loci at h^s again time
 h^s in these clone? h^s h^m

On making purification before
next next loci (pop. forms
were selected)

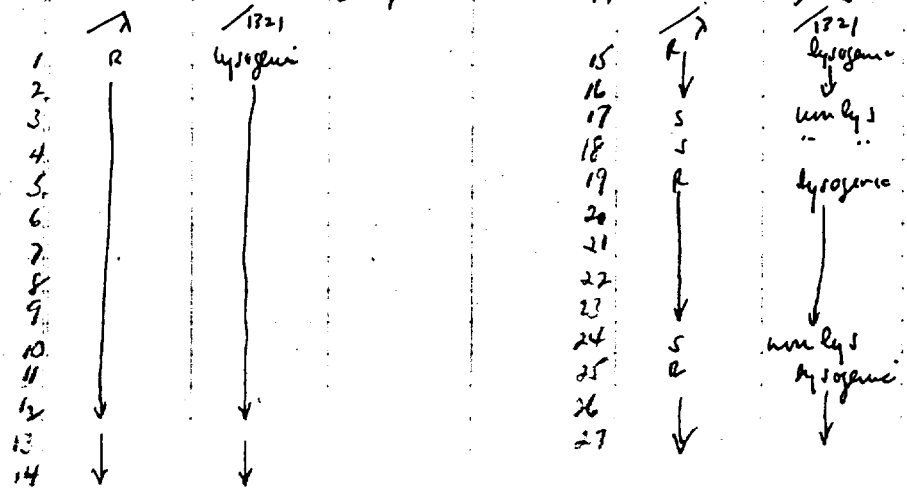
See if pop occur in 3 on B(0)
Also if this cult. auto, in region?

2308 x-2070 - to see if Gal_6^- is distinct from Gal_7^-

B

no lysate 2070 x(0.1) 30 (lysate)

27 pushed - all appear stable



2360 t K12 - to see incidence of stable (+) and lysogenicity
 trans. n_{gen} 304 2000

transl.	↑ R	1321 lysogenic	Sagittaria
1			+
2			+
3			+
4			-
5			+
6			-
7			-
8			+
9			-
10			+
11			-
12			-
13	S R	non lys lys	+
14			+
15			+
16			-
17			+
18			+
19			+
20			+
21			+
22			+
23			-

t = 420 total 8 non-lys
14 seg

2070 t⁺ - for making HFT 6⁻ onto 2175

311-10 (red gate) ① no add 11 97 - this titer higher than on 2308 t⁺, but again this may be indication of decay of lysates in standing (only measured on t⁺ cells) or a particular effect between 6⁻ and 7⁻

- 311-13
- ② pick and streak to obtain HFT 6⁻
 - ③ 13 seg titer, 2175 on steel, 2 possible HFT, #10, #12
 - ④ 17 " titer / 2175 Bgal, 1 HFT appears HFT

311-2

1/26/54

283-1 against 7⁻ - of the order is $\frac{1}{4}+$, then only double crossover
 since $g_{10} (+)$ #3 must be $\frac{1}{4} + \frac{1}{4} + \frac{1}{4} = \frac{3}{4}$

A

no. of = 0
 0.1 (1-1000) ¹⁰
 309-1 ~~1000~~
 (1/2) ¹⁰ ~~1000~~ ¹⁰ ~~1000~~
 1 single (+) → (-)
 3 c. 10 (-) & ¹⁰ ~~1000~~ ¹⁰ ~~1000~~
 4 +
 5 single (-)
 6 c. 10 (-)
 7 single (+)
 8 " "
 9 2 (-)
 10 +

1	4	7	Genotype
-	-	-	1-7-2-4-14?
-	-	-	-
-	-	-	2-7-1-7-2-4-14?
-	+	-	1-7-
-	+	-	1-7-
-	+	-	1-7-
-	-	-	1-4-7-2-4 (1-4)?
			3 1-7-
			3 (1-4)?
			3 gal(+)

283-1
1000

2070 arm brushed with HFT ~~1000~~ - Gal₆ destruct from Gal₇
 1 11
 2 11
 4 111
 7 1111

1/25/54

307-1 X-4- to obtain order $\frac{1-x}{1-x^2} = \frac{1-x}{(1-x)(1+x)} = \frac{1}{1+x}$

- ① The 4- (2472-1) by date deleted 1-100 gave c. 5×10^3 pop - Control (line E)
- ② 24 popules picked, checked, ③
- ③ (-) taken

1-4-5- where double crosses are required on order 1/2

	1	4	1
1	-	+	-
2	-	+	-
3	-	+	-
4	-	+	-
5	-	+	-
6	-	+	-
7	-	+	-
8	-	+	-
9	-	+	-
10	-	+	-
11	-	+	-
12	-	+	-
13	-	+	-
14	-	+	-
15	-	+	-
16	-	+	-
17	-	+	-
18	-	+	-
19	-	+	-
20	-	+	-
21	-	+	-
22	-	+	-
23	-	+	-
24	-	+	-

24 1-7-

The prospects appear the same as above, but the absence of the 1/3 should be noted. Other possibilities

DATE: 1/26/54

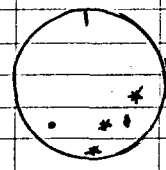
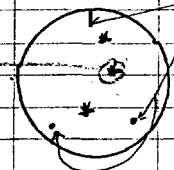
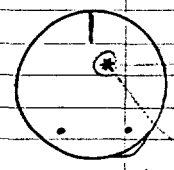
REF:

Repeat examination of Expt 301 - to see if hard particles - NFT
lysets are distributed "as mutant" in clone.

1. Cultures of 1210 753
2. Overnight culture of w1210² replicated to 3 petri and 13 fal plates and incubated c. 45 minutes. At this time the three plates were read. 20' with u.v. (10 cm). Reincubated about 30 minutes and then replicated to plate (previously spread E. coli, 30 min) and the plates incubated 3 days.
3. Confirming of the replica's aided by marking original 1210 plate with 3 streaks of w1210² (just) there. (1)
4. Result.

w1210-irradiated plate. (Bottom view)

w750 replica E 7210-1 plate



○ related fold

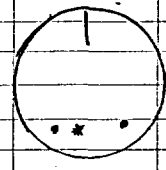
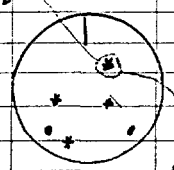
* get (f) pattern

all 3 found stable after (2)

all 3 found stable after (2) - on a waikiki

w1210-2

w750 rep.



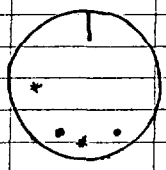
all 4 found stable (2)

found stable (2)

No suggestion of sp. colony on streak (1)

w1210-3

w750 rep.



17 found stable (2)

2 found stable (2)

When are the (+) clones from?

10

20

30

40

50

143 x 100 (100 x 100) 1.4x10⁵

DATE: 2/1/54 Inoc of 241-44 med - Q. J. Cole

REF:

1	2	3	4	5	6	7	8	9	10
Dose	spnt. = 3 ppp 2279 (10 ⁸)	spnt = 2 ppp 759 (10 ⁸)	spnt = 0 Plaque 2279 ↓ 10 ⁸ dil	spnt = 0 ppp 2779 10 ² dil	Exam of stabil/total	Populus Gul Sca	8 Kamda Kocher	9	10
0	28	67	155	1	1/1				
15	54	140	151	15	0/15	3 ⁺ mthly-14 ⁺	2 Lp ⁺ , 7 Lp ⁺ , 3 Lp ⁺		
30	78	184	172	22	17/18	r	- 14 Lp ⁺ , 2 Lp ⁺		
45	108	293	66	25	18/18	-	- 17 Lp ⁺ , 1 Lp ⁺		
60	176	434	28	22	17/18	r	- 16 Lp ⁺ , 1 Lp ⁺		
75	230	638	39	19	17/18	r	1 Lp ⁺ , 16 Lp ⁺ , -		
90	221	711	14	24	17/18	s: shal ⊗	- 17 Lp ⁺ , -		
105	237	536	17	37	18/18				
10	120	252	958	12	10 ² 31	17/18			
135	184	798	10 ¹ = 45	10 ¹ 201	18/18				
150	163	699	und: 305	und 247 (1/2)	18/18				
165	163	717	und: 231	ca. 4	18/18				
180	148	589	und = 141	ca. "	17/18				

↓
dil added

0 dose
1 ml = 155 x 10³ plaques
= 900 x 10² hand
1-10 dil per = 15500 plaques
9000 hand
1.6 x 10⁸ per ml
1.55 x 10⁵ x 10³ = 1.55 x 10⁸ ml
6.7 x 10⁶ hand/ml

10⁴ x 10³ = 1

probably
2279

Multiplicity Assay.

(1) 2279 x 0.1 ml 10² 0 dose plated. To 1/2 of plate 0.1 ml 759-2 & control (Per Plate) see also
 plaques on control (no 759-2) = 37
 populus on 759-2 half = 175
 populus on control = 2
 214 > than the highest value for plaque
 or hand. given above = 0, 15, 30 (155, 157, 172)

(2) 0 dose 10² dil
 (1) 0.5 ml + 0.5 ml cell (c. 10⁸) adsorbed 10'
 (2) add 0.5 ml 759-2 (= 1/3 dil of original), about 5'
 (3) plate 0.1 ml. no. populus = 108 x 1/2 = 282

DATE: 2/5/55

REF:

298-8 see last page and 304

1. 298-8 made lysogenic with λ from 84
2. To complete the study of order.
 - a. 298-8 X HFT 1
 - b. 298-8 X " 6
 - c. " " " 7

undelinked 0.1ml

3. Results. no plaques on any of the plates. What goes?

2279 against HFT 6, 7 to do heri recip. test of prot. effect.
0.1ml e. coli cells + 0.5ml lysate diluted 1:100 \rightarrow 5' incubated

1. 2279 control 3 plates c. 600 colonies/plate are (-)
2. 2279 X 6-

W2862
also had
derivative

Failed to harvest.
Overseed. Inoculate too
thick was very high here

317C-4

	2	7	ϕ	heri
1	0	+	2-	
2	0	+	2-	
3	0	+	2-	
4	0	+	2-	
5	0	+	2-	
6	0	0	2-	317C-6
7	0	+	2-	
8	0	+	2-	

8 additional
segments
from the same
transductions

2175 against HFT 6, HFT 7- to make double (-) for prophage gene under test

① Showy results between 6-, 7- and 2

② 12 colonies picked from early

	6-	7-	8 heri
1	+	0	1 cell
2	0	+	1 amp
3	0	+	
4	0	0	
5	0	+	
6	0	+	
7	0	+	
8	0	+	
9	0	+	
10	0	+	

	2	7	ϕ	heri
1	0	+	2-	7 heri
2	0	+	2-	2 cells
3	0	+	2-	
4	0	+	2-	
5	0	+	2-	
6	+	0	2-	
7	+	0	2-	
8	0	+	2-	

DATE: 2/8/55

REF:

Cross to see distribution of (-)

1. 578/902-1 x 2274
 all embryos ready to hatch at 578/902-1
 & cross plate

10 7505902-1 x 2274

1. control sh. (B gen), 2274 (-), 7505902-1 mixed (mult, +)

2. Poor cross plate - heavy bkgrd of small (-) colony, very shaggy
 c. 50 pult per plate

12 columns mixed - after ③ 2 reg. - All stable (+)

1	2	3	4	5	6	7	8	9	10
1	u	u	u	u	u	u	u	u	u
2	u	u	u	u	u	u	u	u	u
3	u	u	u	u	u	u	u	u	u
4	s	s	s	s	s	s	s	s	s
5	s	s	s	s	s	s	s	s	s
6	s	s	s	s	s	s	s	s	s
7	s	s	s	s	s	s	s	s	s
8	s	s	s	s	s	s	s	s	s
9	s	s	s	s	s	s	s	s	s
10	s	s	s	s	s	s	s	s	s

7. 902-1 x 2274

30

277-3 x 132-1

1. control sh (B gen) (132-1), 277-3 mixed (mult, +)

2. Poor cross plate - heavy bkgrd of (-) colony, very shaggy
 about 20-30 weath (+) per plate

12 (+) change plates - after ③ 10 stable, 1 segregating. Only 2 shaggy (+)

40

one of these segregating

1	2	3	4	5	6	7	8	9	10	11
1	u	u	u	u	u	u	u	u	u	u
2	s	s	s	s	s	s	s	s	s	s
3
4										
5										
6										
7										
8										
9										
10										
11	s	u	u	u	u	u	u	u	u	u

50

DATE: 2/10/55

REF:

On the order of the loci
308-5 = 1-6-

(A) a *Corynebacterium* (c. 0.01mc) of HFT 7 - treated on 1/2 plate P fal
control 1/2 = 0 (a faint bank forms of very minute pop)
HFT 7 lysate 1/2 = 31 (a single gal (-) HFT 7 phage gave you c. 10⁴)

The low titre here suggests the order 1-7-6 and that the pop are the result of double crossover.

B. The segregants. Of the 30⁺ examined, 18 were stable Gal⁻ at (C) RETESTS AFTER PURIFICATION

	1	6	7	6	7	6	7
1.	-	-	0	1-6-7-	0	0	(2pp) 0 * 18-?
2.	-	+	0	1-7-?	0	+	5pp * 1-
3.	parental pop. (-)	-	0	1-6-7-	0	0	0 * 1-6-7-
4.	-	-	+	1-6-7-	+	+	0 * 7-
5.	-	-	0	1-6-7-	0	+	0 * 1-7-
6.	- val +	-	0	-	0	0	2pp * 1-6-?
7.	-	-	+	1-6-7-	+	+	0 * 7-
8.	+ 617	+	-	7-	2pp 0	+	0 * 7-?
9.	-	-	-	1-6-7-	0	0	0 (1pp) * 1-6-?
10.	-	-	-	1-6-7-	0	0	0 (4pp) * 1-6-?
11.	-	-	+	1-6-	0	0	0 (2pp) * 1-6-?
12.					7 (1-6-7-)	amphiblytic	1-6-? = 5
13.					7 (1-7-?)	amphiblytic	1- = 1 amph
14.					1 (1-6-)	idios	1-7- = 1 amph
15.					1 (7-)	amph	7 = 3 amph
							1-6-7- = 1 amph

(B) a *Corynebacterium* (c. 0.01mc) of HFT 4 treated on 1/2 plate P fal

control 1/2 = 0
HFT 4 lysate 1/2 = 500-1000 - This high titre suggests the order 1-6-4 } the overall order appears now to be 2-1-7-6-4
and these pop. are the result of single crossovers

1. The segregants. - After (C) 6 of 24 stable, The segregating products in each case have only 1 or 2 (-) in their composition from atypical? Shunting of order is correct

	1	4	6	1-6-
1.	0	+	0	1-6-
2.	0	+	0	
3.	0	+	0	
4.	0	+	0	
5.	0	+	0	
6.	0	+	0	
7.	0	+	0	
8.	0	+	0	
9.	0	+	0	
10.	0	+	0	
11.	0	+	0	
12.	0	+	0	
13.	0	+	0	
14.	0	+	0	
15.	0	+	0	
16.	0	+	0	
17.	0	+	0	
18.	0	+	0	

Most of these have many small pop. in control and checker. remainder of 2433x-4)

DATE: 2/2/55

REF:

On the order of this loci

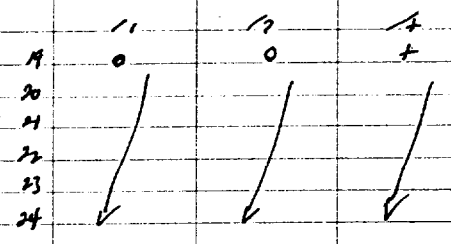
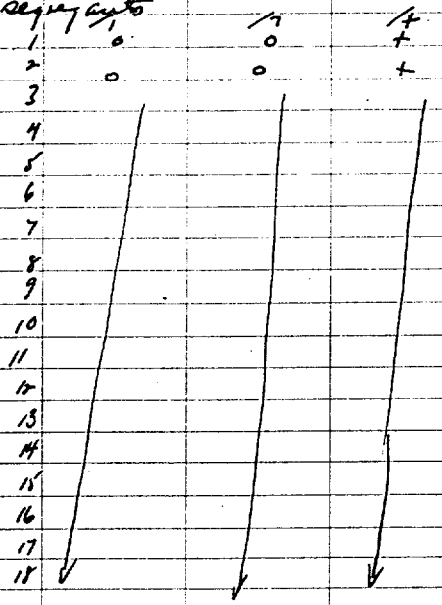
To complete the analysis of the positive aspect.
With regard to the array of data by date or other (-) see 305, 504

1. 307-1 x (+)
no odd = 0
O. line 1612 = 294
(11/12/54)

This finding ~~indicates~~ indicates the structure $\frac{11}{++}$
is of gel +

2. The segregants

A₁₀



24 177

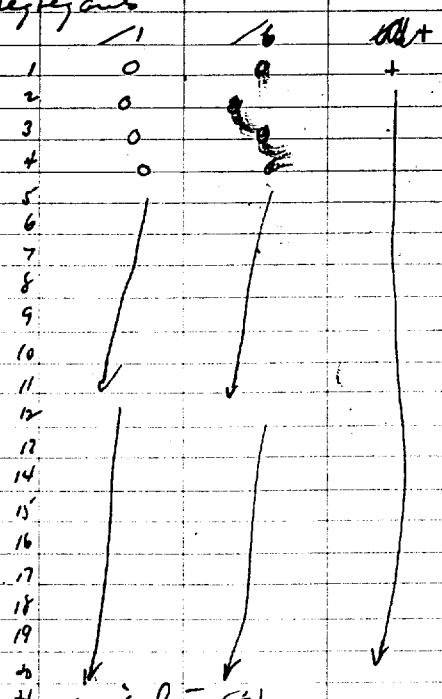
1. 308-5 x (+)
no odd = 0
O. line 1612 = 238
(11/12/54)

The structure $\frac{16}{++}$ is of gel (+)

3708-1
presumed
heterozygote
= 42876

2. The segregants

B



mixed (+)

20 16-

3 stable (+)

20

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40

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DATE: 2/17/57

REF:

Gal₁ - Gal₆ - interaction. Gal₂ - x - Gal₁ - Gal₆ - , is the product Gal₁?

W1210

no odd = 0
o.l.m. by 308.5 = 50

W1210 = gal₈

Seq

1

2

3

4

5

6

7

8

9

10

Genotype

1	+	0	+	+	1-6-
2	+	0	+	+	2-
3	0	+	0	+	1-6-
4	+	0	+	+	2-
5	0	+	0	+	1-6-
6	+	0	+	+	2-
7	+	0	+	+	1-6-
8	+	0	+	+	2-
9	+	0	+	+	1-6-
10	+	0	+	+	2-
11	+	0	+	+	1-6-
12	+	0	+	+	2-
13	0	+	0	+	1-6-
14	+	0	+	+	2-
15	0	+	0	+	1-6-
16	+	0	+	+	2-

<321-3 M-1-6-

W2864

17

18 }
19 } table(t)
20 }
21 }
22 }

12 2- idio
4 1-6- also
7 other(t)?

10

20

30

40

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DATE: 2/21/55

REF:

Multiplicity Effects

Reagent

1. 750 \pm λ (ad prep) $10^7 = 34 \ 41 = 3.8 \times 10^8$
2. cell Array = $10^7 = 172, 188 = 1.8 \times 10^9$
3. T- and. by rate = 10^7 dil of 0 area $241-14$ of 316, = 15500 plaques/ml

Procedure

10

	1	2	3	4
Ry 3 μ nd	1	2	3	4
Penney	0.5	0.5	-	-
Rx 2 cells	-	-	0.5	0.5
	0.5	0.5	0.5	0.5

Incubate 10
at room temp.

dilute as follows, 0.1 ml of above to

0.4 ml 750 λ	0.1	-	0.1
0.4 ml Pen	0.1	-	0.1

Incubate 5'

20

and then
plate 0.1 ml
+ 0.1 ml of Rx 2 cells

Plots: 246v

area, plot
and no of \odot

cf. area
partially lysed

Count

187 plaques	-	-
6 plaques	12 plaques	6 plaques
X 10 X 10	X 10 X 5 X 10	no plaques
18700 plaques	6000	0

30

40

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DATE: 2/4/55

REF:

Multiplicity Effects and also reciprocal case of protein effect between 1-6, 1-7

2279 deleted = 16¹ cell

		tube	1	2	3	4	5	6	7	8	9	10
	cells		+	+	+	+	+	+	+	+	+	+
	HFT 6-	0.5 ml	+	+	0	0	0	0	0	0	0	0
	HFT 7-	0.5 ml	0	0	⊕	⊕	⊕	⊕	⊕	⊕	⊕	⊕
10	admt. 10 min											
	del 10 ²	(45 r. 323)	0	+	0	+	0	+	0	+	0	+
	cell 0.1 ml + 0.9 ml	753.2	+	0	+	0	+	0	+	0	+	0
	cell 0.1 ml + 0.9 ml	penicillin	+	0	+	0	+	0	+	0	+	0
	admt 5 min											
	plate 0.1 ml											

Plasma:

24hr	col. (1), slight	(-) col. del	1-1-1	1-1-1	(-1) col
	condensed	partially gtd	323	323	236 (-) col.
26hr	256(-) col	287(-) col	1 col. pp. dptg	1 col. pp.	236 (-) col.
			246 (-) col. on	306 (-) col	
			partially gtd.		
48hr	1 col. E 2	1 col. E 5	no further	no further	
	cond. pp. ⊕	cond. pp. ⊕	change	change	

Stocked +
323-2
successful
Sept 1, 1964
Contract firm's
revised

Shaded out parent (-) picked
in each case to strain (+)

323-3 323-4 ← Stocked +
successful
Sept 1, 1964

No (+) obtained

22

23

323-4 = 1-6-6 for protein effect analysis
1 Segments 1/22 +ⁿ

323-2 = 1-6-6 as above
1 segment 1/23 +ⁿ

6 Segments

	1	6	tr β	Change
1	0	+	sew	1-
2	+	0	sew	6- 323-5
3	+	0	sew	6- 323-6
4	0	+	sew	1- 323-7
5	+	0	sew	6-
6	+	0	sew	6-

4 seg.

	1	2
1	+	0
2	+	0
3	+	0
4	0	+
	all 1-2	

For further
qualitative seg
7-
7-
7-
1-
see 393B

W2894

W2895

6- mutations
both stable gtd

g. 6- gtd mutations,
1 unstable

W2791

DATE: 2/27/55

REF:

324

10
 Is 1210 really gal⁻ or a closely linked (-) In the past, particularly with W2350 and W2760, Gal⁻ Gal⁻ derived from W1210 it has been noted that tests against HFT2 (and HFT1 lysate are derived from the (-) of W902) gave (delayed) positive response few in number. Purification of the culture did not always show this effect. It may be (despite the crossing data (11,000 prototrophs) between W18 x 902 that 1210 gal⁻ is not a recurrence of gal⁻. To add further information 3 separate gal⁻ HFT lysate derived from (W902 gal⁻), known to have originated from separate single gal⁻ HFT colonies were put onto 1210

	lysate	1210
2342	3/17/54 lys	c 50 pop after 3 days
2342	3/20/54 lys
	241-14 lysate

20

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Mal-4⁺ breakthrough - also selection for a mutant - Serial selection
 296-1 gal⁻ neg : 296-1 = HFT 2342
~~HFT~~ (not purified) exposed to 0.1 ml 2342 (2/17/54) for these.

2nd add = 23
 0.1 ml HFT2 = 78 x 8 = 624 slightly higher than previously but this is probably a different lysate

40

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DATE:

REF:

325

Observations on Erotia Induction

1895^{HT} 1 pt X 2234 F- 4p^s

- Freshly grown cultures, by diluting overnight culture
- 5.0 + 5.0 ml

Pre mix

Port - Mix

0.5 ml samples

	1895	2234	Plaque	Col	Plaque	Col	Plaque	Col	Plaque	Col
10	2	72	0	23	17	17	17	17	5	23
	5.1 ml		0	0	1					
	2234	0	99	19	23	23	23	23	17	17
		dit								
		= 1/4								

Suggestive -

2nd run.

Dilutions the same throughout

	1895	2234	Plaque	Col	Plaque	Col	Plaque	Col	Plaque	Col
20	3	6	0	0	11	18	19	24	21	31
	2234									
	1895									

③ After 9th culture diluted and plated a 8 gal to one of and plate one by 2. 3 plates

Not suggestive

replication of this plate shows 2-10 protospores.

1, 2, 3 about same

On streaking out some of these protospores, 28(-), 2(+), most of the (+) were greatly contaminated with (H - the original source of) since the culture was diluted before plating and there should have been few (H) protospores. The cross plates and the protospores were discarded - R to do -

30

1210 & 84 to make HET of (210 (-))

1. no add 2
 PA (old) 22

40

2373 - to run stability of Kaid.

1. no add 8
 o. l. no K12 250
 (7/29/54)

d. as control the same / 2279

2279
 no add 0 } many small
 o. l. no K12 281 } protospores
 (7/29/54)

50

DATE: 3/5/55

REF:

307-1 10^5 (mode E UV) = gal⁻, gal⁺. To determine order and base of
 reaction is operating against certain ² should show
 1. Transduced with both 4⁻, 6⁻ (HFT) = 2478-1 for 4, 311-2 for 6-

	1	2	3	4	5	6	7	8	9	10
one night out	A. Control not exposed to h.				(-)	(+)	population			
↓	exposed to both			plate	1	79	0	0		
↓					2	77	0	0		
↓					3	87	0	0		
[0.2 and 0.2 ml HFT]										
↓	B. Exposed to HFT 4				1	47	4	0		
↓					2	48	5	1		
↓					3	69	0	0		
↓						164				
↓	C. Exposed to HFT 6-				1	32	0	0		
↓					2	23	0	0		
↓					3	37	1	0		

[Handwritten signature]

306C -
 Analysis of 332

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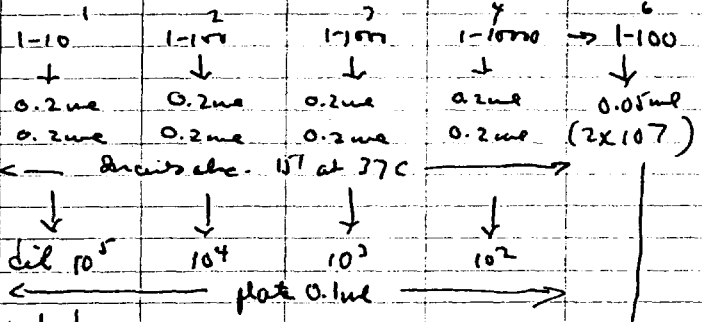
DATE: 3/18/53

REF:

2344 MI - at J.L. Request to test multiplicity of food and to get information for run.

1. Cerealid 2344 → dil. from overnight

2346A (2/17/54) →



Id. 1x10⁷

not done, no plates available

Result

no. colonies in plate(s)	-	46 47	37 38	35 30	134
	-	3 7	2 2	2 2	120
no cells / tubes / 1/2 HFT. / tube	-	2.54x10 ⁷	2.54x10 ⁶	2.54x10 ⁵	2.54x10 ⁹
No. survivors	-	4.7x10 ⁶	3.8x10 ⁵	3.3x10 ⁴	
Frac. surviving	-	0.36	0.29	0.25	
Frac. survivors	-	0.11	0.053	0.061	
Frac. transd.	-	0.04	0.015	0.015	

Plasm. found. 328-118

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DATE: 3/12/55

REF:

2344 MI X — 2346 A 3/17/54 Repeat to see frequency of hazard.

Quoted 2344 MI

↓

1-100 →

Contra

0.3

EPA

0.3

Birth

0.3

2346 X

Admits

15 min

37c

Al

104

10²

↓
0.05 ml = plated →

0.05 ml

Colony:

Plates

1	2
---	---

(+)

0	0
---	---

(-)

43	26
----	----

total

43	26
----	----

35

1	2
---	---

9	2
---	---

229	199
-----	-----

229	199
-----	-----

184/428 = 4.2% (+)

10

20

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DATE: 3/14/55

REF:

Repeat analysis of 285-2 - 24 pgs. (-) picked and checked to obtain a page
 gdx (+) from each one.

	1	2	3	4	5	6	7	8	9	10
	Gas +	Stability ①	Stability ②	i	+	LP ^{RK}				
	1	u	✓	parental (-)						This is wrong cell - as LP ^{RK} is not LP ^{RK} should have had 285-1
	2	u	✓	0	+	ans 1-				
	3	s	1							
	4	u	✓	0	+	ans 1-				
	5	s	2							
10	6	s	3							
	7	u	✓	0	+	ans 1-				
	8	s	4							
	9	s	5							
	10	u	✓	0	+	ans 1-				
	11	s	6							
	12	s	7							
	13	s	8							
	14	u	✓	0	+	ans 1-				
	15	u	✓	0	+	ans 1-				
20	16	s	9							
	17	s	10							
	18	s	11							
	19	s	12							
	20	s	13							
	21	s	14							
	22	u	✓	+	0	ans 4-				
	23	u	✓	- parental (-)						
	24	14 s	13 s			6 1-				
		8 u	9 u			1 4-				

30

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DATE: 3/18/55

REF:

1321 X 902 FT - to obtain improved cross data. Previously 518 X 902

1. Control streaks ok on 8 gal
2. Cross plates bad - many small (+), ca. 20 large col., probably (+)
 Don't know what to make of this? Entire undant..?

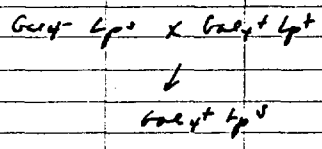
10

518 X 2035 to look for "unstable" recombinant class - namely gal+ Lp^s
 5 gal.

1. Over plate ok. ca. 90% gal+ - no record of control plate
2. 50 (+) col. picked streaked out B₉ gal, same suspension also streaked

amphib streak #	against 2 gal. streaks 40 (-) in	Tests of the (+) on the streak
# 3	1/ca. 50	sem. sem.
# 5	0/100	sem. sem.
# 20	0/ca. 200	sem. sem.
# 41	5/ca. 100	sem. sem.

4/50+ are recombinants between Lp and val



replied to S gal
 all perturbation

20

296-1 - 1412 E AFT²

30

1. Hoping to find 1 best range mutant. Repeated transd. of gal- seg from this heterozygote

1. transd. with gal+ HET 1 → gal- obtained → Gal+ HET → apt. #3
 #1 #2

from #3 8 trans. picked and single gal- seg obtained. Plated with 0.1 ml MFT gal+ 1 to see if phage received in USA.

40

1. On one (#3) single faint plaque (?) noted.
2. on one (#6) severe " plaque (?) observed near edge

observed after the 4th passage.

50

334

DATE: 3/18/55

REF:

W2790

1	2	3	4	5	6	7	8	9	10
---	---	---	---	---	---	---	---	---	----

W2580 = gal - the 1/2^s - modeled and by skinned

↑
W2580
↳

10

20

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DATE: 3/20/55

REF:

	1	2	3	4	5	6	7	8	9	10	
	Lysate						9/20/55				
	811 t 902		algae	Transduction's (H ₂ O)		H ₂ O = 2x10 ⁴				lysozyme	
		Plaque (10 ⁵)	filter	901	2580	2x10 ⁵	H ₂ O / 811	2580	2580	5.0	
	335-1	1. 110, 116	2.3 x 10 ⁷	34	30	19 = 3.8 x 10 ⁶	22	20	++	+	
	2	2. 57	1.5 x 10 ⁶	26	20	9 = 1.8 x 10 ⁶	24	17	0	0	
	3	3. 129, 134	2.6 x 10 ⁷	20	23	12 = 2.4 x 10 ⁶	21	25	±	++	
	4	4. 148, 158	3.1 x 10 ⁷	19	26	15 = 3.0 x 10 ⁶	41	23	++	+	
10	2580 t 811			2x10 ⁵		H ₂ O = 2x10 ⁵					
	5	1. Plaque (10 ⁵)		115	16	when	13	18	0	0	
	6	2. 229 = 2.3 x 10 ⁷			33		1033	78	± 2x10 ⁶	++ 4.4 x 10 ⁶	
	7	3. 211 = 2.1 x 10 ⁷		69	78		28	96	1.2 x 10 ⁶	1.4 x 10 ⁷	
	8	4.		35	32		12	16	0	0	
	Squibb			29	11		22	26			
20	Better data from Peatich										
	2236 for HPT 3-										
30	1. no add = 2		when these were tested, initially few percentage unstable (12/12? or 17/14?)								
	2. 902 λ ⁺ = 342		plate stored in fridge c. month, then 24/14, 17/12 found stable								
			why? see next page also/511								
	2062	1. unadd	40								
		2. 902 λ ⁺ =	85	81							
40	2580 against PH λ		6/1/54	Rpt 9/28/55							
	1.	no add	11	9							
	2.	0.1 ml λ	140	170							
50	2580 λ ⁺ against PH λ		6/1/54								
	1.	no add	19								
	2.	0.1 ml λ	4 x 906 = 3994								

DATE:

REF:

	1	2	3	4	5	6	7	8	9	10
	327C-1 = 1-6-7-7 - From 24 exp. presumed parental (-), checked out - so actually were. From each parent (+) a couple + obtained.									
	C.O. (1)	stability (1)	(2)	1/1	1/6	1/1				
	1	o	o	_____						
	2	o	o	_____						
	3	o	o	_____						
	4	o	o	_____						
10	5	o	o	_____						
	6	o	o	_____						
	7	o	o	_____						
	8	u	u	o	o	2 exp				
	9	u	u	o	o	1 exp				
	10	u	u	o	o	o				
	11	u	u	o	o	o				
	12	u	u	o	o	o				
	13	u	u	o	o	1 exp				
	14	u	u	o	o	3 exp				
20	15	u	u	o	o	o				
	16	u	u	o	o	2 exp				
	17	u	u	parental (+)		o				
	18	u	u	o	+	6 exp				
	19	u	u	o	o	1 exp				
	20	u	u	o	o	1 exp				

This culture presumably
48h
4h

all exp
except
parental (-)

6 - 1
7 - 1
1 - 1

134
7 A.

30

SA by 902 A *

40	1. no add	29	
	2. 0.1 me A	88	60

SA by 902 A

50	1. no add	45	
	2. 0.1 me A	234	185

DATE:

REF:

337

Linearity of HFT at high dilution

7SD v 1-10 dil of 241-14 (0 dose - rad. opt.) ^{see 316}

6/22/55 1-10² dil
Plates exposed W2792

1	2	3	4	5	6	7	8	9	10
	me dish	page	Δ						
	0	1	0	10.1			0	0	0
	0.02	16	14	70			0.02	90	
	0.04	38	36	88			0.04	183	
	0.06	68	66	110			0.06	257	
	0.08	74	72	90			0.08	354	
10	0.10	95	93	92			0.10	459	

2279

undil of 241-14 (0 dose rad. opt.) ^{see 316}

Plt. 418155

Plt. III

Plt. II

see plate, back @ 37
total plate addition
made up to 0.2 with
buffer

1	2	3	4	5	6	7	8	9	10
	me by side	1-2	Δ						
	0	1	0	4	3	0		0	2
20	0.02	28	27	total percentage of cells = 35	15000/0.1ml		30	37	35
	0.04	61	60		66	63	37	73	71
	0.06	93	92		140	137	68	145	143
	0.08	201	200		154	151	90	135	138
	0.1	316	315		157	156	238	158	157
				0.12	175	172	312		
				0.15	287	284	(0.14) 2350		

30

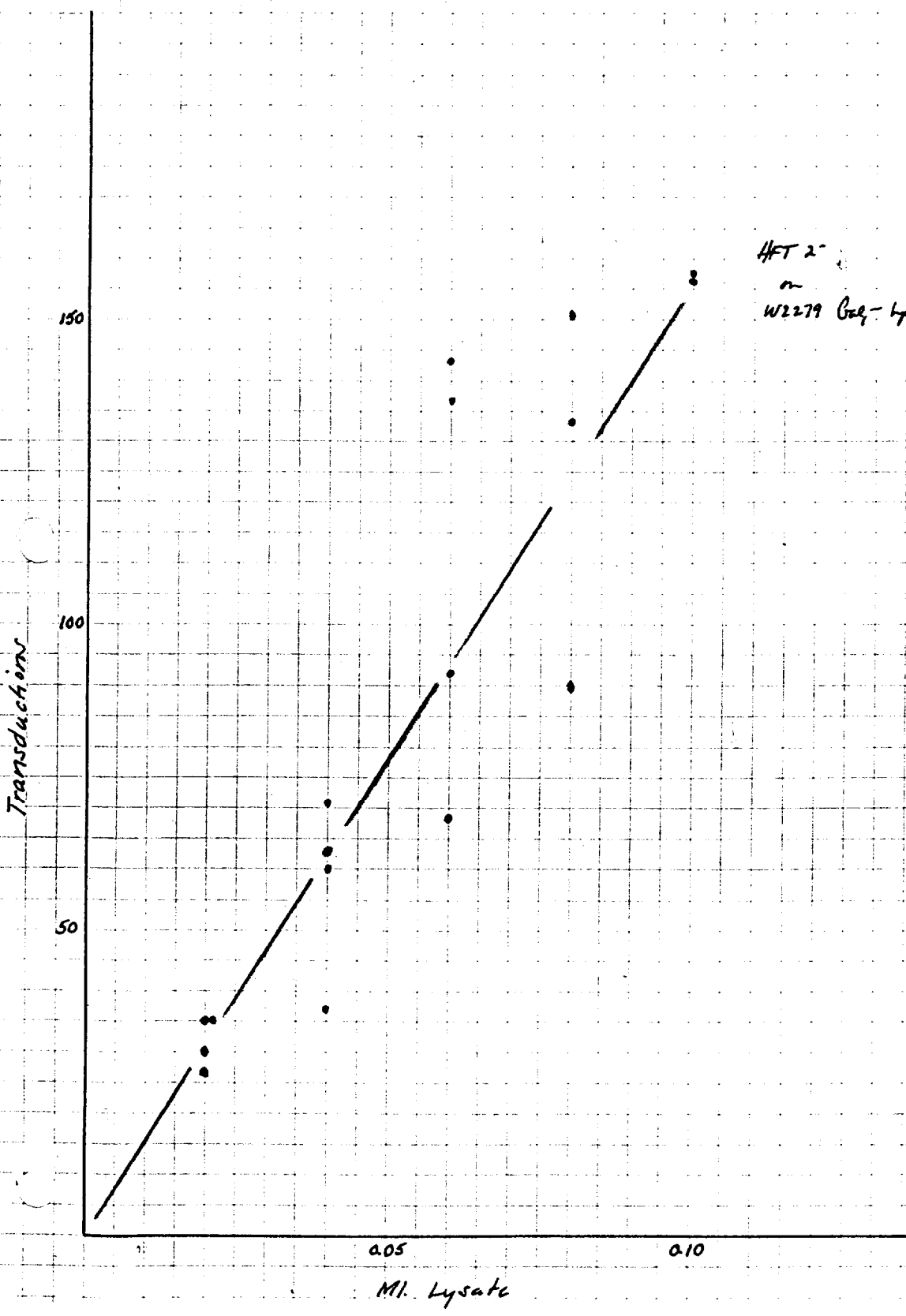
40 1402 X 902F+

7. units on B gel etc.

2.	(-)	(+)		
Platograph	521	1		
①	512	1		
②	603	1		
③	466	2	3771	$\frac{0.0018}{7.000}$
④	538	0		
⑤	575	0		
⑥	556	2		
	3771	7		3771 322.90

= 0.18%

50



DATE: 3/23/55

REF:

Linearity with NPT.

1. 84 overnight cure, wt conc. lysate = K12 11/12/54

	lysate	lipidase	Δ	
	0	23	0	-
	0.02	63	40	20
	0.04	112	89	22
	0.06	151	128	21
	0.08	plate contaminated		
10	0.10	234	211	21

2. 2580

	0	22	0	K12	11/12/54
	0.02	163	141	70	
	0.04	270	248	62	
	0.06	345	323	55	
20	0.08	460	438	55	
	0.10	527	525	53	

3. 2580¹

	0	10	0	K12	11/12/54
	0.02	1467			
	0.04				
30	0.06				
	0.08				
	0.10				

528

	0	49	0		
	0.02	499	450		
	0.04	714	665	186	} plate contaminated
40	0.06	888	839	139	
	0.08				
	0.10				

50

DATE: 4/8/55

REF:

1437 to see if Matt 4×2 (A-2 rest) are also bound. by HFT since the bound - type may all be temperature dependent (etc).
 checked out on B used - mixed.
 1. 4 (+) } checked / A-2, melt = A-2 rest.
 2. 4 (-) } melt = A-2 rest.

1439 spread with A-2 to obtain Matt A-2 rest.

20 Tests of some HFT cultures (2-) obtained on 293. Status checked on B gel - all pure (-). Columns prepared and tested / HFT for HFT using u.v.
 (+) = HFT, O = NPT

293-1A.

○	○	⊕	⊕
⊕	⊕	⊕	○
○	○	⊕	⊕

 O may be the result of too few cells
 HFT 1/4 gel +^a aggregating at ②

293-2A

⊕	○	⊕	⊕
⊕	⊕	⊕	⊕
⊕	○	○	⊕

 O = too small inoculum (no growth on spot plate)
 HFT 2/3 gel +^a aggregating at ②

293-2B

⊕	⊕	⊕	⊕
⊕	⊕	⊕	⊕
⊕	⊕	○	⊕

 HFT 1/2 1/4 1/4 2-
 O = good inoculum
 HFT 2/3 gel +^a aggregating at ②

293-11A

⊕	⊕	⊕	⊕
⊕	○	⊕	⊕
⊕	⊕	⊕	⊕

 O = good inoculum
 HFT 3/4 gel +^a aggregating at ②

1A

⊕	⊕	⊕	⊕
⊕	⊕	⊕	⊕
⊕	⊕	⊕	⊕

 ⊕ good inoculum 1 2 1 1 2-
 2 3 1 1 2-

2A

⊕	○	⊕	⊕
⊕	⊕	○	⊕
⊕	⊕	⊕	⊕

 O good inoculum 2 0 2 1 2-
 0 0 2 1 2-

DATE: 4/13/55

REF:

1	2	3	4	5	6	7	8	9	10
1736 - originally (see pg 137, 113) 1736 C/12 gave -1 or weak (+) result.									
- suspicion that something unusual here. Re-examine									

①	incub	c. 40
	L12)	c. 1000

②	Streaks.	12/12	strong	fact,	organizing
---	----------	-------	--------	-------	------------

10

20

2799 =	malt λ -2 ^F 2279	no	break through:
--------	-------------------------------------	----	----------------

Disruptive λ 4^R

1. 10 add = 3

2. 0.1ml

2342 3/17/54 = 4

30

241-14 mal -	Spontaneously produced λ - HFT?
--------------	---

1750

1. no add (P) 6

2. 10² (0.1ml) 41

3. 10³ (0.1ml) 3

35

$35 \times 10 \times 10^2 = 35 \times 10^4$

opt.

40

2279	10 ² (0.1ml)	0 plaque
		5 plaque

50

DATE: 4/20/55

REF:

2580 for checking a diploidy of the markers after crossing over a *ginc aestivus* HFT plant (-)

341C-control
341/4
341E/2

1. origin of markers in 335
2. a second plate made see p 335
3. to aid in understanding phenotype of their cultures on arabidopsis comparison made between K12 +

17 cag clones obtained. ^{Spotted on} W995 (100%) int. -
^{W945} int. -
^{W2570} int. -

Previously diploids for having dist are followed by 2580 after 2 days peculiar phenotype of this cross. ^{W995}

10

Summary

⊕ F11	⊙ ⊙ ⊙ ⊙	⊙ F11	⊙ ⊙ ⊙ ⊙	⊙	⊙ ⊙ ⊙ ⊙	⊙	⊙ ⊙ ⊙ ⊙
341-9	→ ⊕ ⊙ ⊙ ⊙	HFT 2	⊙ ⊙ ⊙ ⊙	some of 0 have control pop	⊙ ⊙ ⊙ ⊙	most of these have c. 20	341-9 rechecked on
⊙ 2580	⊙ ⊙ ⊙ ⊙		⊙ ⊙ ⊙ ⊙	341-12	⊙ ⊙ ⊙ ⊙	specific WHY?	Bgal 4/6 rechecked 4
⊕ = HFT	⊙ ⊙ ⊙ ⊙		⊙ ⊙ ⊙ ⊙	(Phenotypic all before tested)	⊙ ⊙ ⊙ ⊙		Bgal 4/6 " "
⊙ = contaminated	⊙ ⊙ ⊙ ⊙		⊙ ⊙ ⊙ ⊙		⊙ ⊙ ⊙ ⊙		Bana 4/6 " "
⊙ = HFT	⊙ ⊙ ⊙ ⊙		⊙ ⊙ ⊙ ⊙		⊙ ⊙ ⊙ ⊙		Bana 4/6 " "

21/4/56
LFT seq of 341-9
P/S 2580 -
5/5 genⁿ of LFT of HFT

20

2062 - see 335 for details.

30

1. check of HFT derivatives for diploidy for protein
2. 5 separate hand-picked segregants tested

Segregants from hand #

	1	2	3	4	5
	F11	F11	F11	F11	F11
	2580	2580	2580	2580	2580
	⊙	⊙	⊙	⊙	⊙
	⊙	⊙	⊙	⊙	⊙
	⊙	⊙	⊙	⊙	⊙
	⊙	⊙	⊙	⊙	⊙
	⊙	⊙	⊙	⊙	⊙
Expt. 1	⊙	⊙	⊙	⊙	⊙
	⊙	⊙	⊙	⊙	⊙
	⊙	⊙	⊙	⊙	⊙
	⊙	⊙	⊙	⊙	⊙
Expt. 2	⊙	⊙	⊙	⊙	⊙

ALL HFT, get things on 2580 plates 10-20 pop. Do this an indicator how are 4-?

as above

40

50

DATE: 4/30/55

REF:

342

1. Are there position effects between 4, 6, 7?
2. The direction.

A. Overnight broth culture diluted 1:100 in 0.1 ml cells + 0.1 ml lysate → adsorb 10' at 4°C
 Dilute 1-100, 0.5 + 10 ml, and plate 0.7 ml on 2 B gal plates.

③ lysate of
 247B-1 = 4
 311-2 = 6
 309-1 = 7

③ -x Gal₇ - 2308?
 20 hrs

10

		(-) col	(+) "Hpt"
A. broth		302, 260	0
HPT 4 ⁻ B. 247B-1 A		192, 198	6, 5
HPT 6 ⁻ C. 311-2 A		254, 169	5, 4

same as above (+) were + in 24 hrs.

phenotype

The phenotype here

may be weaker (+)

popul. (-) ?

342-A1 → X

* 342-B1 → X 4

④ -x Gal₆ - 2070

20

A. Broth		483, 431	0
B. HPT 4 ⁻		256, 251	4, 2
C. HPT 7 ⁻		283, 332	3, 3

pop (-) ? * 342C1 → X C 4

pop (+) ? * 342D1 → X D3

⑤ -x Gal₄ - 518

30

A. Broth		108, 124	0
B. HPT 6 ⁻		199, 99	2, 1
C. HPT 7 ⁻		85, 105	0, 1

pop (-) ? * 342E1 - R E2

intermed (+) ? 342F1 -

* on 2nd streak gave papulating colonies.

40

50

DATE: 5/11/55

REF:

HPT 7 - X 6- 342 D - 07 24 P.E. (-) picked for (+), 4 were true (-), 20 P.E. (-)

Procedure on 343

P.E. #1	(+) W. reg?	Seq. reg. 1/6	Seq. reg. 1/7	1/4		
2	+	7	0 ✓	+	+	6-
3	+	30				marked
4	+	3	0 ✓	0 ✓	+	6-7
5	0	-				
6	+	12	0 ✓	0 ✓	+	6-7- ← 344-6
7	+	7	0	+	+	6-
8	+	4				P.E. (-)?
9	0	-				
10	+	4				P.E. (-)?
11	0	-				
12	+	9	0	+	+	P.E. (-)?
13	0	-				
14	0	-				
15	0	-				
16	+	4	0	+	+	6-
17	+	5	0	+	+	6-
18	+	730	0	+	+	6-
19	+	750				
20	+	10				P.E. (-)?

13 seq - one lost
7 more

5 6-
+ 6-7
4 P.E. (-)?

(A) 344-6 above, the 6-7, K12A 7/27/54

no. add = 0
o. line = 44

see p. 325 for previous use of this code

Seq.	1/6	1/7	1/4	Handtype
1	0	0	+	6-7
2	0	0	+	
3	0	0	+	
4	0	0	+	
5	0	0	+	
6	0	0	+	
7	0	0	+	
8	0	0	+	
9	0	0	+	
10	0	0	+	
11	0	0	+	
12	0	0	+	
13	0	0	+	
14	0	0	+	
15	0	0	+	

(B) 344-6 - X 2580

no. add = 10
o. line = 215

Seq.	1/6	1/7	1/4	Handtype
1	0	+	+	2-
2	0	+	+	2-
3	0	+	+	2-
4	0	+	+	2-
5	0	+	+	2-
6	0	+	+	2-
7	0	+	+	2-
8	0	+	+	2-
9	0	+	+	2-
10	0	+	+	2-
11	0	+	+	2-
12	0	+	+	2-
13	0	+	+	2-
14	0	+	+	2-
15	0	+	+	2-
16	0	+	+	2-
17	0	+	+	2-
18	0	+	+	2-
19	0	+	+	2-
20	0	+	+	2-

50

DATE: 6/20/55

REF:

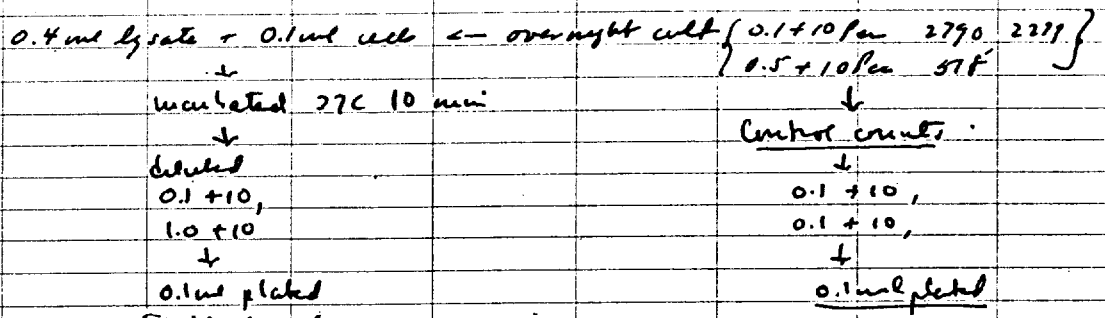
1. Testing for re-infection of *Sp* in isolated heterocysts.
Also effect of temperature shift on frequency of transduction.

A. Overnight cultures 2279, 2790, 518 at 37C in incubator.
4 cultures divided and 'pulses' placed at room temp for ca. half hour.
PM temp cells, 37C cells.

B. Lysates
Gels - 1 for 2279 518 = 2342 3/20/54
Gels - 1 for 2790 = 2346

a. Lysate 0.4 ml ~~placed~~ in small tubes, caps off incubated at 37C for half hour.

C. Technique



37C	Culture	Half	fraction (1)	3	Culture	Control	Counts	37C
			2			RM temp		
	W518	1/443	3/460	0/428	no more ch. also plated.	W518 =	112,103 (1.1 x 10 ⁷)	119,130 (1.2 x 10 ⁷)
30	W2790	1/460	1/372	1/422	← also culture with plaque	W2790	35,42 (3.9 x 10 ⁶)	173,184 (1.8 x 10 ⁷)
	W2279	8/173	1/166	5/192	← just about every colony has plaque - multiplicity too high.	W2279	180,144 (1.6 x 10 ⁷)	338,290 (3.1 x 10 ⁷)

No gels on these dates

RM temp.	Aut.	Plate 1	fraction (1)	2	3
40	W518	2/ ca. 50	3	4	
	W2790	2/263	1/292	0/262	
	W2279	1/55	3/56	0/50	

← these plates have contamination - no count of trial made.

← these colonies all are plated - mult. too high

The multiplicities in this experiment were too high to be used for the purpose of testing of *Sp*.

50

DATE: 6/27/55

REF:

7 Observations on segregation rate

- A. Culture = 348 edis type
- B. Streaked out into single pure col. inoculated into Pan.
- C. Diluted $10^2 - 10^4 - 10^6 - 10^8$

10
 before 60 min. by.
 this way 3 min.
 from 7:15 AM to 7:30 AM
 2nd - leave
 until 1:00 PM
 2 = (256x)

5 0.1 ml samples
 plated B. gal

1. 100	} therefore ave. of 1.2 cells/plate = 3.6 cells/plate
2. 108	
3. 134	
4. 136	
5. 108	
<u>586 = 117</u>	1.05 ml 0.6 cell

10. 0.3 ml samples incubated 37C 8:30 AM
 Plated 0.05 ml of 1:00 PM

1. 361
2. 61
3. 219
4. 122
5. 855
6. 1279
7. 471
8. 676
9. 319
10. 336

20 6/27/55

- 2. Repeat experiment using 780B12-1
- A. Othrylet culture from stool.
- B. diluted $10^2 - 10^4 - 10^6 - 10^8$

array 0.05 ml

1. 98, (2)
2. 112, (5)
3. 106, (4)
ave 106 (4-)

212 (2-)
100

5 0.1 ml samples incubated in horizontal 0.2 ml pipette at 37C. 9:15 AM
 20 in number 1:45 PM
 Placed (and inoculated)

2 cells/0.1 incubation, 0.08 Gal-

30

①

Sample	(-)	total
incubated 1	*	*
in	2	6
pipette	3	3
	4	4
	5	0
	6	* = *
	7	23
	8	* 103

② At the same time 5 0.1 ml samples plated in deep form on B. gal, resuspended after plating ca. 2:15 PM by adding 0.1 ml broth & spreading

	(-)	(+)
1.	0	89
2.	0	52
3.	7	1295
4.	22	1222
5.	52	730

40

7	319	1622
10		
11		
12		
13		
14	15	1/4 = 141
15	9	1113
16	all (-)	
17	22	1766
18	15 (18)	1/4 = 479
19	est. 100	est. 2000-2000
20	19	897

Calculations on next page - also notes -

50

*. These platings have ca. 3000 cells
 the no. (-) appears to vary from ca 10-200

$$a = 0.602 (\text{no. mutant}) / (N \log N)$$

DATE:

REF:

Inoculation	in pipette	3	4	5	6	7	8	9	10
No. (-)	Total cells	$N \log N$		$0.602 R$		$S/3$	Recalculated		
6	1169	$1169(3.0684) = 3580$		3.612		1.0×10^{-3}	1.0×10^{-3}		
3	595	$595(2.7745) = 1650$		1.806		1.1×10^{-3}	1.1×10^{-3}		
4	251	$251(2.3996) = 600$		2.408		4.0×10^{-3}	4.0×10^{-3}		
23	1252	$1252(3.0976) = 3880$		13.846		3.6×10^{-3}	3.6×10^{-3}		
9	1113	$1113(3.04508) = 3400$		3.418		1.1×10^{-3}	1.6×10^{-3}		
19	897	$897(2.95274) = 2650$		11.438		4.5×10^{-3}	4.3×10^{-3}		
0	237	-		-					
100: Resuspended									
7	1295	$1295(3.11227) = 4030$		4.214		1.0×10^{-3}			
12	1222	$1222(3.09207) = 3780$		13.244		3.5×10^{-3}			
52	730	$730(2.86332) = 2085$		31.304		1.5×10^{-2}			
0	81								
0	52								
103	2750	$2750(3.439) = 9430$		62.00		6.6×10^{-3}	6.6×10^{-3}		
317	1622	$1622(3.21005) = 5200$		49.2 192		3.7×10^{-3}	3.6×10^{-3}		
20	22	$1966(3.29358) = 6460$		13.2		2.0×10^{-3}	2.8×10^{-3}		

Estimated P_0

Using 200
$$a = \frac{2.3}{200} \log \frac{1}{\frac{1}{12}} = (0.012)(\log 12) = (0.012)(1.08) = 1.3 \times 10^{-2}$$

Using 1024
$$a = \frac{2.3}{1024} \log \frac{1}{\frac{1}{10}} = \frac{2.3}{1024} \log 10 = \frac{2.3}{1024}(1) = 2.2 \times 10^{-3}$$

$$\frac{3.20}{7} = 2.4 \times 10^{-3}$$

30

40

50

DATE: 6/22/55

REF:

yield of HFT from hetero source

1. Culture 241-14 (HAI/A) - streaked from stock and 10 colonies picked, tested for HFT against 2279 - Culture showed Pen cross freq

1 0 0 0 0 0
0 0 0 0 0

2. Reacti^{HFT} not apparent 24 hours - select most likely appearing Col and proceed. Points incubated 34 hours - incubated later 10^H 9

3. 4 ppt. a. dilute 1.0 ml per 6 plates ca. 1000 col/ml in Dulbecco. Inoc 35 seconds and plate ⁶⁰⁰ 600 gal with 750 Col. - Make per mod. count b. Inc. 2 hours - respread 3 plates

4. Results:

Per mod count / 0.1 ml = 67
50
46 } ave = 53

Per mod plaque / 2279
0.1 ml = 22
12
14 } ave 16

Per mod. p4p / 737 (1) 2
NOT SPREAD (2) 5
(3) 2 } ave 3

SPREAD (1) 3
(2) 2
after 2 hours (3) 3 } ave 2

Col	PAF	2/5 + ^H
5	3	(1) 3/5 + ^H
8	4	(2) 5/8 + ^H
5	2	(3) 2/5 + ^H

52B	52C	52D
1	1	1
2	2	2
3	3	3

10

20

30

40

50

DATE: 6/22/55.

REF:

Low multiplicity, ductins - does λ segregate?

1. Culture - 2277 overnight
 $\lambda = 241-14$ (10/12/54) titer 5.3×10^8 phages } see p 308
 1.7×10^8 heads }

2. @ 0.4 ml 2277 + 0.1 ml HFT 2-
 " " " 0.1 ml burst

3. incubate 37C 10 min.

10 4. dilute $10^3, 10^4, 10^5 \rightarrow$ plate 0.05 ml
 λ - 6 plates

5. Control 3 plates - 53, 47, 49 one Col - $50 \times 10 \times 10^6$ - 1.0×10^9

6. E₁ phage plates (+) titer

1	66
0	54
0	40
0	41
0	54
0	36
(1)	311

no (-) found cultures with λ or phage lysed.

tested against 7279 and found not byogenesis

353-1

minimum Rpt.

1. Culture = 2344 JL's δ stock Col₂ - by 5 Hft
 taken from his stock culture labeled δ

2. Lyt₁ = 7506 R2-1 1/28/55 titer =

trans. titer = 2.6×10^7 see 311

3. Sept. 0.1 ml Lyt₁ + 0.1 ml cells inc. 37C 15' - 10 ml Pen added; diluted $10^3, 10^4 \rightarrow$ 0.1 ml

4. Results - A. control plating - exposure to both

Plate 1 237 }
 2 326 } $2.82 \times 10 \times 10^4 \times 10^2 = 1.41 \times 10^9$ cell/ml overnight cult.
 3 283 }

B. Lyt₁ lysed

	+	*	(-)	(-) ^{partially} lysed	total
1.	3		92	0	95
2.	2		87	4	90
3.	0		76	1	77
4.	3		71	4	78
5.	1		61	0	62
6.	0		94	1	95
7.	0		96	2	98
	9		9	12	595

from P₁ trans. 750 R2-1
 titer = 5.6×10^7

* the + appear to be in two categories: (1) appearing ca. 15 hours and represented by intact (not lysed) colonies that are almost all (+). (2) appearing in (-) colonies, λ contaminated and partially lysed, after ca. 24 hours. In the above there were 5 were early, 3 late. This correlation was noted.

After	lysis of phage	Frocher Col ₁
early	2/5	predom. (+)
late	3/3	" (-)

} one trans. but mixed - Hft - δ

C. Plates - primary stocks of (+) given to # 1-5 early - 2/5 P₁ trans
 6-8 late - 3/3 P₁

EML 900
 give letters A-H

50

DATE: 6/30/55

REF:

Crossed rate in P.E. heterog. 342E2 6-4-4
 Sept. as in 357 with 710CKR-1

1. Overnight culture from single colony
2. Dilute 10⁻² - 10⁻⁴ - 10⁻⁶ - 10⁻⁸

incubated 9:30 - 2:30

Plates all 20-40
 low because
 very
 used

10

3 plates for estimate of total viable count

10 ⁻²	10 ⁻⁴	10 ⁻⁶	10 ⁻⁸	CFU	(+)	Total	Max Prob
1. 4	67	(8 col.)	-	7	0	0	-
2. 4	78	(7 col.)	-	2	0	1406	1.3 x 10 ⁻⁴
3. 6	78	(8 col.)	-	3	0	ca. 3000	6.0 x 10 ⁻⁵
	ave 71	0.71/plate	-	4	0	1/2 = 193-1544	-
		0.14	-	5	0	1/2 = 193	6.0 x 10 ⁻⁵
			-	6	0	0. contamination	-
			-	7	ca 1000	ca 3000	-
			-	8	0	ca. 150	-
			-	9	4	1/2 = 643	ca. #3 1.3 x 10 ⁻⁴
			-	10	0	ca. 4000	ca. #3 4.1 x 10 ⁻⁵
			-	11	2	1/2 = 841	6.3 x 10 ⁻⁵
			-	12	5	2266	4.0 x 10 ⁻⁴
			-	13	0	0	-
			-	14	1	1/2 = 370	3.0 x 10 ⁻⁵
			-	15	0	ca. 7000	2.5 x 10 ⁻⁵
			-	16	ca 1000	ca 2000	-
			-	17	0	0	-
			-	18	0	2245	-
			-	19	0	748	-
			-	20	0	34	-
			-			total 29128/10 = 2913	

Using milk method and N = 5950 (from culture #4)

20

$$a = \frac{2.3}{5950} \log \frac{1}{\frac{1.7}{17}}$$

$$a = (0.00039) (0.176) = 7.0 \times 10^{-5}$$

$$a = \frac{2.3}{2.9 \times 10^3} \log \frac{1}{\frac{1.7}{17}} = \frac{0.405}{2.9 \times 10^3} \times \frac{4.05 \times 10^{-4}}{2.9} = 1.4 \times 10^{-4}$$

30

Reynolds 342C1 No above - except plate picked to isolate pen and this given overnight

Time	Count	Total
10:30 AM	5950	577
1:45 PM	2	595
	0	556
		583 = 5.8/0.1 ml

(+)	Total	Max Prob
1	2	728 276 (1003)
2	0	818 62 (880)
3	0	116 110 600
4	0	1/2 = 86 688
5	0	1/2 = 50 640
6	0	1/2 = 49 392
7	0	1/2 = 50 670
8	0	1/2 = 91 728
9	0	1/2 = 85 644
10	0	1/2 = 149 1192
11	0	1/2 = 62 496
12	0	1/2 = 83 664
13	0	1/2 = 137 1096
14	0	1/2 = 71 592
15	0	1/2 = 90 720
16	0	1/2 = 152 1216
17	1	185 85
18	0	1/2 = 87 696
19	1	1249 1.4 x 10 ⁻⁴
20	0	1/2 = 62 496

Using milk method, and N = 1200 (A.F.)

40

$$a = \frac{2.3}{1200} \log \frac{1}{\frac{1.7}{17}} = (1.7 \times 10^{-3}) (0.13) = 2.6 \times 10^{-4}$$

$$a = \frac{2.3}{779} \log \frac{1}{\frac{1.7}{20}} = \frac{2.3}{779} \log 1.2 = \frac{2.3}{779} \times \frac{0.087}{7.79 \times 10^2} = 2.4 \times 10^{-4}$$

50

50 | 15588 1559 (779)

DATE: 6/30/57

REF:

1924 x - 2346 A to look at single heterocyst

1. 0.1 ml overnight culture + 10 ml

0.1 ml + 0.1 ml 2346 A 9/11/54

↓
incubated 37C - for 10"

↓
add 10 ml Pen → die 10², 10⁴ → 0.1 ml sample

2. Best - control
2346 A 9/11/54 - 2 cell.

10

20

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DATE: 7/6/55

REF:

Seiyaku from W2730

1. Sept 6 before - see page 357

2. Overweight count

$10^2 - 10^4 - 10^5 - 10^8 \rightarrow 20.0 \text{ line sample}$

10

o.l.m.s (#)	total
1. 17	122
2. 19	168
3. 21	151
19	441 = 147

= 1.5 cm/plate

28/19

Using null factor

$$a = \frac{2.3}{100} \left(\log \frac{1}{p_0} \right) = \frac{2.3}{100} \left(\log \frac{1}{0.5} \right) = 0.012$$

Estimated clone

size \rightarrow 40, 80, 200, 400, 40, 80, 150, 100, 30

ave est. 150

$$= 1.2 \times 10^{-2}$$

20

(-)	total	o.l.m.s
1. all	323	8.1×10^{-3}
2.	176	3.0×10^{-3}
3.	0	0
4.	1668	2.8×10^{-4}
5.	317	2.2×10^{-3}
6.	1236	8.1×10^{-3}
7.	10	0
8.	0	0
9.	ca. 200	ca. 700
10.	36	1055
11.	0	0
12.	0	0
13.	3	299
14.	0	0
15.	0	0
16.	6	386
17.	0	0
18.	0	0
19.	55	1965
20.	-	0

$$\frac{45.2}{5.1}$$

30

- 8. failed to grow
- 9. contaminated with Galt
- 10. contaminated with Galt & Galt

40

50

DATE: 7/13/55

REF:

1. Rpt 2344M1 (Sh. shole) ~~HT~~ \times HFT \neq
 Lysea = 293-125 -

2. 0.3mc Lysea + 0.2mc overnight cult.

↓
 add some broth
 ↓
 $1-10^7$
 ↓
 0.1mc

broth @ control.

10 Plates bad
for cult.

3. Resuel

A. Control.

$\frac{1}{10} \cdot 10^7 \cdot 10^2 \cdot 10 = 6 \times 10^7$

- 1. 457
- 2. 540
- 3. wet plate

5.0×10^7

? W(?)

B. EPH

(+)

total (-)

control (-)

Deny:

1.

2

322

2

02 01

20

1

ca 300

✓

03

(-5)
 Part of plate theme seen in center counts -
 above the plane or above - (after 2 days)
 some (H) appeared - maybe reversions - This
 culture seems to revert more than the Gals -

30

40

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DATE: 8/10/55

REF:

	1	2	3	4	5	6	7	8	9	10
	Permanay growth									
	λ	00	Test tubes - Blank = water - Pet4 filler							
			Airtight (ca. 10% w/v)							
	650	0.051	0.74							
	600	0.071	0.81							
	570	0.11	0.90							
	500	0.19	1.04							
	450	0.365	1.23							
	400	0.630	1.45							

10

20

Lysates	1210	K12	227C
10:30 AM 0	0.58	0.73	0.80
1:10 PM 2:20	0.75	0.52	0.80
1:25 PM 2:35	0.67	0.44	0.67
1:45 PM 2:55	0.67	0.40	0.53
	0.67	0.40	0.48

$\frac{0.2}{20.0} \times 5 = 0.05\%$
RNA

30

9/25/55
750 K12-1 grown on B gal + 0.05% RNA

1. no RNA 5 gal - + ca 10 gal + ca 60 gal +
2. RNA 6 gal - + ca " + ca 20 gal +

} No effect of RNA at this conc

40

1210 λ x 750 in search of HFT f

no add 2
0.1% 18 from lysate

HFT test	No. tests	against?	HFT?
1	10	750, 2341	possibly no/2341 - lysate made 358-1
2	10	750, 1210	possibly no/750 358' 2

} none HFT - new lysate of bits made - both still

50

2279 + 750 K12 8/10/55

1. 750 at 142x: 568
2. 2279 at 0.6x: 426

Plaque low titer on λ - this lysate has been at room temp since it was made

no control

359A

29/15 used for 4th rx

DATE: 9/28/55

REF:

	1	2	3	4	5	6	7	8	9	10
Rev of second strand	359A ¹ Dose 2 ¹ 2777	359A ¹ Dose 2 ¹ 2777	359A ¹ Dose 2 ¹ 2777	359A ¹ Dose 2 ¹ 2777	359A ¹ Dose 2 ¹ 2777	359A ¹ Dose 2 ¹ 2777	359A ¹ Dose 2 ¹ 2777	359A ¹ Dose 2 ¹ 2777	359A ¹ Dose 2 ¹ 2777	359A ¹ Dose 2 ¹ 2777
1	S W 0	Lp ¹	V S +	V S +						
2	R S 0		V S +	V S +						
3	S W 0		V S 0	V S 0						
4	S W 0		V W +	V S 0						
5	S W 0		V S 0	V S 0						
6	10 ¹ R S 0	Lp ¹	V S 0	V S 0						
7	S W 0		V S 0	V S 0						
8	S W 0	Lp ¹	V S 0	V S 0						
9	S W 0		V S +	V W 0						
10	S W 0		V S 0	V S +						
11	S W 0		V S 0	V S 0						
12	S W 0		V S 0	V S 0						
13	S W 0		V S 0	V S 0						
14	S W 0		V S +	V S 0						
15	S W 0		V S 0	V S 0						
16	27 ¹ no growth		V S 0	V S 0						
17	S W 0		V S 0	V S 0						
18	S W 0		V S 0	V S 0						
19	S W 0		V S 0	V S 0						
20	S W 0		V S 0	V S 0						
21	S W 0	Lp ¹	V S 0	V S 0						
22	S W 0		V S +	V S 0						
23	S W 0		V S 0	V S 0						
24	S W 0		V S 0	V S 0						
25	R S 0	Lp ¹	V S 0	V S 0						
26	S W 0		V S 0	V S 0						
27	S W 0		V S 0	V S 0						
28	S W 0		V S 0	V S 0						
29	R S 0	Lp ¹	V S +	V S 0						
30	S W 0		V S 0	V S 0						
31	S W 0		V S 0	V S 0						
32	S W 0		V S 0	V S 0						
33	S W 0		V W +	V S 0						
34	S W 0		V S +	V S 0						
35	R S 0	Lp ¹	V S 0	-						
36	S W 0		V S 0	-						
37	S W 0									
38	S W 0		9 Lp ⁺	2 Lp ⁺						
39	S W 0		28 Lp ⁺	32 Lp ⁺						
40	S W 0									
41	S W 0		2nd Lp ⁺	33 S Lp ⁺						
42	S W 0		34 S Lp ⁺	1nd Lp ⁺						
43	R S 0	Lp ¹								
44	R S 0	Lp ¹								
45	S W 0									
46	S W 0									
47	S W 0									
48	S W 0									

Reverse 10/3/55
 o.l.m. 10th Dec / 2777
 C = 25 ca. Spl: 1E
 359A0 0 = 355¹ L₆₀
 359A1 1 = 672 ca 2A: 1E
 2 = 678 ca 1A: 1E
 Array of tubes for prot. 1, 4th rx

10 Lp⁺ = 9
 60 Lp⁺ = 38

3/12
2010-2010

35913

DATE: 10/6/55

REF:

Continuation of 359 359A

Plating of lead 359-1 by sale on 2580 (Lpt) - 0.1ml 10⁵ dilution of
 Vulture Repeats Assay/2391 (same dia ->) mediated by sale

Counters	Repeats	Net	Assay/2391
0	31	44	35
1	75	142	
2	183	347	
3	388	545	
4	576	689	253
5	710	832	
6	863	951	

#10 streaked
 out - 10 ul.
 Agard 2391
 all numbers
 359B(4)
 7m dia
 #10

5-54
 45-400
 to 400
 0-400

on the
 not used

359B1
 also 359B

359B10
 359B11

10

40

50

1	2	3	4	5	6	7	8	9	10
1	ac +	v s +	2-Lpt	1-Lpt	2-Lpt	7-Lpt	7-Lpt	2-Lpt	7-Lpt
2	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-7-Lpt	2-Lpt	2-Lpt
3	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
4	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
5	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
6	us +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
7	us +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
8	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
9	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
10	us +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
11	s +	v us +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
12	s +	v s +	7-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
13	us +	v us +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
14	s +	v us +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
15	us +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
16	s +	v us +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
17	s +	v s +	7-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
18	us +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
19	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
20	s +	v us +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
21	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
22	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
23	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt
24	s +	v s +	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt	2-Lpt

* Cl...
 other...

14 ends 2 etc	14 ends 1 etc 1 amplif	16 ends	16 ends 1 etc	16 ends 1 etc 1 amplif	17 ends 1 etc	110 6 2
------------------	------------------------------	---------	------------------	------------------------------	------------------	---------------

55 17
 7 1
 1 2

DATE: 8/18/55

REF:

1924 x - HFT 1 to study lyogenization + crossing over
 1-ly⁺ x 1-ly⁺

1. Transd. plates not needed - ca. 300 colonies/plate. No gal⁺ on control, lysate exposed.
2. After 2 days on the lysate plate several reappearing colonies
3. Results

Reappearing colonies streaked out and from the primary streaks 8 (-) and 8(+) tested for lyogenization against 2279

Grant
 Give to
 Brodick

Rep. Cd.	8 gal ⁺ Gal ⁺ + ly ⁺ ?	8(-) ly ⁺ ?	8(+) ly ⁺ ?
1	no ly ⁺	no ly ⁺	no ly ⁺
2	no ly ⁺	no ly ⁺	yes
3	no ly ⁺	reprod.	no ly ⁺
4	no ly ⁺	no ly ⁺	no ly ⁺
5	no ly ⁺	no ly ⁺	no ly ⁺
6	no ly ⁺	yes	yes
7	no ly ⁺	yes	yes

} discarded

4. 360-2 = 24 Gal⁺ colonies from 24 diff. $\frac{+}{-}$ colonies. Tested 12279 for lyogenization. All found but lyogenic.

Row	360-2	12279	12279	HFT	Genotype	Lyogenization	Gal ⁺ still?
1	360-2	no ly ⁺	no ly ⁺	1	1-4-	ca. 1000 plates	-
2	360-2	no ly ⁺	no ly ⁺	1	1-4-	"	-
3	360-2	no ly ⁺	no ly ⁺	0	1-	"	(1) 1 found other
4	360-2	no ly ⁺	no ly ⁺	0	1-	"	(1) 1 found other
5	360-2	no ly ⁺	no ly ⁺	0	1-	"	(2) 2 found other
6	360-2	no ly ⁺	no ly ⁺	+	4-	"	(3) 3 found other
7	360-2	no ly ⁺	no ly ⁺	+	4-	"	(3) 3 found other
8	360-2	no ly ⁺	no ly ⁺	0	1-4-	"	-
9	360-2	no ly ⁺	no ly ⁺	0	1-	"	(1) 1 found other
10	360-2	no ly ⁺	no ly ⁺	1E(-)	4-	"	(5) 5 found other
11	360-2	no ly ⁺	no ly ⁺	+	4-	"	(1) 1 found other
12	360-2	no ly ⁺	no ly ⁺		4		
13	360-2	no ly ⁺	no ly ⁺		4		
14	360-2	no ly ⁺	no ly ⁺		3 Ampli		

Uncultured cultures (ex. 112) read. and incubated without air density ca. 10⁸

360-3
 360-4
 1 found other
 1 found other
 2 found other
 3 found other
 7 found other

DATE: 8/25/55

REF:

Observation of erotic induction:

1895 ⊗ 2790

antigen added
Aerated, cultures 3 hours from overnight

1.0 ml + 5.0 ml + 4.0 ml Pe - aerated in Antigen (4. x 10⁸ K₁₂)
2. x 10⁹ F

10², 10⁴, 10⁶

0.5 ml
1. 206
2. 206
24

Samples taken

deluted 10², 10⁴

0.01 ml plated

1. / 2229 B Gel
2. / D(0)

Optical Density
Read / 10 ml Pe
at 6.0

10² - 10⁴, 10⁶

0.05 ml

1. 209

2. 157

396 / 198

Time

0
15
30
45
60
75
90

0.0

0.95
—
1.10
1.7
2.2
2.0
2.2

Plages/20%

7 *
19
36
69
147
ca 110

Perh. 0(0)

3
1
0
3
11
6

Whitkop
Lp

35
15
—
35
11.5
6.5

Ratio of $\frac{K_{12}}{F} = \frac{20.6}{99} = \frac{1}{5}$

4.2 x 10⁷ - 4.1 x 10⁹

4.00 x 10⁹

* Since these plates are plated on B gel
estimation of 18% G₁₂ content
can be made
appear to be ca 10⁸ at 0
and no reduction in
numbers up to 90 min
noted.

7 x 10⁹ x 10⁴ =

140 x 10⁴ = 1.40 x 10⁶ / ml at 120

$\frac{150}{7} \times 1.4 = 29 \times 10^6 = 2.9 \times 10^7$

$\frac{29}{1.4} =$

$\frac{84}{2.1} = 29.4$

$\frac{2.9}{11} = 72.7 \text{ } \frac{10^7}{\text{hr}}$

$\frac{0.07}{410} = 17.00$

40 2341 ⊗ 2308 Mal - Inoc. 2341 to look for translocation and seg behavior of K₁₂ to

- 2341 aerated wet. inoc. 60 sec. - Mixed equivalent volumes of 10² F - inoculated 20 min 37C deluted 170x and plated B gel, S Mal - 5 Mal plates discarded - difficult to plate from Mal.
- 20 G₁₂ plate high on S Gel picked and tested / 2779 for isogenicity all non isogenic

The 20 G₁₂ are divided into several groups on degree of G₁₂-ness

- 12 full G₁₂ - these do not segregate
- 3 intermed G₁₂
- 5 weak G₁₂ - these contain a few full G₁₂

} all appear to be seg.

363 B

50

DATE:

REF:

	1	2	3	4	5	6	7	8	9	10
	Segregant display. Test by reversing segregant, ✓ stability									
	Cushion.			Rev. stab	0 = no	+ = yes				
	1	5186142		0						
	2	5186750		0 (ult +)						
	3	5186902-1		0						
	4	5186902-2		0						
	5	21756811-1		0 (ult)						
	6	21756811-2		-						
10	7	7506412-1		0						
	8	7506412-2		0 (ult +)						
	9	7506902-1		-						
	10	7506902-2		0						
	11	8167902-1	all +	no						
	12	8167902-2	all +	0 (ult)						
	13	286-1		-						
	14	286-2	guided to gene	-						
	15	217567301		0 (ult)						
	16	217567302		-						
20	17	290-1		-						
	18	295-2		-						
	19	302-2	7x2-	-						
	20	33448	2:1-67-	0						

10/10. oncom for test, no seg = 10 seg int display

311-2 = HFT 6-

obtained out 2 Galt recessive obtained

both segregating G_{alt}⁺/G_{alt}⁻

2 HFT seg obtained by 1x 12 16
+ + a = 6-

HFT
6-
+/-

309-1 = HFT 7-

shed out 8 G_{alt}⁺ recessive obtained

2/8 segregating G_{alt}⁺/G_{alt}⁻

HFT
7-

2874

looking for G_{alt}⁻ in 583 level

1st fruit 8 seg = G_{alt} 7-
2nd " " = "

50

DATE: 8/31/55

REF:

Segregation - non disjunction? Obtain clone with early seg and test for Gal+ component to see if it segregates 50% homozygous.

7. Culture W2869 - from single (+) colony -> overnight per. dilute

13.3/0.01 ml
10⁷ - 10⁶ - 10⁵ - 10⁴

10 (-) Spent 0.01 ml samples B Gal 0.01 ml incubate 3 hours and respond. - look for clones with many
0.1 ml for Gal+ Gal- plates

1.	19	117
2.	17	142
3.	19	139
55-18		398-133/0.1 ml
7		3

Reproduction

1. failed to grow or vigorous - 19
2. one gal+
3. one gal-
4. mixed Gal+ + Gal- - 15

nois used for cog. study

$\bar{x} = 1.3$, $e^{-x} = 19.4$
Gal+ $\bar{x} = 1.2$, $e^{-x} = 21.6$
Gal- $\bar{x} = 0.18$, $e^{-x} = 59.76$

Plate	Colony	Gal	Other	Notes
1	1	13+	no other (-) on plate	1+
20	46	2	38+	+
2	11	3	9+	2+
	45	4	37+	+
3	8	5	7+	3+
	39	6	24+	+
	24	7	14+	+
	41	8	31+	+
4	33	9	32+	4+
5	11	10	10+	5+
	9	11	7+	+
30	36	12	22	+
6	12	13	11+	6+
	5	14	4+	+
	one 12	15	11+	+

2 Gal+ reversions selected from each all found Gal+ still localization of clone on plate and relationship to other Gal- in case of spreading from one streak to another

40 Segregants from three Gal+
#6 col A = 1 (2-), 5 (4-)
B = 3 (2-), 3 (4-)
C = 2 (2-), 1 (4-)
D = 1 (2-), 4 (4-)

Segregants from #1 Gal+
A. 8 (2-), 5 (4-)
B. 2 (2-), 3 (4-)
C. 4 (4-)
D. 5 (4-)

OPNET
A1 (2-)
24 (4-)
B1 (4-)
82 (2-)
85 (4-)
50
C1 (4-)
C2 (2-)
D1 (2-)
D3 (2-)
D4 (4-)

Gal+ D
Reversions (2) others from three and tested for stability
2 des of A1 mixed (6+) Gal. United 750
B1 (4) still
K1 B1 K1 unstable - NFT segregants obtained % Gal-
4 cultures (2+)
5 NFT
7 HET
10% col. tested 1 HFT/81
- NFT segregants obtained % Gal-

DATE: 9/25/55

REF:

A

293-12 (which are) give increased frequency of HFT Gal⁻ Use to test for disorder of distal markers on a subset of C.O. between Gal's.

1. A mal obtained by 1² passage in B mal.
2. 12 seg obtained, tested for HFT against 2580, PH

control	2580	84
0 0 0 0	0 0 0 ⊕	0 0 ⊕ 0
0 0 0 0	0 ⊕ 0 0	0 0 0 0
0 0 0 0	0 ⊕ 0 ⊕	0 0 0 0

Total
13 HFT 4⁻
37 +
17/49 ~ 16/79

⊕ = control
⊕ = HFT

1/12 HFT 4⁻

mult passages

3. Streak HFT 4⁻ in B mal, 1/6 mult² stbl, 1/6 mult¹ stbl, #10 1/6 mult¹ stbl, #12 1/6 mult¹ stbl
4. Test the seg 1/2⁻ 1/4⁻ 1/4⁻ + (these are mod -)

A. Despite mal reaction - after 3 days at 37C, reaction obtained - one appear to be transduced by +, 2⁻, but not by 4⁻ - means all Gal⁻ - Gal⁻ is in response
B. Reversions of #4, 6, 10, 12 obtained to check HFT.

The HFT⁺ above (A1) and 3 from A2 grown with 1177 24 hours and plated in S.Mol. Env. no growth - indicating these cultures not HFT⁺.

Reversion above say culture not homo-geneous ~~why?~~

Subpt. on 12 additional seg.

control	2580	84	1/6 mult ¹ #3	1/6 mult ¹ #6	1/6 mult ¹ #7	1/6 mult ¹ #8
0 0 0 0	0 0 ⊕ 0	0 0 0 0	0 0 0 0	0 0 0 0	0 0 0 0	0 0 0 0
0 ⊕ 0 0	0 ⊕ 0 0	0 0 0 ⊕	0 0 0 0	0 0 0 0	0 0 0 0	0 0 0 0
0 0 0 0	0 0 0 0	0 0 0 0	0 0 0 0	0 0 0 0	0 0 0 0	0 0 0 0

CONTINUED PAGE 366

A2
The Gal⁻ are from mixed

6. 293-12 associated in Bgal, Bacc, Bantl. - 1/4⁺ stbl, (1 with T?) not gal⁻/gal⁻

2852 = 365B
Gal⁻ Gal⁻ x - 2 - to obtain (- -) for stpt. to show

crossing over between parents, as $\frac{+}{-} \times \frac{+}{-} = \text{not Gal}^+, \text{C.O. between parents to give } \frac{+}{+} \text{ yields Gal}^+$

1. 2852 checked / HFT 2⁻
2. 10 streak made from cross bank. 7 yielded 1/4⁺ from 1. 3 yielded 1/4⁺ from 2. 1/4⁺/1/4⁺ same seg 1-4⁻ 1/4⁺
3. Obtain 1-4-2/4⁺ 1-4-1/4⁺ 1-2-4⁺ 1-2-4⁺ → mixed Bgal → 7 seg obt. 6 = 1-2-4⁺, 1-4-4⁺

Reversion 365B-1 mix of 365B-2

B

10

20

30

40

50

DATE: 10/8/55

REF:

2341 $4^r/4^r$ $2/2$ X — (+) 750 CK12-1

1. 2 day cult. 2741 → 1-10

↓
0.1 ml + 0.1 ml 750 CK12-1 → $10^2, 10^4$ → 0.1 ml
inc. 10^7 37C

2. same with both as control →

10

		(-)	(+)
Control	1.	150	0
	2.	167	0
Sept.	1.	173	1
	2.	158	2
	3.	162	2
	4.	129	2
	5.	189	2

(3-4) several partially lysed colonies/plate

One streaking

Cont	Sub	SE	Strik
1 4^r	4^r	→	4^r + -S
2 4^r	4^r	→	4^r + -S
3 4^r	4^r	→	4^r + -S
4 4^r	4^r	→	4^r + -S
5 4^r	4^r	→	4^r + -S
6 4^r	4^r	→	4^r + -t
7 4^r	4^r	→	4^r + -t
8 4^r	4^r	→	4^r + -t
9 4^r	4^r	→	4^r + -t

A

20

These discarded

365B2 = H⁻ Gal⁻ Gal₂⁻ Gal₂⁻ 4^r X HET 7 to see quadruple (-)

1. Procedure as with 2341 above. 309-1 lysate used.

30

		(-)	ly. col.
Control	1.	428	0
	2.	454	0
	3.	503	0
Sept.	1.	171	15
	2.	189	27
	3.	196	20
	4.	206	33
	5.	200	46

There are low, perhaps 50% with (A0%) colonies partially lysed

B

40

50

DATE: 10/18/55

REF:

368A

1. Examination of the later grades - 6 gals, 6 gal - taken from the 1st streaking out of the isolated (single colony) heterocysts; and streaked against λ , 2279

	1	2	(bridge)	4	5	6	7	8	9
Gal +	1	+	+	+	R	+	+	+	+
	2	+	wt	+	R	+	+	+	+
	3	+	+	+	R	+	+	+	+
	4	r ?	+	+	R	+	+	+	+
	5	+	wt	+	R	+	+	+	+
	6	+	+	+	R	+	+	+	+
Gal -	1	+	+	S	S	S	+	+	+
	2	S	S	+	S	+	+	+	+
	3	S cutan	S	+	S	S	+	+	S
	4	+	+	wt	S cutan	wt	+	+	wt
	5	wt	+	+	S	wt	+	+	+
	6	S cutan	S	+	S	wt	+	+	S cutan

Reason of note on which this was based: which shows good λ R₁ #16, the plate shows self λ on plate.

	#10	#12	#13	#14	#15	#16	#18
Gal +	1	+	+	+	+	R	+
	2	+	+	+	+	R	+
	3	+	+	+	+	R	+
	4	+	+	+	+	R	+
	5	+	+	+	+	R	+
	6	+	+	+	+	R	+
Gal -	1	S	S	+	wt	S	+
	2	S	S	+	S	S	+
	3	+	S	+	S cutan	S	+
	4	+	+	+	S	S	+
	5	+	S	+	S cutan	+	+
	6	+	S	+	S	+	wt

FOR #4 #16 } put from the original culture (from the first plate) grown up in broth and spotted (put centrifuge) on 2279. One set of plates mixed, one not. both showed λ , the mixed. λ uninduced.

See 371

To AR. ^(W) new Hfr by X Gal₂ - (241-14) W
 also Gal₄ - (247R-1) - X
 From the Gal₂ - X 4 Gal₂ (Blau) 3 gal₂ 1 gal₂ re λ found by +

368A-3

50

DATE: 10/19/55

REF:

370

1. One step with 241-14 - using M62 using 2343 Gal. - plus a cell
 2915 as plaque assay
 1. 241-14 streaked out - 10 colonies picked to broth, tested for HFT.
 6 found HFT, broth of one used for test

2. Dilute fresh cult. from broth to $ca. 5 \cdot 10^8$ / ml. saline
 Pre. med. assay.

			Plate	Count	3 days	2nd	H62
10	1.	spout rev. 2343	0.1 ml in M62 =	✓ 1	0	3	+
	2.	HFT lambda assay	0.1 ml				
		+ 0.9 ml 2343 cult. -	0.1 ml =	✓ 2	0	8	+
	3.	lambda plaque assay	0.1 ml				
		+ 0.9 ml 2915	0.1 ml =	✓ 3	7		

3. Port. Inoc. = 40 seconds. ADD 5ML + 5ML 2X PEN

Time			Yield		
0'	1.	Trnd. loop	0.1 ml + 0.9 2343	= 4. ✓	0
	2.	lambda	" " 2915	= 5. ✓	348 (9)
15'	3.	Trnd.		6. ✓	0 (8)
	4.	lambda		7. ✓	341
30'	5.	T		8. ✓	0 (10)
	6.	lambda		9. ✓	ca 300-400
45'	7.	T		10. ✓	1 (6)
	8.	lambda		12. ✓	321
60'	9.	T		13. ✓	1 (2)
	10.	lambda		14. ✓	360
75'	11.	T		15. ✓	0 (8)
	12.	lambda		16. ✓	740
90'	13.	T		17. ✓	1 (5)
	14.	lambda		18. ✓	7417
105'	15.	T		19. ✓	
	16.	lambda		20. ✓	

3a. 6 M62 plate 0.1 ml each
 ✓ To 3 odd 0.1 ml 2343
 After 90' add 0.1 ml 2343 Respread 3 rows.

Time	Yield
28-28	ca 600 all over - 4, 2, 3
28-28	35, 8
28-26	122, 91, 105

2915 x K12 lysogenization - does lysog. occur in 10^8 / 10^9 ?

at low multiplicity - ca. 3-4 contain. cells per 1000-1500
 7 16 cylinders picked for exam. - 10 colonies from each streaking

#	Lp	+ S R	+ S R	+ S R
1	9	1 0	8	0 10(1) 0
2	3	7(1) 0	9	2 9(1) 0
3	3	7(5) 0	10	0 10(1) 0
4	10	0 0	11	7 3(1) 0
5	10	0 0	12	with base
6	5	5(4) 0	13	0 9(1) 0
7	0	10 0	14	0 10(6) 0

* = with Lp media. Streaked out and
 2 colonies picked from each.
 all Lp + media's streaked

contains 5 in percent ()

Bolt in
 ca. 600 (9)
 or char

with HFT selected
 after streaking out
 2 for
 each

B

DATE: 10/22/55

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371

6.5 x 10⁴

6.5 x 10⁴ / 100 = 650

lp⁺/lp⁻ heterogeneity - always accompanied by λ⁺?

1A. 2279 Gal⁻ - lp⁺ from HFT Gal₂ from ~~241-14~~ 241-14 Mal-ly soln. titrated 1.3 x 10⁸

A. Overnight 2279 die 0.4 ml + 10 Pa - 0.5 ml + 0.1 ml 241-14 Mal-

2. Result:

die 10² - 10⁴ → 0.1 ml sample P₁ P₂

	1	2	3	4	5	6	7	8	9	10	
10	A. Booth control.		fast	fast	wt	Percent	Count				
	1	0	624	0			+	lys	0	ml	
	2	0	684	0							
	B. 241-14 ly soln		1	6	633	26		+	lys	0	ml
			2	5	608	18		+	lys	0	ml
			3	7	592	25		+	lys	0	ml
			4	2	498	16		+	lys	0	ml

Analysis
→ Continue

372

7 0 13 +

○ Omitted from further analysis because of the fact that it is not typical of the other cultures.

30

Same as above but W518 Gal⁻ - lp⁺

A. Booth control.		0	90	0
2	0	62	0	
B. 241-14 Mal. ly soln		0	59	1
		2	47	5
		3	131	9
		4	42	2

The lp⁺ clone from 371A, 371B grown up in broth, centrifuged and spotted on 2279. One set of spots made, one non-made (20' W) none of them spouting: show λ lysis. At the same time 368A-4, 368A-16 showed λ lysis.

These cultures retested / 2279 (not used) on 12/2/55. No evidence of phage lambda found.

40

	Heterogeneity / 2279	From 1st streak	6 Gal ⁻
1	+	6 Gal ⁺	6 Gal ⁻
2	0	6 Gal ⁺	1(+) 5(5)
3	+	5/5 Gal ⁺	6(5)
4	0	5/5 Gal ⁺	3(+) 2(5)
5	0	1 Gal ⁺	4(5)
6	0	6 Gal ⁺	1(5)
		6 Gal ⁺	6(5)

← Given to JL 11/24/55

50

4 lp⁺
2 lp⁻

DATE: 10/28/55

REF:

372

371A Continued - Analysis of (+) and (-) from methylmercury

+ = Lys, 0 = unLys

	#1	#2	#3	#4	#5	#6	#7	#8	#10
6(+)	R +wk	R 0	R +wk	R +wk	S 0	L +wk	S 0	R +wk	R +wk
	R +wk	R 0	R +wk	R +wk	S 0	L +wk	S 0	R +wk	R +wk
	R +wk	R 0	R +wk	R +wk	S 0	L +wk	no test	R +wk	R +wk
	R +wk	R 0	R +wk	R +wk	S 0	L +wk	no test	R +wk	R +wk
	R +wk	R 0	R +wk	R +wk	S 0	L +wk	S 0	no test	R +wk
	R +wk	R 0	R +	R +wk	S 0	L +wk	no test	no test	R +wk

	#1	#2	#3	#4	#5	#6	#7	#8	#10
10	R +	R 0	S 0	R +	R 0	R +	R 0	R +wk	R +
6(+)	R +	R 0	no test	-	R 0	R +	R 0	-	S 0
	R +	R 0	R - 0	-	R 0	R +	R 0	-	S 0
	R +	R 0	-	-	R 0	R +	R 0	-	S 0
	R +	R 0	-	-	R 0	R +	no test	-	R +
	R +	R 0	-	-	R 0	R +	no test	-	R +

↑
(-) neg. (+) result
change than
Oppt +

↑
Gut + out
Gut + in
change +

↑
(-) Exchange
man Gut
A R

↑
Oppt +
and
but =
order reversed

	Lp (+)	Lp R	Lp +	Lp +	Lp R	Lp +	Lp R	Lp +	Lp +
1st chb	Lp (+)	Lp R	Lp +	Lp +	Lp R	Lp +	Lp R	Lp +	Lp +

	#13	#14	#15
1	R 0	R +wk	R 0
2	R 0	R +wk	R 0
3	R 0	R +wk	R 0
39	R 0	R +wk	-
5	R 0	R +wk	-
6	R 0	R +wk	-

1	S 0	R +	S 0
2	S 0	R +	S 0
3	S 0	R +	S 0
4	S 0	R +	S 0
5	S 0	R +	S 0
6	S 0	R +	S 0

40	Lp R	Lp +	Lp R
----	------	------	------

DATE:

REF:

374.

1. Enlarging the P.E. loci data

Combination of 1⁺ & 4⁻

360-3

1. 360-3 - 24 P.E. (-) picked at random

9 belt total 10 belt - say

192444FI

10

	1	2	3	4	5	6	7	8	9	10
					11 ⁻	14 ⁻				
					P.E. (-)					
				1	0	0	?			
				2	0	0	?			
				3	0	+	comb.			
				4	PE					
				5	0	0	?			
				6	PE			3	PE.	
				7	0	+		3	1-4-?	
				8	0	0		3	1-	
				9	0	+		1	4-	
				10	+	0				

2. 368-1 No above 9/15 belt data

20

6 belt - say data

	1	2	3	4	5	6	7	8	9	10
					PE.					
				2	0	+				
				3	+	0		3	1-	
				4	0	+		2	4-	
				5	+	0		1	PE.	
				6	0	+				

3. 366-2 No above 6/28 belt data

30

12 belt - say data

← This class Lp^R/Lp^S

	1	2	3	4	5	6	7	8	9	10	11	12
					PE.							
				1	0	+						
				2	0	+						
				3	PE							
				4	comb.							
				5	PE.				4	1-		
				6	+	0			3	4-		
				7	0	+			2	1-4-		
				8	+	0			2	PE.		
				9	+	0						
				10	0	+						
				11	0	0						
				12	0	0						

40



These are Lp^S except P.E.

6 belt plates were tested against all Lp^S also recheck slms. Galt dice

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376

1 2 3 4 5 6 7 8 9 10

Running run in 1-4 PE also say of possible

A 368-1 overnight from single (E.C.) CR. Inc. 9:50 AM → 2:20 PM
 B 368-2 10⁶ - 10⁴ - 10⁶ - 10⁶ → 10 0.05^u sample

368-1
 1. 0
 2. 0
 3. 1
 4. 2
 5. 3

75
 94
 96
 $\frac{75}{3} = 25$

368-2
 1. 1
 2. 5
 3. 4

3
 0
 0

82
 50
 69
 $\frac{69}{207} = 0.67/2 = 0.34$

Plat. 1000
 1. 5 493
 2. ug
 3. ug
 4. ug
 5. ug
 6. 0 ca 300 ca 600
 7. 0 " "
 8. ug
 9. ug
 10. 0 549 549

Wash 340 375 ← NOTE change in ratio

multiplicity: 0.45/sample

(+) Total

1. 2ug
 2. 0 ca 300 ca 600
 3. 0 8 794
 4. 0 1 379
 5. 346 0 ca 500
 6. 0 0 297
 7. 2ug
 8. ug
 9. 0 0 208
 10. ug

Plat. on above culture following day. Spl. done by (+) Total
 spotting 0.05^u B pet and Inc. 5.5 hrs. Ratio 1. 10 0 65
 (+) Total 17 0 108
 Cst. 173

1. 2ug
 2. 0 6 42
 3. 0 5 314
 4. 0 2 71
 5. 0 226 229
 6. 0 0 2
 7. 0 9
 8. 0 0 51

40

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7/2 - ^{Cont} 7-7 also byogenizahic of 2308 by HFT 7 (309-1) dos
 at give more $1/4/4$ than (42)?

1. Sept. procedure as on 376 E 2279 x - HFT (2-241+4 wad-)

2. Mate	Gen-Ad cont	Gen-	Total
1.	1	35	36
2.	1	29	30
3.	2	37	39
4.	1	37	38
5.	2	26	28
	7		171

171 | $\frac{0.00}{7.00} = 4\%$
~~7.00~~
~~7.00~~

10

3. From each primary check 10 clones picked and tested, 2279, 1412

Index 4	1	2	3	4	5	6	7
1. H	s	cont	ne	ne	ne	ne	ne
2. H	s	ne	ne	ne	ne	ne (cont)	ne
3. cont	s	ne	ne	ne	ne (cont)	ne	ne
4. H	s			ne	ne	ne	ne
5. H	s			ne	ne	ne (cont)	ne
6. H	s				ne	ne	ne
7. H	s				ne	ne	ne
8. H	s				ne	ne	ne
9. H	s				ne	ne	ne
10. cont	s				ne	ne	ne

ne = normal
 s = sex

ne = population,
 normal

30 cont. cont. cont.
 with d

40

50

DATE: 12/18/55

REF:

2341X - HFT + from 750CK12-1

1. overnight 2341 ~~2341~~ 0.4 ml + 0.1 ml HFT (+) → 2341 streaked out 3 gel
 ↓
 0.1 ml plated 10 column plates,
 undiluted for count took against 2
 ca 20 5 Lp 1/2
 2 Lp

del 10 ⁷	10 ⁴ 10 ⁵	0.1 ml
Gel (+)	Gel - 1/2 gel	Gel -
1 0	0	1/2 = 117 w/ plate
2 1	2	1/2 = 130 w/ plate
3 1	4	308
4 0	2	373

10

2. The Galt ³ streaked out - 6 + and 6- clones streaked 1, 2, 2, 2, 2, 2

Galt 1	2	3	4	5	6
1	2	3	4	5	6
2	2	2	2	2	2
3	2	2	2	2	2
4	2	2	2	2	2
5	2	2	2	2	2
6	2	2	2	2	2

20

Galt 1	2	3	4	5	6
1	2	3	4	5	6
2	2	2	2	2	2
3	2	2	2	2	2
4	2	2	2	2	2
5	2	2	2	2	2
6	2	2	2	2	2

30

These clones do not carry a 1⁺ particle? That entire man.

40

from stock cult. collection
 2234 for resequencing over 373, 375 X 117
 heterozygous

0.1 ml 10⁻⁴ dil HFT Gel (+) 750CK12-1
 ca. 3000 plaque
 3 small plaques - apparent to be segregating

50

Testing P. gel pour method to study segregation - P.E. clones - scoring both Galt, Gel. Also complete recovery
 1. Galt readily scored 1st day. Supposed
 2. After 2-3 days P.E. colonies Galt, Gel - colonies dark, but with light edges. Pick same to check
 17 picked - 16 pure - , 2 mixed +, -, mostly -

DATE: 12/14/55

REF:

A

Analysis of Some P.E. clus. Gal₁-//Gal₂-

~~21~~ Culture 307-1A. 21 Gal + obtained from 21 separate P.E. Gal-

5 apparent stable Gal₁ - ~~21~~ 16 ~~to~~ seg Gal₂ -

24 The Gal- /HFT 1- /HFT 2-

1	0	0
2	0	0
3	+	0
4	+	0
5	0	0
6	+	0
7	0	+
8	0	0
9	+	0
10	— P.E (-) —	
11	0	+
12	0	0
13	0	0
14	0	0
15	0	0

stke (+) - Dilute cultures made by med. overnight cult in saline adding 0.1 ml to 7.5 ml Pea - incub 3 hrs at 37

App. Stke +	stke. B Cell #	Plate cleared
1	seg	nr
2	w seg	yes
3	w seg	yes
4	w seg	nr
5	w seg	yes

* stke or cult used for plate

2 Gal₁-
4 Gal₂-
8 Gal₁-Gal₂-

HFT 1 - x Gal₂ - 270 P

2. Culture 307-1C 20 Gal + obt. from 20 sep. P.E. Gal-

4 apparent stable +

16 seg Gal-

This heterozygote is sp²/sp¹

Stke +	As above T	stke. Dgar	lysis clear	Gal Staked Tested
1	w seg	nr		S
2	w seg	nr		S
3	w seg	nr		S
4	seg	nr		R

23 The Gal- /HFT 1- /HFT 2-

1	0	0
2	+	0
3	0	+
4	+	0
5	0	0
6	0	0
7	+	0
8	0	0
9	0	0
10	0	0
11	0	+
12	0	+
13	— P.E (-) —	
14	+	0
15	+	0
16	+	0

3 Gal₁-
6 Gal₂-
6 Gal₁-Gal₂-

B

50

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1 2 3 4 5 6 7 8 9 10
 $L_p^+ \otimes L_p^+ \rightarrow L_p^+ \times L_p^+ ?$
 81 x 7035 - throwing replica plotting to test.

1. The parents:

	no. cut	no. colonies not growing $\frac{100\%}{100\%}$
W 7035	51	9
	54	12
	30	8
	39	7
	47	4
	<u>221</u>	<u>44</u>
W 811	75	0
	85	2
	73	8
	76	0
	62	6
	<u>88</u>	<u>5</u>
		<u>21</u>

no random streaks/2279

no random streaks/1279

	2035	↑	811
1	+	811	+
2	0	811	all
3	0	811	+
4	0	811	↓
5	0	811	
6	0	811	
7	0	811	but
8	0	811	may
9	0	811	really
10	+	811	so
11	0	811	
12	0	811	
13	0	811	
14	0	811	
15	0	811	
16	0	811	
17	0	811	
18	0	811	
19	0	811	
20	0	811	
21	2 up to 18 up some		

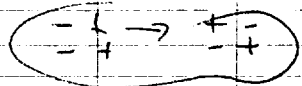
one of these
 became 2035
 = W3171

Rpt. test of 323-2. possible effect heterozygous 7: - x 1-

24 P.E. (-) picked → 21 → 21 Galt → 8 seg → 13 Galt. etc.

1 of the 8 progenies
 obtained is P.E. (-)
 Obtained after two single
 colony isolations. Indicals

Seg.	1	17
1.	0	0
2.	+	0
3.	0	+
4.	0	0
5.	0	+
6.	0	+
7.	0	0



17-
 31-
 217-

383B

DATE: 12/26/55

REF:

Yield from LFT cultures.

1. Irradiated 2550 (F-Gal⁻) — X W3005 (F-Gal⁻ Gal⁻)

2. Expt. (Prelim. expt showed feasibility of expt. Inad. amounts used as starting culture for expt.)

1. 2550 ca 5000/line → centrifuged, resuspended, volume - 40 records UV.

2. 3005
↓ spread for control = 109

Time	cpd/plate	0.1 line	cpd/plate
10	247	1. 200	697
20	227	2. "	585
30	230	3. "	704
40	198	4. "	ca 700
50	354	5. (Respread)	574
60	390	6. (after)	719
70	369	7. (2 hrs)	746
80	579	8. "	ca 600
90	546		

measured plates

✓
↓
↓
↓
↓
↓
↓
↓

2886 / 4 = 672

2639 / 4 = 658

Expt. 2/10/56

4/5

30

40

50

Alleles - Selfing

Cross of $Hfr \times F^-$ to establish that selfing does not give $Coel$ recombinants.

1. Technique. Overnight pen cultures → Parent $1.0 \text{ ml} + 1.0 \text{ ml}$ Parent

Transfer → incubate 3 hours.
 1. dilute $10^2, 10^3 \rightarrow$ plate 0.05 D(G)
 2. plate 0.05 ml undiluted H(G)

2. Counts.

	Hfr	F-
Co_1	2345	2957
Co_2	2694	945
Co_3	2487	1405

3. Expt. 1 -

	D(G) 10^2	H(G) 10^2	H(G)
(A) $Co_1 \times Co_1$	1. 22	192	80 Co_1
control spread	2. 22	ca 200	80 Co_1
Stk	3. 18	ca 200	ca 100 Co_1
	Ave. 21		
	$420 \times 10^3 = 4.2 \times 10^5$		
	no recombination.		

(B) $Co_2 \times Co_2$ - both parents Co_2 . i Co_2
 (C) $Co_2 \times Co_3$ - apparently 2877 but same Hfr population (Other lab people without this necessity)

control spread	1. 0	0
Stk	2. 0	1
	3. 0	2
	Ave. 0	Ave. 1

20×10^2

Expt. 2

(A) $Co_1 \times Co_2$	1. -	60	1	0	Hfr
both parents Co_1	2. -	74	2	0	(-)
for single (-) on Bgal	3. -	65	3	0	Stk
		199	67		

$134 \times 10^3 = 1.34 \times 10^5$

new stocks prepared 387 →

Sept. 3

(A) $Co_1 \times Co_2$

control Bgal

Stk

1.	73	0
2.	72	0
3.	91	0
	236	79

$108 \times 10^3 = 1.08 \times 10^5$

(B) $Co_1 \times Co_3$

control Bgal

Stk

1.	72	0
2.	59	0
3.	34	0
	175	57

1.14×10^5

DATE: 12/30/55

REF:

386

1. Phage content of some l_p^+ cultures - Cultures grown up and centrifuged. The supernatant centrifuged and shaken with dialysate. Inoculated on 2279 on B gel.

1	1924	no plaques
2	2035 l_p^+	contains 2 (45)
3	1898	no plaques
4	1027	contains 1 (60)
5	2537	no plaques

Phage - make sure

10

20

Segregation of l_p^+ from $\times l_p^+$ populations.

1. Culture 373-3 grown in Pen - diluted and plated on B gel

2. Second day replica'd to B(0) spread with 2279. Irradiated UV 25' serial.

Plate	Col (+)	Col -	no. not l_p^+ 2279
1.	179	13	0
2.	153	13	0
3.	221	11	0
4.	194	8	0
5.	186	13	0
6.	291	10	0
7.	225	6	0
F.	189	8	0
	1578	82	0

7th Expt

on 300 μ plate ca 20-30

0

40

X20 plate = 6000 colonies total

50

DATE: 1/10/56

REF:

388

2341 x 230c $\frac{1}{2}$ - $\frac{1}{2}$ x F $\frac{1}{2}$ - $\frac{1}{2}$

1. 1.0 me of each parent to 10 per - Jones ca. 4 lines,
2. 0.5 me parent 1.0 me
3. Colony counts 3 days:
 1. 38 Gnet
 2. 46 Gnet } seen small
 3. 35 Gnet

10

40 Gnet	40 Gnet	40 Gnet	40 Gnet	40 Gnet	40 Gnet	40 Gnet	40 Gnet	40 Gnet	40 Gnet	40 Gnet
1	2	3	4	5	6	7	8	9	10	11
1	we	we	n	n	+	+	+	+	+	+
2	we	we	n	n	+	+	+	+	+	+
3	we	we	n	n	+	+	+	+	+	+
4	we	we	n	n	+	+	+	+	+	+
5	we	we	n	n	+	+	+	+	+	+
6	we	we	n	n	+	+	+	+	+	+
7	we	we	n	n	+	+	+	+	+	+
8	we	we	n	n	+	+	+	+	+	+
9	we	we	n	n	+	+	+	+	+	+
10	we	we	n	n	+	+	+	+	+	+
11	we	we	n	n	+	+	+	+	+	+
12	we	we	n	n	+	+	+	+	+	+
13	we	we	n	n	+	+	+	+	+	+
14	we	we	n	n	+	+	+	+	+	+
15	we	we	n	n	+	+	+	+	+	+
16	we	we	n	n	+	+	+	+	+	+
17	we	we	n	n	+	+	+	+	+	+
18	we	we	n	n	+	+	+	+	+	+
19	we	we	n	n	+	+	+	+	+	+
20	we	we	n	n	+	+	+	+	+	+
21	we	we	n	n	+	+	+	+	+	+
22	we	we	n	n	+	+	+	+	+	+
23	we	we	n	n	+	+	+	+	+	+
24	we	we	n	n	+	+	+	+	+	+
25	we	we	n	n	+	+	+	+	+	+
26	we	we	n	n	+	+	+	+	+	+
27	we	we	n	n	+	+	+	+	+	+
28	we	we	n	n	+	+	+	+	+	+
29	we	we	n	n	+	+	+	+	+	+
30	we	we	n	n	+	+	+	+	+	+
31	we	we	n	n	+	+	+	+	+	+
32	we	we	n	n	+	+	+	+	+	+
33	we	we	n	n	+	+	+	+	+	+
34	we	we	n	n	+	+	+	+	+	+
35	we	we	n	n	+	+	+	+	+	+
36	we	we	n	n	+	+	+	+	+	+
37	we	we	n	n	+	+	+	+	+	+
38	we	we	n	n	+	+	+	+	+	+
39	we	we	n	n	+	+	+	+	+	+
40	we	we	n	n	+	+	+	+	+	+

5/7

20

dispute
the
30
order
completes
class
one
generation

40

ca 1/10/56
13
1/10/56

50

26
13
190

DATE: 1/15/58

REF: MLM 360-2
368-2

	2	14	21	3	4 (1/2)	5	6	7	8	9	10
10	0	+	-	4	4						
	1	+	-	4	4						
	2	+	-	4	4						
	3	+	-	4	4						
	4	+	-	4	4						
	5	+	-	4	4						
	6	+	-	4	4						
	7	+	-	4	4						
	8	+	-	4	4						
	9	+	-	4	4						
	10	+	-	4	4						
	11	+	-	4	4						
	12	+	-	4	4						
	13	+	-	4	4						
	14	+	-	4	4						
	15	+	-	4	4						
	16	+	-	4	4						
	17	+	-	4	4						
	18	+	-	4	4						
	19	+	-	4	4						
	20	+	-	4	4						

I

Separate
Gal- progeny obtained
from a P.C. clone
without going through
Gal+

{ 14 4 }
{ 4 1 }

	2	14	21	3	4 (1/2)	5	6	7	8	9	10
20	368-2(1)	-	+	4	4						
	2	-	-	4	4						
	3	-	-	4	4						
	4	-	-	4	4						
	5	-	-	4	4						
	6	-	-	4	4						
	7	-	-	4	4						
	8	-	-	4	4						
	9	-	-	4	4						
	10	-	-	4	4						
	11	-	-	4	4						
	12	-	-	4	4						
	13	-	-	4	4						
	14	-	-	4	4						
	15	-	-	4	4						
	16	-	-	4	4						
	17	-	-	4	4						
	18	-	-	4	4						
	19	-	-	4	4						

	1	4	1st
368-1	0	3	+
1.	0	0	+
2.	+	0	0
3.	0	3	+
4.	0	4	+
5.	Gal-	4	Gal+
6.	0	2	+
7.	0	1	+
8.	+	0	0
9.	0	2	+
10.	0	2	+
11.	0	7	+
12.	0	3	+
13.	0	1	+
14.	0	1	+
15.	0	2	+
16.	0	3	+
17.	0	2	+
18.	0	1	+
19.	0	4	+
20.	0	1	+
21.	+	0	+
22.	0	4	+
23.	0	6	+
24.	0	6	+

II

Significance
there is that
the way 4-
obtained is by 5-
Can induction
Just the elongation
is up?

As above

Gal- x Gal+

1/2 1/2 = 1/4

III

lysozyme
inhib
New prep being made

{ 21 1 }
{ 3 4 }

low
+
here - indicates a
homogamete

DATE: 1/12/55

REF: 389

For the purpose of obtaining $F^{-}TB_3 = Gal_2/Gal_7$ for cross with Gal_1

1. 307-1A, a P.E. heterozygote of 1+7 / 1-7+ composition (taken and } These are
 Gal - segregant from this time tested for HFT products } segregants for
 HFT 3/1500 } seg. event

P.E. $Gal_1 \rightarrow 1 Gal_1$

	1	2	3	4	5	6	7	8	9	10
Seg	1	+	0	+	0	0				
10	2	+	0	+	0					
	3	+	0	+	0					
	4	0	+	+	0					
	5	0	+	+	0					
	6	+	0	+	0					
	7	0	+	+	0					
	8	+	0	+	0					
	9	+	0	+	0					
	10	+	0	+	0					
	11	+	0	+	0					
	12	+	0	+	0					
	13	0	+	+	0					
	14	+	0	+	0					
20	15	0	+	+	0					
	16	+	0	+	0					
	17	0	+	+	0					
	18	0	+	+	0					
	19	+	0	+	0					
	20	0	+	+	0					
	21	0	+	+	0					
	22	0	+	+	0					
	23	0	+	+	0					
	24	0	+	+	0					
30	25	+	0	+	0					
	26	+	0	+	0					

14 7-
12 1-

0/26

1/26/55

Additional 27 seg tested against 2580 - unclamped

1/22 HFT against 2580 - control opt Gal - = 389-1

1st part to see if 1, 7- (Gal, -)

See 415

389-1

50

DATE: 1/25/56

REF: 390

10. ^{to fill up ms table (again!)} Observations on HFT 7-, 6-

1. 309-1 = HFT 7- stroke.

A. 12 columns tested for HFT/2580

•••••
•••••
•••••

⊕ = HFT
○ = LFT

LFT seq = 9 { ^{1/2} 1 1/4 3 = 1/9 LFT 7-

may be already related

More evidence for hypothesis

ALSO ENTER

STOCK BOOK

MINKE LYSME

HFC-

10

LFT seq revisions: 0/2 LFT seqⁿ segregating at 3rd streak

2. 361-2 = HFT 6- stroke

A. 8 columns / 2580

•••••
•••••

20

LFT seq revisions = 0/3 segregating at 3rd streak

Preparation of Stocks for Kalcha, NHT, for study of the biochemistry of galactose fermentation

30

1. 2637 K - HFT 1-, 2-, 4-, 6-, ?

Overnight culture diluted 1:100, 0.01 ml + 0.5 ml of respective HFT strokes - subsequently (after 15 min. stop! 27C) diluted 1:50 and plated 8 fold - 4 plates each.

2. Results - on all plates, colonies are plaqued - probably > 80% consisting phase 1 cells

40

	rows	columns	(-)	misses	heterozygous	Hft	(-)	misses
A) HFT 1- (2346)	1	192	0	2	of phase kept	E/L 294	0	0
	2	188	0	2		ca 300	0	0
	3	221	0	0		ca 300	0	0
	4	178	0	0	390-1	ca 300	0	0
B) HFT 2- (2342)	1	330	0	0				
	2	402	0	0				
	3	ca 300	0	0				
	4	ca 300	0	0				
C) HFT 4- (233-2A)	1	101	0	2				
	2	86	0	0				
	3	94	0	0				
	4	60	0	1	390-4			
D) HFT 6- (311-2)	1	157	0	0				
	2	205	0	0				
	3	201	0	0				
	4	234	0	0	390-6			

50

2 repeated - no misses, same means (-) missed. Under A 1/2 + 0 slightly reversible?

390-7, 1/2

to become HFT stroke in HFT stroke. This one is probably by 1/4

SEE 396

DATE: 1/30/56

REF:

391-

The cross 2252 X 341-9
 M. Gest 14₁ Gae₂/Gal₂ F- TLB₁ = L₁⁺

to see if Gae⁺/L₁⁺ are generated

1. Parents given to gether 3 hours - placed in vials (containing 3 yeast M)
2. Following day tube (kept at room temperature) replated in Gae

10

(A) M. Gest

	(-)	(+)
1.	ca. 200	ca. 22
2.	"	ca. 31
3.	"	ca. 27
4.	"	ca. 40

(B) Pick Gest and see if any + L₁⁺ L₁⁺

Day	Seq	Worms	L ₁ ⁺	L ₁ ⁺	L ₁ ⁺
1	0	(1)	4 (10)	20 (A)	(2)
2	+	+			
3	0				
4	0				
5	0				
6	0				
7	0				
8	0				
9	0				
10	0				
11	0				
12	0				
13	0				
14	0				
15	0				
16	0				
17	0				
18	0				
19	0				
20	0				
21	0				
22	0				
23	0				
24	0				
25	0				
26	0				
27	0				
28	0				
29	0				
30	0				
31	0				
32	0				
33	0				
34	0				
35	0				

19 S
 12 +
 2 R 1?

DATE: 1/31/56

REF: 392

① HFT from 2580

- ① Overnight culture diluted 1-50 incubated at 37C in serum ca 3-4 hours
- ② Centrifuged, resuspended in saline, recentrifuged, resuspended in saline
- ③ Dilute $10^2, 10^3, 10^4 \rightarrow 0.05$ ml

B gel A 1. 88
B 2. 83

ok Add 0.05 ml und. B gel SM with 3005^R Lpt

④ inoculate to serials \rightarrow
 ⑤ dilute $10^2, 10^3 \rightarrow 0.05$ ml C 1. $\frac{1}{4}$ 150 1200
 B gel D 2. $\frac{1}{4}$ ca 200 800

dilute 1.0 ml + 10 ml Pa - Inc. 37C

Time	plate 0.5 ml	3005 ^R Lpt on B gel SM
	1. unit	2. unit
12:25	0	0
12:40	15	0
Merge again	30	0
too soft - Surface	45	0
plating work	60	0
here after 0.1 ml und. with 1:40	75	1
0.1% sample	90	1
	105	9

RPT.

- 1. Overnight culture diluted ca 1-50 incubated 37C - as above. After production dilute 1-2 instead of 1-10

Serial
 ① ② 3079 on SM
 ③ ④ 3079 on B gel

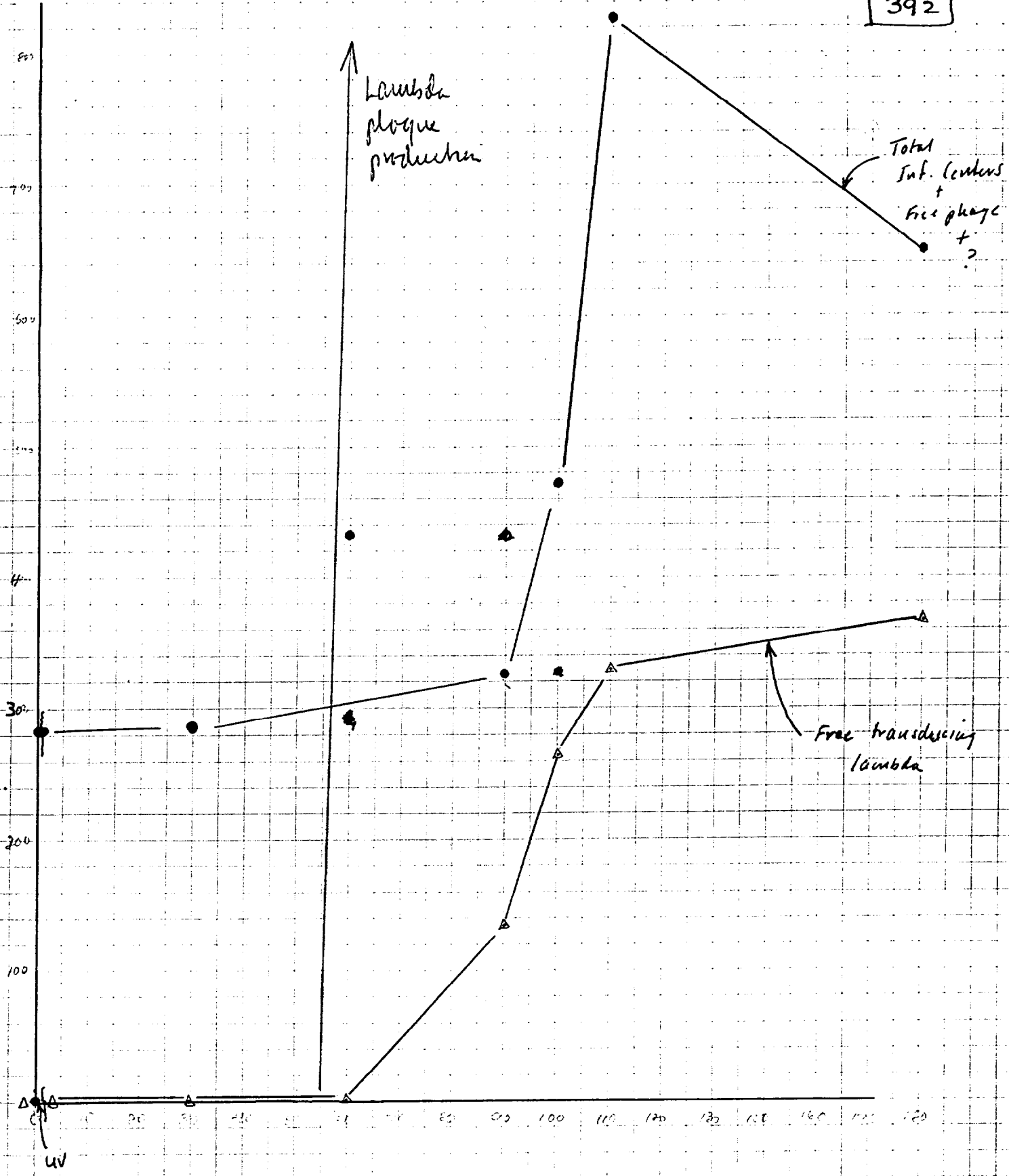
Time	Valid	Counters
30	0 ✓	79
60	13 (14)	11 (12)
90	19 (21)	15 (20)
100	21 (22)	23 (24)
110	21 (20)	27 (28)
180	21 (20)	71 (52)

to follow plaque production:
 a loopful taken out at the following intervals and spotted on 3079 on B gel SM

T	Spot
15	17 plaques
30	23 plaques
60	solid spot
90	" "
100	" "
110	" "

Time	Count	plaque	concentration	log	SM	log	SM
SM 1	ca 100 ⁺ pl.	6	pop. (count)	3.7	SM 7	112	134
4 2	ca 1000 pl.	9	pop. (count)		SM 8	153	
A 3	? count	9		8	SM 9	271	327
B 4	? "	7			SM 10	382	
7.5	ca. 1000	0		0	SM 11	218	263
SM 6	ca 1000	1(?)			SM 12	307	
0.7	count?	252	"	2.85	SM 13	472	471
B 8	count?	321	"		SM 14	470	
SM 9	ca 500 plaques	0		0	SM 15	298	328
SM 10	ca 500 plaques	0			SM 16	358	
B 11	ca?	212		2.88	B 27	744	829
B 12	"	363	up		B 28	913	
SM 13	count	3		2	SM 19	575	365
SM 14	count	0			SM 20	356	
B 15	"	334		4.33	B 31	612	180
B 16	"	522			B 32	686	649
B 17	all plaques	count					

These were
 Surface
 plated
 undiluted
 plates and
 order
 1/30/56



Lambda
plaque
production

Total
Inf. Centers
+
Free phage
+
?

Free transducing
lambda

uv

DATE: 2/11/56

REF:

393

Most of the HET stocks obtained come from ly^+ experiments. ~~But this is not~~
 Has there been selection for a lambda carrying on ly^+ (high) and low on ly^s ?
 This is the usual experimental observation - on cocktail HET lysates assay
 highest on ly^s cultures. Obtain a ly^s stock for HET cultures obtained -
 ly^s recipient. Such a culture is 364A1 - Start culture.

7. 13 colonies picked and tested for HET 1750

10

1. $\oplus \oplus \oplus \oplus$
 $\oplus \oplus \oplus \oplus$
 $\oplus \oplus \oplus \oplus$
 \oplus

20

30

2/11/56

From EML to obtain an HET (-) stock a culture labelled 7H-2961
 which is ly^s ogenic made by 7H-2- ly^s . From the plate received
 24 gals picked and streaked.

1. all requested (-)

a. One (-) was tested for HET / 2580, against 7- ly^s stock (prepared for Kakeba study)

2580	7- ly^s
00000	00000
00000	00000
00000	00000
0000	0000
00000	0000

40

ly^s used in
 all spots.

b. 20 more tested with same results - all are HET on 2580, 7- ly^s above.

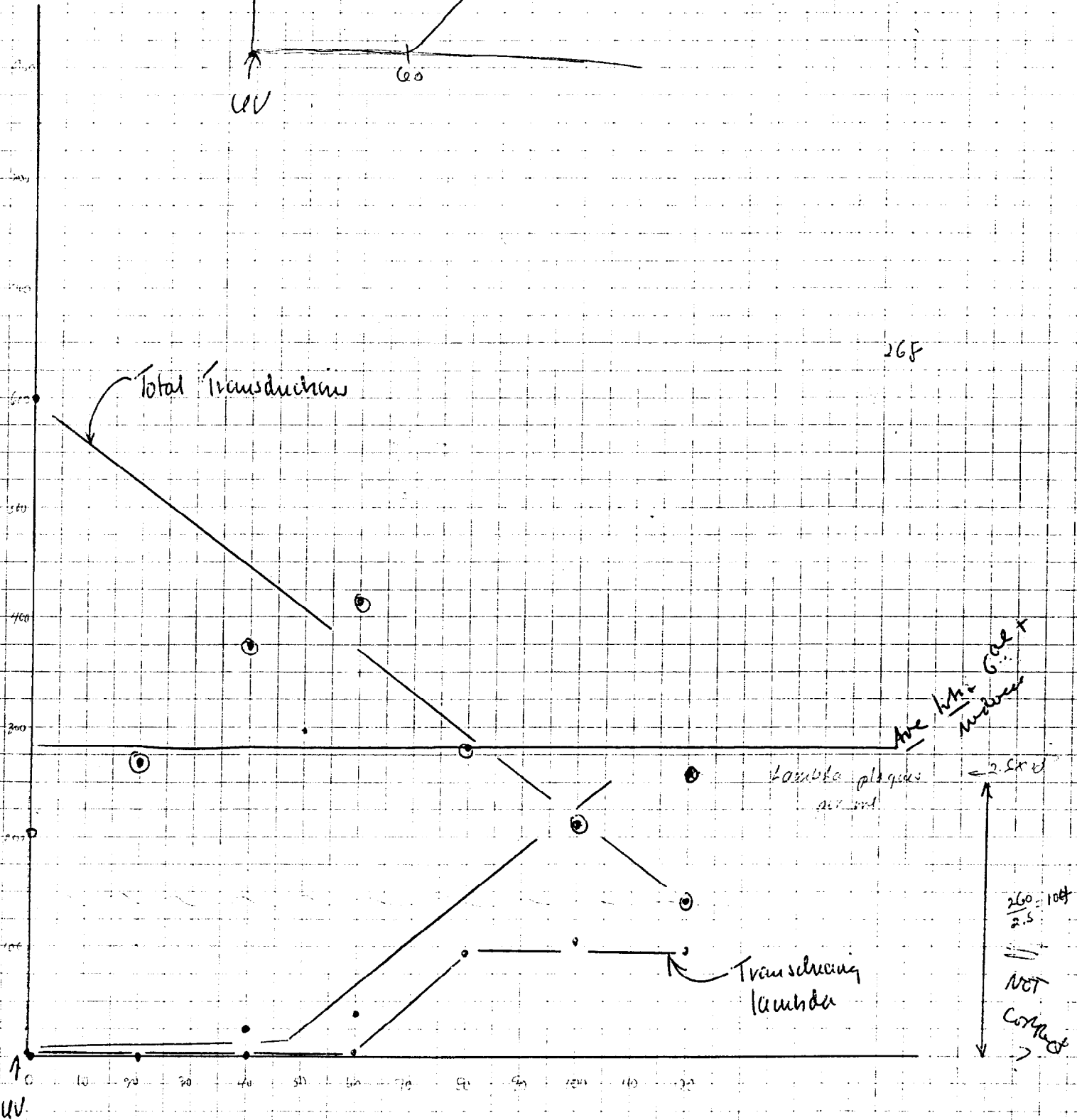
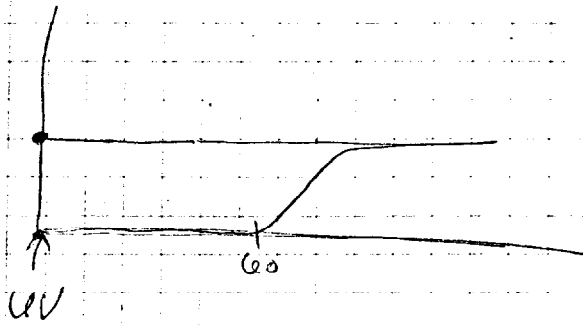
50

From here to 412
B good sm plating were
from method

DATE: 2/15/56

REF: 394

	1	2	3	4	5	6	7	8	9	10	
	2580 - overnight incubated 2.5 hrs with air. Centrifuged resusp in saline - Pre-mad. cell assay								2579	$\frac{0.51}{113.0}$ $\frac{11.9}{20.0}$	57% sum
					Basal dil $10^6 - 0.05$ ① 107 ② 110 $\frac{110}{217 \times 10^7} = 2.2 \times 10^9$		Basal 50%		on Basal ③ 12 ④ 21		
		11	11	trans assay	condensed		⑤ 0 ⑥ 0				
10	Mean from 0 condense S.M. P.M.F. w-2.1 hr.	100 sec	Port. mod. cell assay	dil 10 ⁶ ⑦ 62 ⑧ 70 $\frac{70}{132 \times 10^7} = 1.3 \times 10^9$	Basal λ assay ⑨ 26 plaque ⑩ 45 71			$71 \times 10^7 = 7.1 \times 10^8$ plaque counted $\frac{5}{14} \cdot 71 = \frac{355}{14} = 25 \times 10^8$			
			Port. mod. trans. assay used	⑪ ca 2000x 450 units ⑫ "		⑬ 0 ⑭ 0		Fluorim of content			
	Time 2 1/2 hrs	0 1:20 PM 20 1:40 PM 40 2:00 PM 60 2:20				⑮ 279 ⑯ 252 $\frac{537}{2} = 266$	⑰ 0 ⑱ 0		$\frac{1}{16}$ ca 200 plaque		
						⑲ 350 ⑳ 397 $\frac{747}{2} = 374$	㉑ 0 ㉒ 0		del 10^6 46 + 137 137 162 $299 \times 10^7 = 2.9 \times 10^9$		
						㉓ 419 ㉔ 488 $\frac{907}{2} = 454$	㉕ 1 ㉖ 3		del 10 ⁶ 163 189 $352 \times 10^7 = 3.5 \times 10^9$		
						㉗ 312 ㉘ 248 $\frac{560}{2} = 280$	㉙ 108 (many small) ㉚ 81 $\frac{189}{2} = 95$				
			0.05 ml + 3079			㉛ 175 ㉜ 214 $\frac{389}{2} = 205$	㉝ 95 (many small) ㉞ 109 (many small) 154		centrifuge 3 ml 15 sec + 3079 ㉟ 57 ㊱ 67		
			2 (0) ① 10 ② 10 at 30 sec			㊲ 125 ㊳ 149 $\frac{274}{2} = 137$	㊴ 68 (many small) ㊵ 127 (many small) $\frac{195}{2} = 98$				
									del 10^7 0.5 + 3079 ㊶ 123 ㊷ 139 $262 \times 10^8 = 2.6 \times 10^9$ plaque		
						Total = 1677 Avg $\frac{Total}{6} = \frac{1677}{6} = 279$					
			$600 \times 10^6 \times \frac{1}{3} = 24000 = \frac{2.4 \times 10^5}{2.2 \times 10^8}$ cells = $1 \frac{1}{10}$ cells								
50									$\frac{104}{118}$ $\frac{68}{24.7} = 99$ 99 279 = burst size	$\frac{0.36}{49.0}$ $\frac{277}{23.7}$ 530	



DATE: 2/16/56

REF: 395

1. Cross done 2/13/56 of 300⁺ \times 341-9
 $\frac{1}{2}$ lg^+ cd , lg^+ R TLR = $2/2^- \text{lg}^+$

1. 1.0 ml of each parent added 9.0 ml per, incubated 3 hours, diluted 1-100 with distilled water, 0.1 ml

(+) (-)
 2 \times 7 ca 20
 3 5 ca 300
 4 5 ca 250

10 - volume plate shaded B gel - grew up as
 but on B gel - 10 ^{ca 100} plates to D(0) - all grew therefore
 pathogenic
 The 10 tested against HET 1 - HET 2
 10 + 0 = 602-

3. Out of 20 heads tested

Seq	?	λ	3079	Genotype
1	0	R	+	lg^+
2	0	R	+	lg^+
3	0	R	+	lg^+
4	0	R	+	lg^+
5	+	R	+	lg^+
6	0	R	+	lg^+
7	+	R	0	lg^+
8	0	R	0	lg^+
9	0	R	+	lg^+
10	+	R	+	lg^+
11	0	R	+	
12	0	R	+	
13	0	R	+	
14	0	R	+	
15	0	R	+	
16	0	R	+	

(3+) 16 R 14+ 20

3 seq obtained

	λ	λ	λ
5	+	0	R
7	+	0	R
10	+	?	R

$$2^+ 1^- 5 \times \frac{2^- 1^+ 1^?}{2^- 1^+ 1^?} \rightarrow \frac{2^+ 5}{2^- R}$$

10

20

30

40

50

DATE: 2/17/56

REF: 396

	1	2	3	4	5	6	7	8	9	10
	2637 derivatives from 290									
	labelled - after some arrangement, and re-arrangement									
		Presumed	3079	(number 910)		(Cereulys)				
	W3091	Col ₁ - L ⁺	ne	+	partially S	L ⁺				
	92	Col ₂ - L ⁺	ne	+	R	L ⁺				
	93	---								
	94	Col ₁ - L ⁺	ne	+	R	L ⁺				
	95	---								
	96	Col ₆ - L ⁺	ly. seq.	+	R	L ⁺				
10	97	Col ₇ - L ⁺	ly. seq.	+	R	L ⁺				
	98	---								
	99	---								
	W38 00	Col ₁ + L ⁺	ly. seq.	+	R	L ⁺				
	3101	Col ₁ - L ⁺	ly. seq.							
	3102	Col ₂ - L ⁺	ly. seq.							
	3103	---								
	3104	Col ₁ - L ⁺	---							
	3105	---								
	2106	Col ₆ - L ⁺	partially S.	+	S	L ⁺				
20	3107	Col ₇ - L ⁺	partially S.	+	S	L ⁺				
	3108	---								
	3109	---								
	3110	Col ₁ + L ⁺	partially S.	+	S	L ⁺				

Frequency of transposition among stable (+)

Left by side = K12 8/10/55

7. 2580 spant = 8
o.l.m = 573

2. 3079 spant = 0
o.l.m = 517

3. 750 spant = 3
o.l.m = 356

DATE: 2/28/56

REF: 397

2580 one step using 3077 as indicator
 overnight cult diluted 1-10, aerated 3 hours.

Summed 318 $\frac{0.154}{8}$
 $\frac{149.0}{318}$
 $\frac{172.0}{159.0}$
 $\frac{159.0}{13.00}$

15.4%

PRE-UV

7.6e 10⁶ active
 58
 $\frac{0.0}{10} = 0.002$

148
 170
 318

Pre hand, plaque array, Same plate.

1-10 cu

U.V. 60 records

Add 3 unit 90 use Pen 2K

Post UV

not added to Pen.

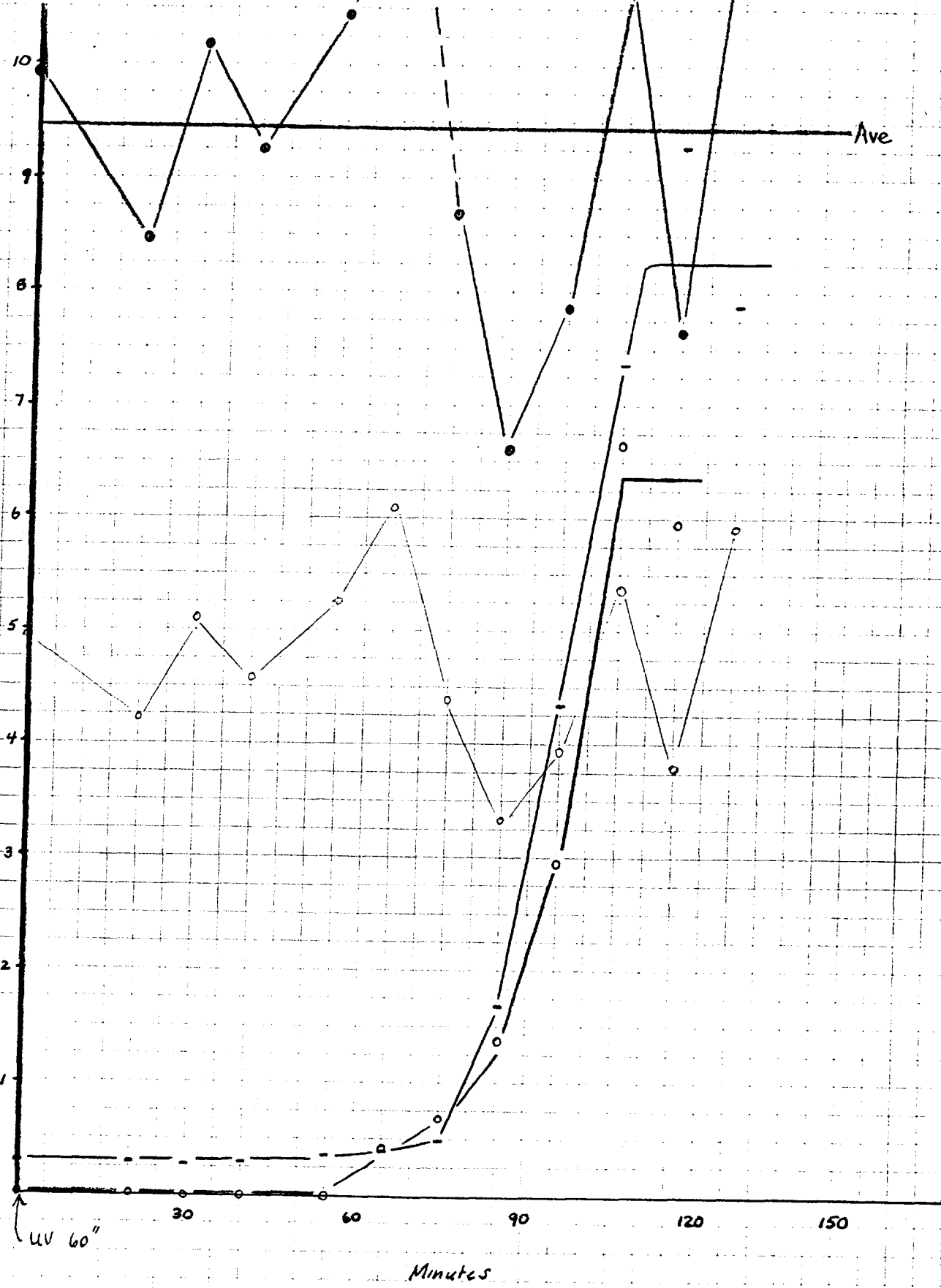
lit 10⁶ - 0.05
 22
 27
 49
 887
 1097
 1984

Summed 58
 64
 61
 1000 plaque
 ca 1000 plaque

Time	Step	Value	Value	Value
20	27	1:30	20	476
	29	56		368
	32	146	30	467
	21	53		552
	27			1019
	38	1:50	40	462
	32			461
	35	60		723
	33	2:05	50 (55)	528
	36	69		517
	35			1045
	31	2:15	60 (65)	515
	38	77		621
	31			1216
	35	2:25	70 75	427
	63	98		441
	49			863
	109	2:35	80 85	296
	22	335		364
	168			660
	452	2:45	90 95	358
	413	865		395
	433			78.3
	772	2:55	100 105	505
	701			566
	737	1473		1067
	960	3:05	110 115	408
	873			350
		1833		753
	873	3:15	120 125	612
	700			303
		1573		1175

163 770

397
2/21/56



Plaque in hundreds
per 0.1 ml. of 10⁶
dilution

UV 60"

Minutes

DATE: 7/22/56

REF: 398

Better transducer indicators?

ly. cell = 367M1 diluted 10⁴ 0.05%
taken

1. Effect of staining. Overweight cultures of 3079, 3080

sedimented \times 1000 in saline sedimented \times 1000 in D(m)
 created 6' hours 0.1 ml used as an indicator. Best under control
 1. Plating Plating Surv Surv Surv Surv Surv Surv Surv Surv Surv
 3079 + 0 0 + 139 (ca) 2 plates grown
 + 0 0 + 158 (ca) 2 plates (6 hours for
 10 + 0 + 31 (ca) 1 plate (probably lower
 + 0 + 24 (ca) 1 plate cell density than
 0 + + 52
 0 + 0 + 164

3080 + 0 0 + 200
 + 0 0 + 222
 + 0 + 65
 + 0 + 149
 20 0 + ? ? 138 } plates
 0 + ? ? 127 } not labeled

These results suggest that stained
 cells may be better as an indicator cell
 densities were not the same and it
 will be better to repeat using a density
 check.

30

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DATE: 2/24/56

REF:

399

One step with 945 (culture from ARC) technique used mentioning on
in 397 with the exceptions.

Summed = $\frac{1}{2} \times 54 = 27$

A. Error

1. Culture of 945 (mal. by P) instead of 2580 (mal. by P)
2. Density of susp. (unadjusted) and worked to 0.25% UV.-60"
3. Diluted 1:10 instead of 1:100
4. Time (absolute) the same - samples taken at 50, 60, 70, 80 etc. instead of 15, 30

(55 ml samples)

10 Time

B. Results

Time	1	2	3	4	5	6	7	8	9	10
0	0	0	0	0	0	0	0	0	0	0
20	6	6	27	27	3	0	0	0	0	0
30	7	7	66	66	2	0	0	0	0	0
40	8	8	47	47	2	0	0	0	0	0
50	9	9	80	80	2	0	0	0	0	0
60	8	8	84	84	2	0	0	0	0	0
70	8	8	93	93	2	0	0	0	0	0
80	10	10	60	60	2	0	0	0	0	0
90	12	12	58	58	2	0	0	0	0	0
100	62	62	88	88	2	0	0	0	0	0
110	74	74	80	80	2	0	0	0	0	0
120	184	184	140	140	2	0	0	0	0	0

Time

3500 SM

Hand counter measured
divide 1-10

130
55 ml samples
plaque
184
145
329
10 = 20

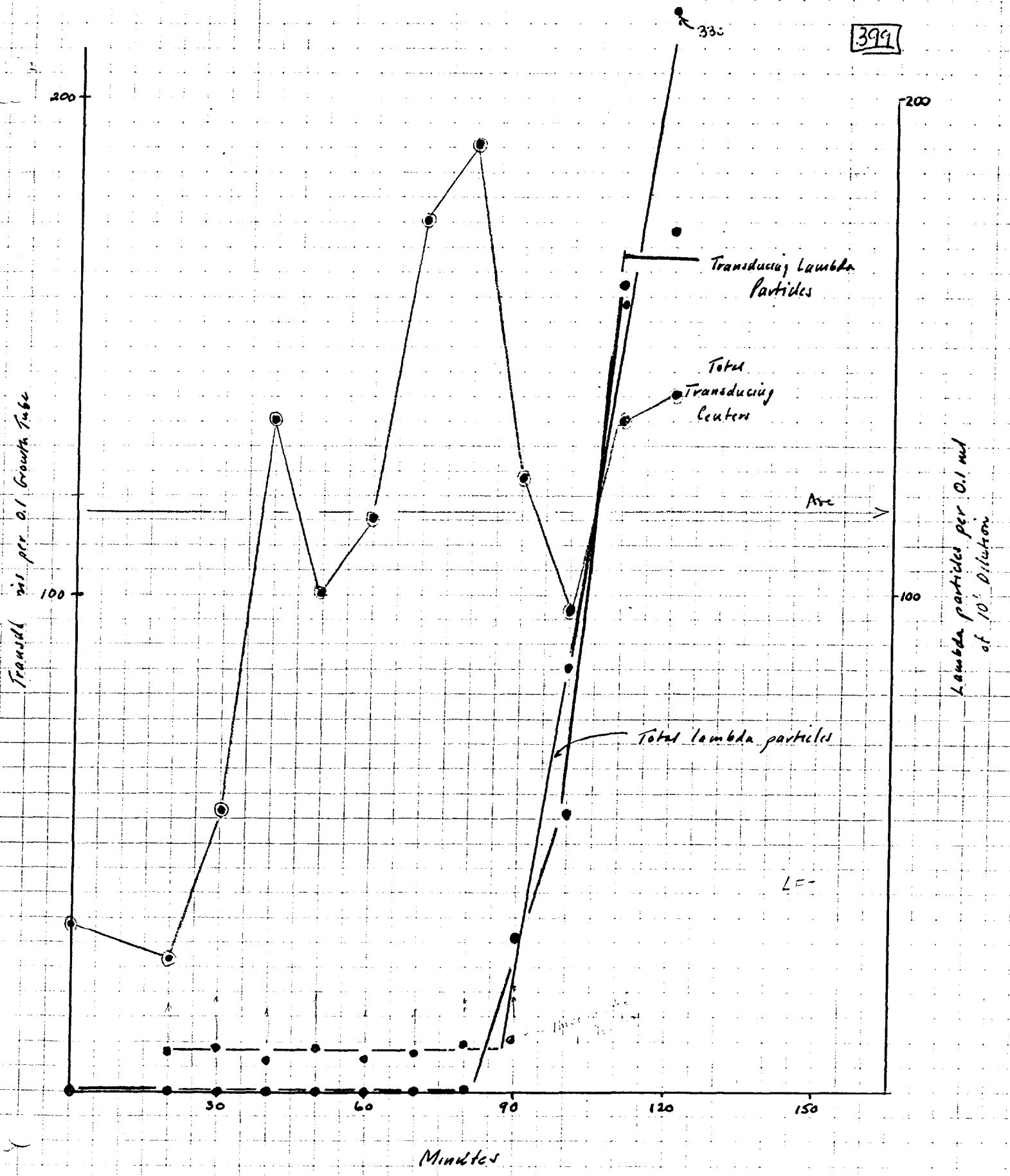
Total
Plaque

17
18
11

Free viral in W3079
since plaque
notified

50

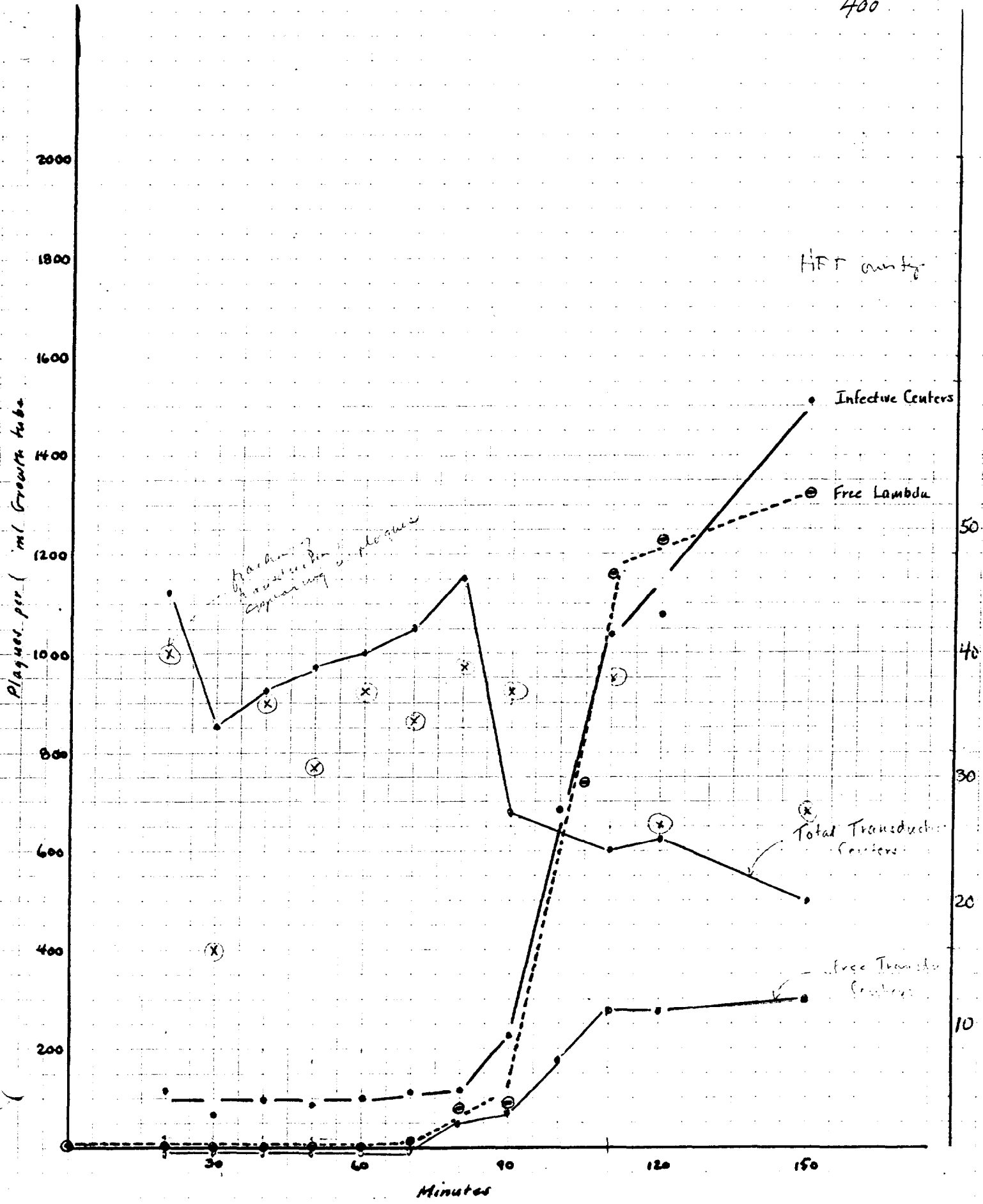
399



DATE: 2/27/56

Survival = $\frac{16}{40} = 0.40$ REF: 400

	1	2	3	4	5	6	7	8	9	10
	One step with 341-9							Plaque count in 9		
	1. Overnight culture diluted ca 1-10, incubated in rotator at 37C for 2 hours. Centrifuged, resuspended volume to 0.05							2. 58		each of 2 of these LFT
	2. Pre assay (A) $10^2 - 10^4 - 10^6 \rightarrow 0.05$ surface for volume							ASSAY THESE COL. FOR HFT \rightarrow		
	3. Inoculate 60 sec							10 LFT		
	12:15	10	$10^2 - 10^4 \rightarrow 10^5$	0.05	surface for HFT	76	76	76	76	
	17:35	20	Inoculate - plates at various time intervals			57	57	57	57	
	12:45	30	Cells disappearing, 40-16 = 24			22	22	22	22	
	12:55	40	Infect. centers produced = 12			23	23	23	23	
	1:05	50	30 = 6			20	20	20	20	
	1:15	60	40 = 10			26	26	26	26	
	1:25	70	50 = 9			12	12	12	12	
	1:35	80	70 = 11			21	21	21	21	
	1:45	90	80 = 11			18	18	18	18	
	1:55	100	90 = 10			22	22	22	22	
	2:05	110				23	23	23	23	
	2:15	120				17	17	17	17	
	2:25	130				13	13	13	13	
	2:35	140				14	14	14	14	
	2:45	150				7	7	7	7	
						12	12	12	12	
						9	9	9	9	
						11	11	11	11	
						7	7	7	7	
						8	8	8	8	
						3	3	3	3	
						4	4	4	4	
						6	6	6	6	
						5	5	5	5	
						7	7	7	7	
						4	4	4	4	
						4	4	4	4	
						8	8	8	8	
						13	13	13	13	



DATE: 3/8/56

REF: 401

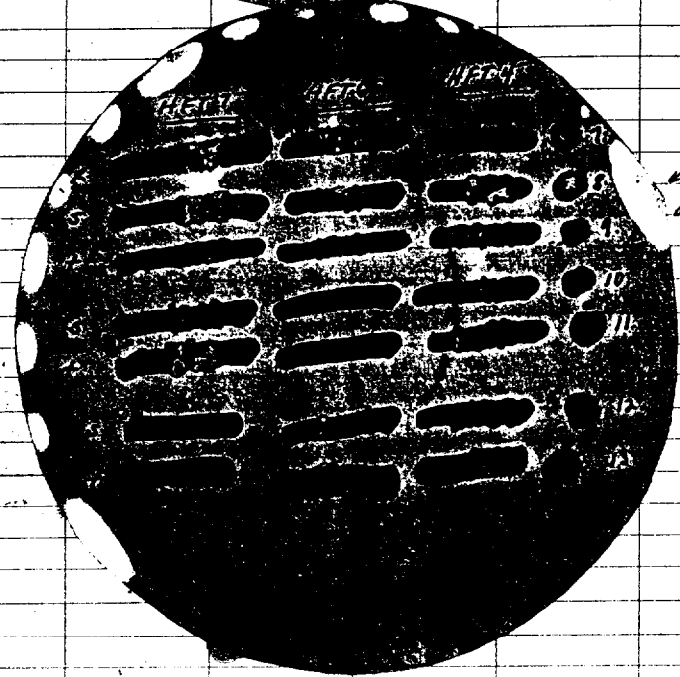
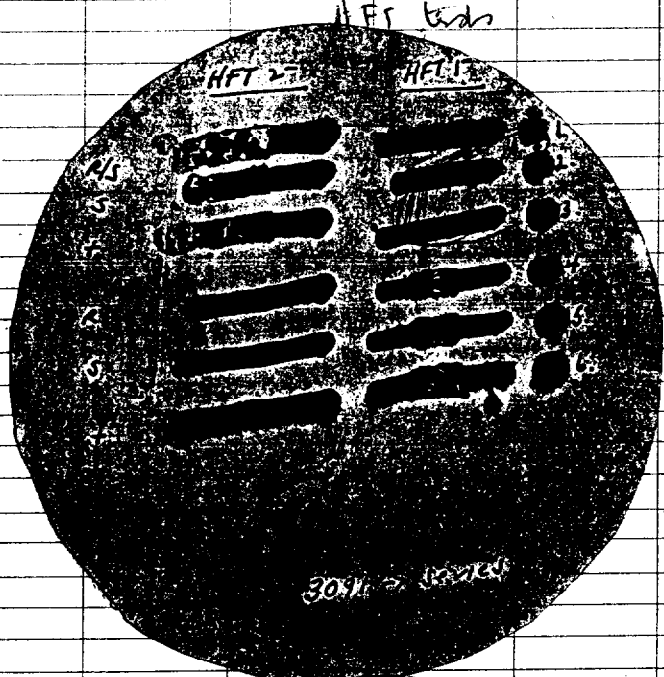
Preparation of pertubosin F stocks and the gal_2^- markers - Kalesha stocks

1. The numbers proposed for these are $3091 \rightarrow 98$, $3101 \rightarrow 3107$, $3110 \rightarrow ?$
 $3100 = gal^+ Gal^+$
 $3110 = gal^S Gal^+$

2. Beginning with 3110 (= F-1491) - the cells from an overnight culture exposed to HFT's and diluted and plated. See page 390

3. After all cultures obtained, they were tested:

3079	λ	gen ⁺ D ₁₀ ⁺	Original gen. description
1. wt	R	+	1-1/1
2. wt	S	+	1-5
3. gal_2	R	+	1-+
4. wt?	R	+	2-+
5. wt	S	+	2-5
6. gal_2	R	+	2-+
7. wt	R	+	4-+
8. wt	S	+	4-5
9. gal_2	R	+	4-+
10. gal_2	R	+	6-3
11. wt	S	+	6-+
12. wt	S	+	7-3
13. gal_2	R	+	7-+
14. wt	S	+	+ 5
15. gal_2	R	+	+ +



Notes

- ① The Gal_2^- derivatives appear to be revertants
- ② The cultures obtained by serial 6- gal_2 (to obtain gal_2^+) appear no longer to be Gal_2^- ; and the gal_2^+ derived is also not Gal_2^-

Redo Gal_2^-
 $Gal_2^- gal^S, gal^+$

THE CULTURES FIRST ISOLATED FOR Gal_2^- and gal_2^- STOCKS #s 4, 5, 6, 7, 8, 9 above given number 401-4, 5, 6, 7, 8, 9 etc

Comparison of lysate from homogeneous 364A1 - a homogeneous obtained in a homogeneous line originating in a 4^+ recipient. If the higher titer of HFT lysates on 4^+ recipients (as compared with 4^- recipients) is due to presence of phage particles entering 4^+ recipients (as HFT homogeneous have been obtained in 4^+ recipients), then there may not be as high on 4^- as on 4^+ when the line originates in 4^- recipient.

1. 6 lysates made starting from single HFT colonies. (see page 10)

used 60 sec., incubated 4 hours.

delivered $10^2, 10^4, 0.05$ ml plated with 3079, 3080, on B gal STT.

lylate	3079 plaque*	3079 Transduct.	3080 Transd.
1	ca. 15	2 (10)	29
2	20	7 (17)	28
3	29	4 (10)	7
4	26	3 (27)	25
5	17	5 (23)	14
6	7	5 (14)	19

It appears that HFT lysates obtained in 4^- recip. this also may be higher on 4^+ recipients

2×10^5

$3.4 \times 10^5 = 3.4 \times 10^6$

$\frac{3.4 \times 10^4}{20}$

1.7×10^3

$\frac{3.4 \times 10^6}{5}$

1.1×10^6 transd

87×10^4

1.4×10^4

1.4×10^6

* plaque assays low because plates were dry

30 Observation that Gal- grow on M bal - tabs made to see what the phage may be.

Comparison - using 1st phage homogeneous, 1st recip. derived from it; 2nd 4^+ recip. derived in M balin

M (unsel)

no growth - crystal on 1st recip. (364A1-9), and a very low no. of this 2nd recip.

M ph 0.01% ph

growth by acc - colonies up to 1 mm across

40 M gal (obtained by plasmid)

growth by acc - 2's better than one - colonies up to 2 mm in diameter

M bal stock

growth by acc 2's better than 1's - colonies up to 2 mm

The Kallikan stocks - Rgt.

3094, 3104 - 3092, 3002 needed.

1. Chicken 390-4 = 4- x 3100 (last 4^d) → heterozygote. This heterozygote not tested directly, but previous segregants found 4^R (see 401) in checker of this ^{sex} gave a non-only 4^d , also evidence of some pleige.

2. Segregant re-obtained from 390-4. 4 picked from first stock and they're closely related.

	HET(+)	HET 4	4^{RX}
1.	+	0	Sex ← 403-1 = 4^d (3104)
2.	+	0	"
3.	+	0	"
4	+	0	"

Sex genotype not expected - indicates that this heterozygote is $4^R/4^d$ - if so when did pleige and sex come from in first step? Certain which genotype of 401-8?

WB3104

20

30

40

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DATE: 3/15/56

REF: 404

~~Production of HFT by irradiation of HFT~~

Irradiation of HFT fams do - lysate approx. titer 2.4×10^6 total / ml - see page 402
 Lysate ml 5 from 364A1. Dilute the lysate, 0.1 ml + 10 ml D(10) in alcohol and
 add 0.1 ml samples to 1.9 ml Pen. Samples (0.05 ml) removed from vials with dilution
 and plated, by grazing on B. sec with indicator

	DOSE	Plaque/3079	Transf/3079	Transf/3080	Plaque 3079	3080	Transf 3079
	1	0	314	13	101 (purple)	104	155
	2	5 sec.	344	34	188	265	56
	3	10 sec.	283	74	208	170	38
10	4	15 sec.	205	83	256	142	68
	5	20 sec.	164	101	371	99	89
	6	25 sec.	151	136	378	98	103
	7	30 sec.	83	128	ca 390	136	133
	8	35 sec.					

Rpt 3/26/56

	1	36	16	4
	2	68	56	0
20	3	14	11	0
	4	26		0
	5	14	16	0
	6	11	8	0
	7	31	32	7

3080
apparently
contain.

Reason for
the variability unknown.

These lysates have been exposed to
a variety of conditions - including
moving, etc. storage at room temp,
refrig, etc.

40

50

400

300

200

100

Transductions
on
1st Receptor

2nd Receptor
1st Receptor
1st Receptor

Transductions
on
2nd Receptor

Plaque

Seconds UV

5

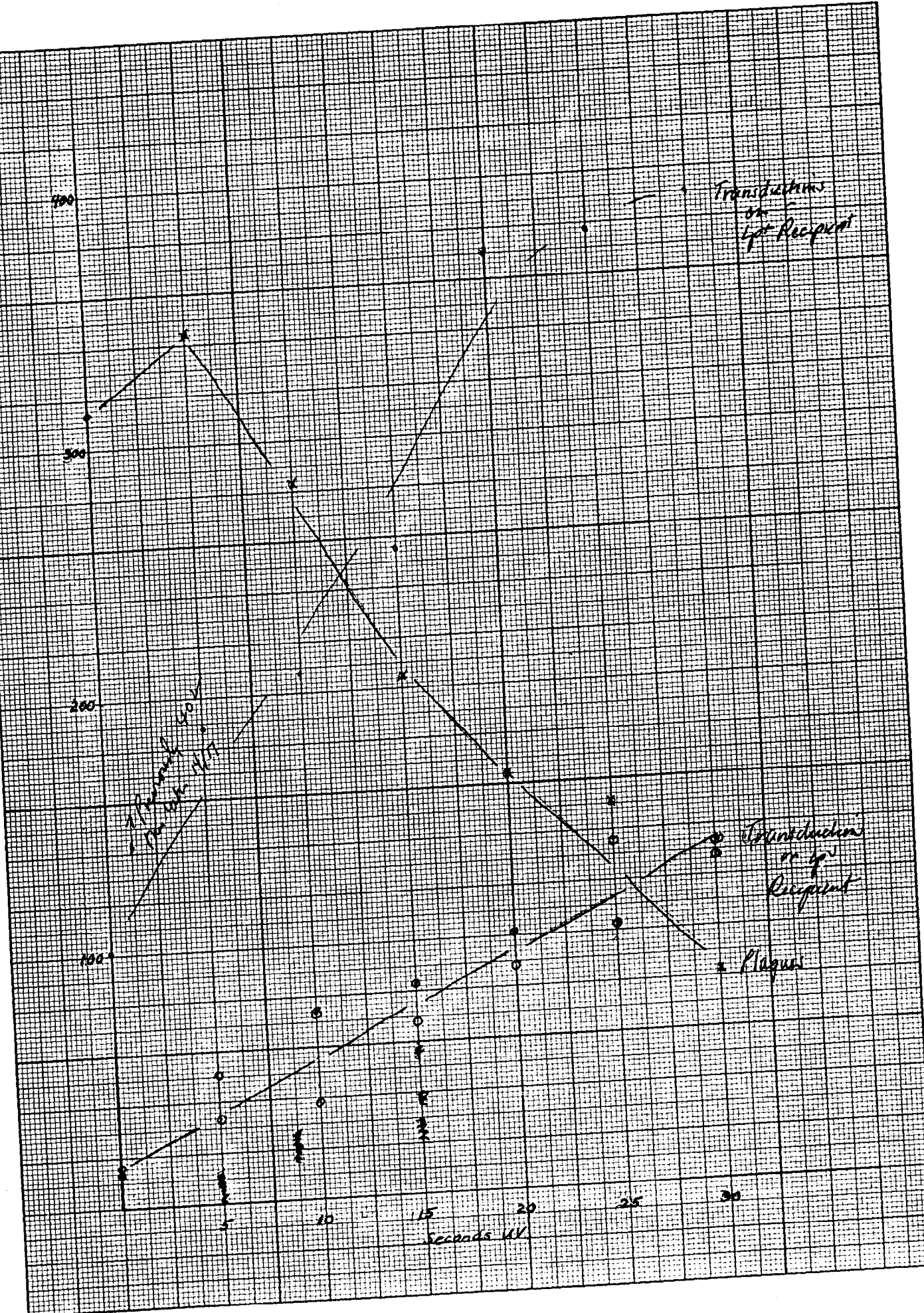
10

15

20

25

30



5 x 10⁵ 1 x 10⁵
3 x 10⁵

DATE: 3/16/56

REF: 405

Production of Transducing λ - LFT - Dura gel study.

2. 945 grown overnight on acetate - dil. 1:4, re-inoculate 2 hours - Centrifuge, resuspend in saline. Adjust density to 0.58 dilute 1-20, inoculate into UV-5000.

Undiluted B_{gal} dilute 10¹, 10² plate

3079 3079 3079

0.05⁵ in Controls

Time	1	2	3	4	5	6	7	8	9	10
10	1. 945	15	30	45	60	75	90	105		
	2. 3079	15	30	45	60	75	90	105		
	3. 3079	15	30	45	60	75	90	105		

Plaque assay

These 100 x 2 x 10⁵

Ppt. undiluted - overnight, diluted 3 + hours after 1-10, centrifuged, resuspend. saline, O.D. = 0.6

3/28/56 dilute 2/9 - procedure as before - remove 0.05 ml + 4 cells, 10² → 0.05 for diluted; 0.05 as before for undiluted - Numbers as above.

Time	1	2	3	4	5	6	7	8	9	10
30	1. 945	15	30	45	60	75	90	105		
	2. 3079	15	30	45	60	75	90	105		
	3. 3079	15	30	45	60	75	90	105		

Plaque assay

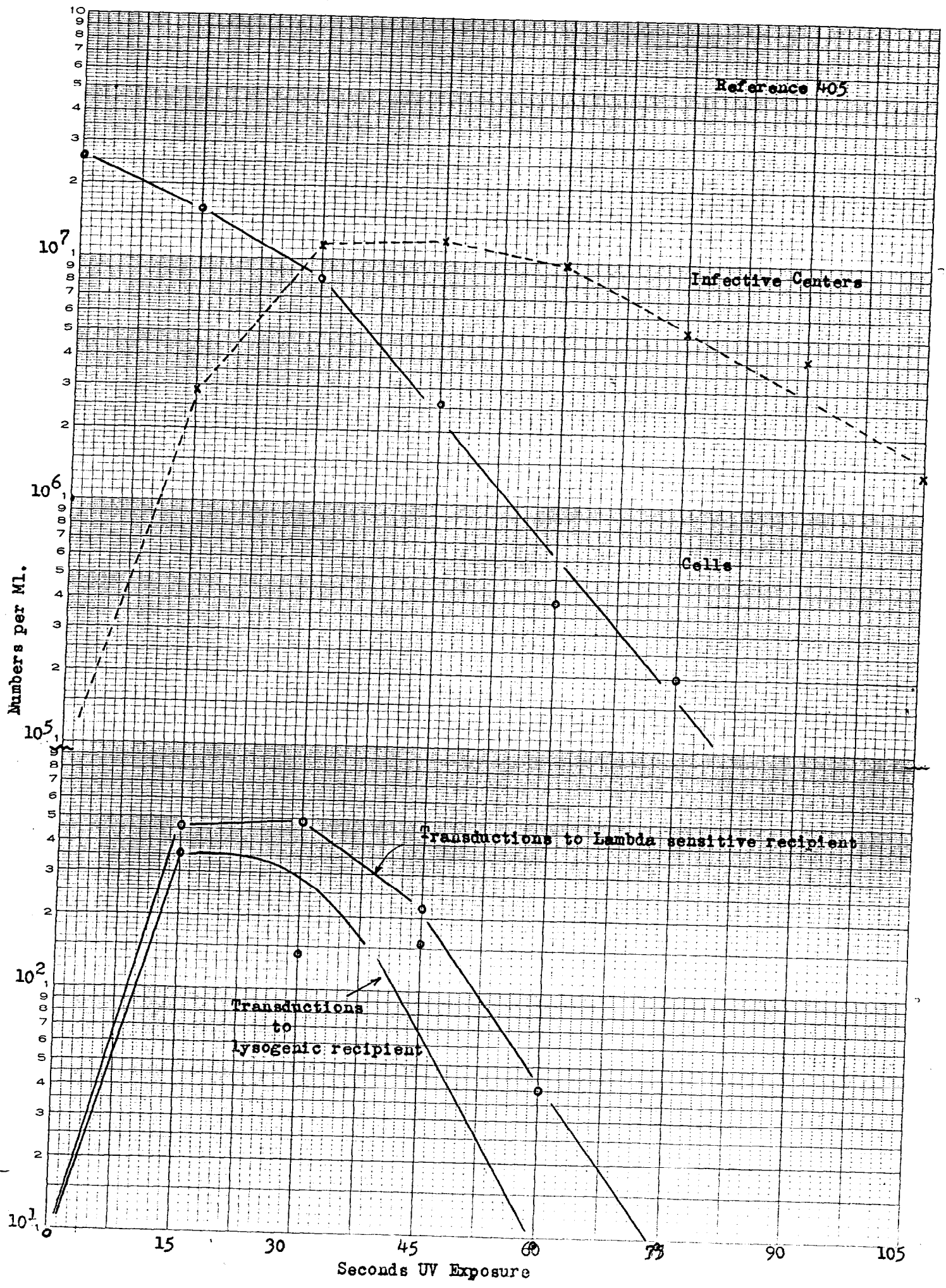
These 100 x 2 x 10⁵

Fraction of cells yielding transductions

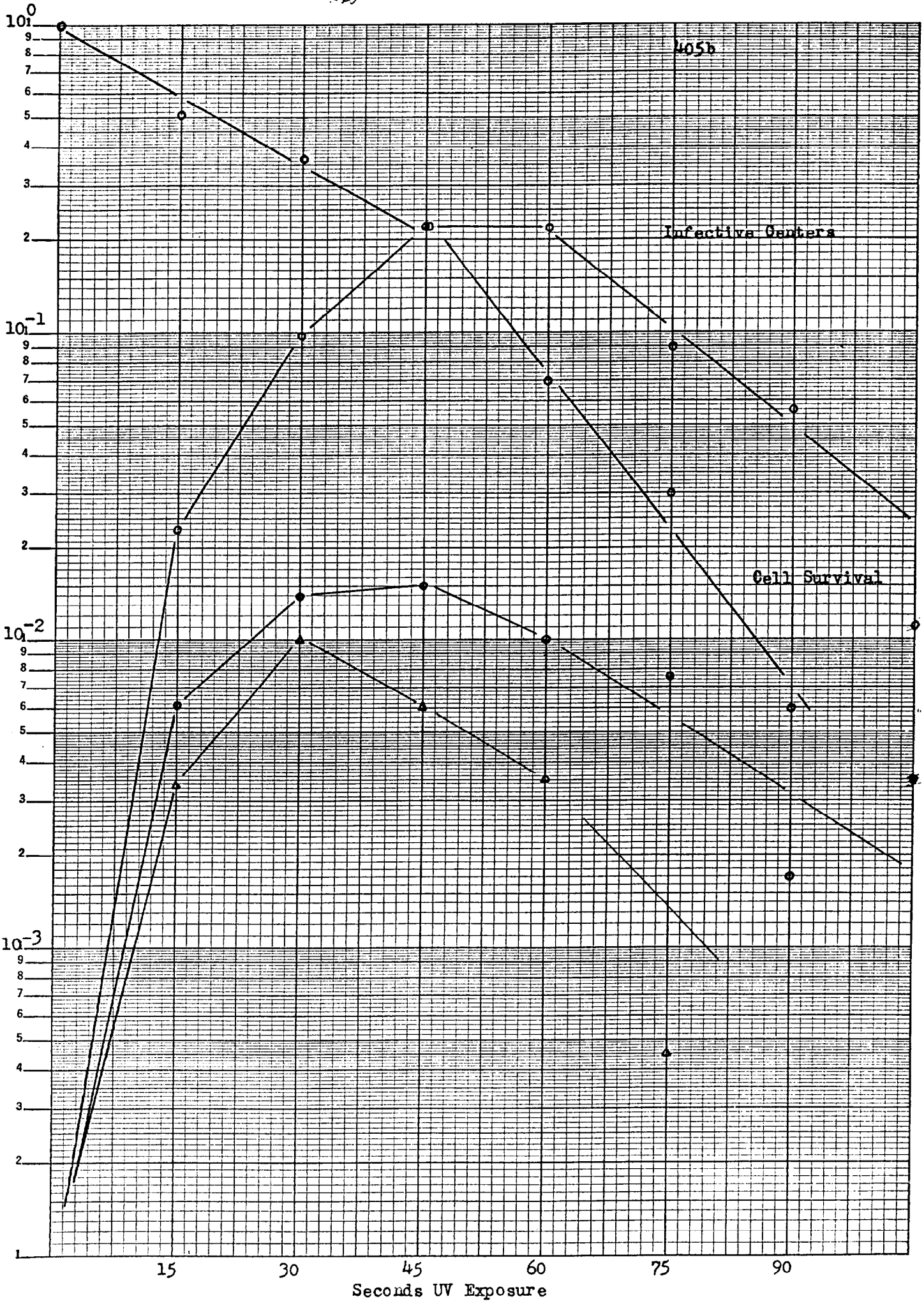
4 x 10⁵ 20 hand
in cell = 472 x 10³
3.1 x 10⁸

NO. 340-LS12 DIETZGEN GRAPH PAPER
SEMI-LOGARITHMIC
5 CYCLES X 12 DIVISIONS PER INCH

EUGENE DIETZGEN CO.
MADE IN U. S. A.



1652



KEUFFEL & ESSER CO., N. Y. NO. 348-81
Semi-Logarithmic, 4 Cycles X 10 to the Inch, 5th lines accented.
MADE IN U. S. A.

Comparison of U.V. Resistance of haploid and heterogametic clone.

1. Set up. 341-9, a 2/2- leucogenetic streaked out, and a LFT segregant obtained. A galt reversal of this segregant by selection on B gal. The comparison for U.V. survival is between these segregant and 341-9
2. Experimental.

① 341-9 streaked from HFT + clone (detected on Bgal-HFT test)
341-9(+) from single colony.

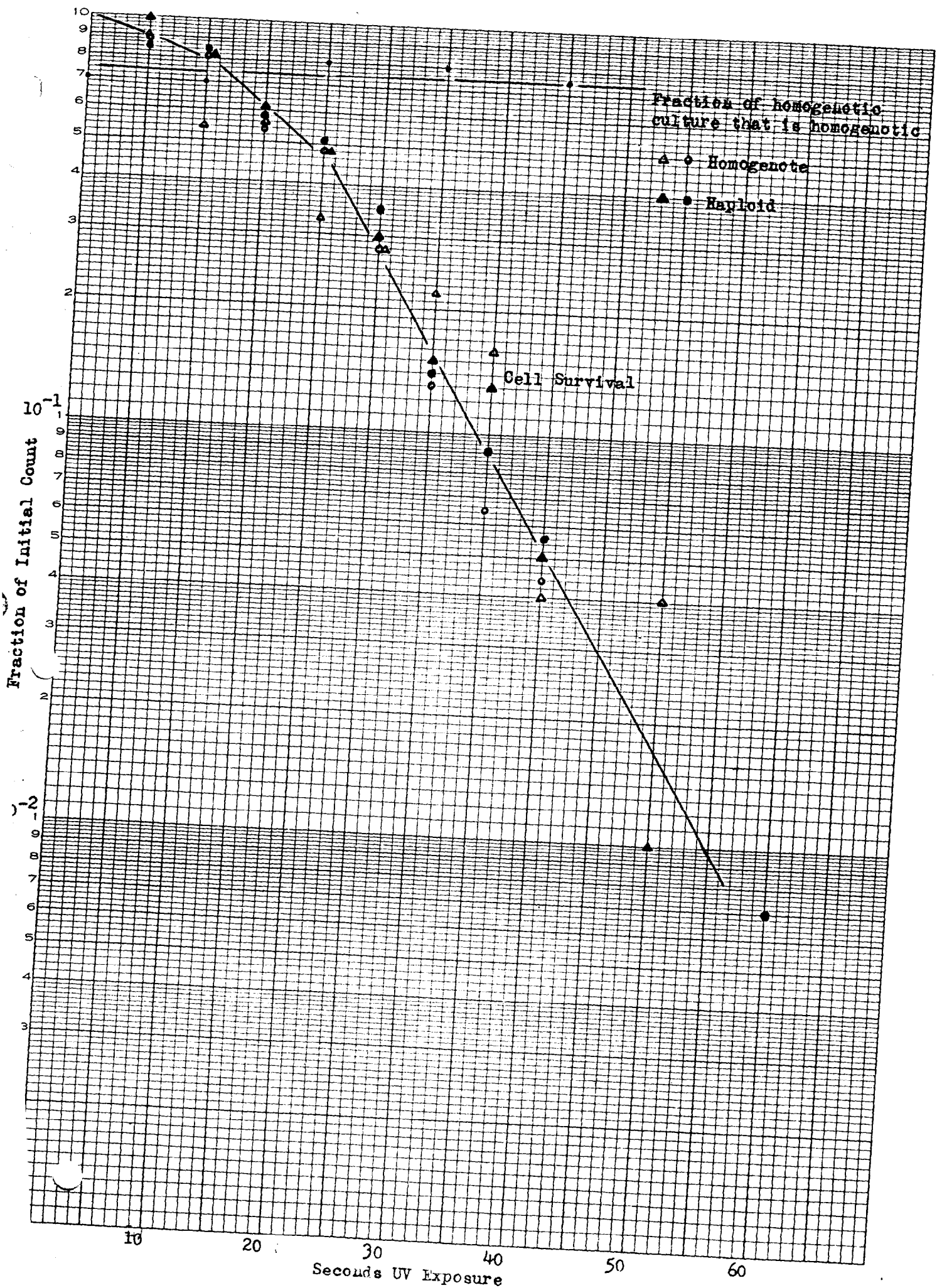
② Overnight cultures diluted ca 1:20 incubated 17 in rotator for ca. 3 hours. Diluted 0.2 + 10 ml solvent - each into the first tube, and then 10³, 10⁴ additional. The first tube incubated, and samples, 0.05 ml taken out and plated on B gal

Time	Galt colonies:		Total	Fractn	Gal - colonies:		Total	Fractn	Fraction of Gal - Colony HFT + / 3080	
	①	②			①	②			14/20	0.7
0	226	234	460	1.0	147	129	276	1.0	14/20	0.7
5	193	214	407	0.89	116	119	235	0.85	14/20	0.7
10	173	203	376	0.82	116	117	233	0.84	14/20	0.7
15	138	110	248	0.54	66	95	161	0.58	—	—
20	110	111	221	0.48	76	64	140	0.51	14/20	0.7
25	55	73	128	0.28	53	43	96	0.35	—	—
30	38	22	60	0.13	28	11	38	0.14	14/20	0.7
35	16	14	30	0.06	17	8	25	0.09	—	—
40	4	16	20	0.04	5	10	15	0.06	15/20	0.75
60	0	0	0	0	1	1	2	0.007	1/2	0.5

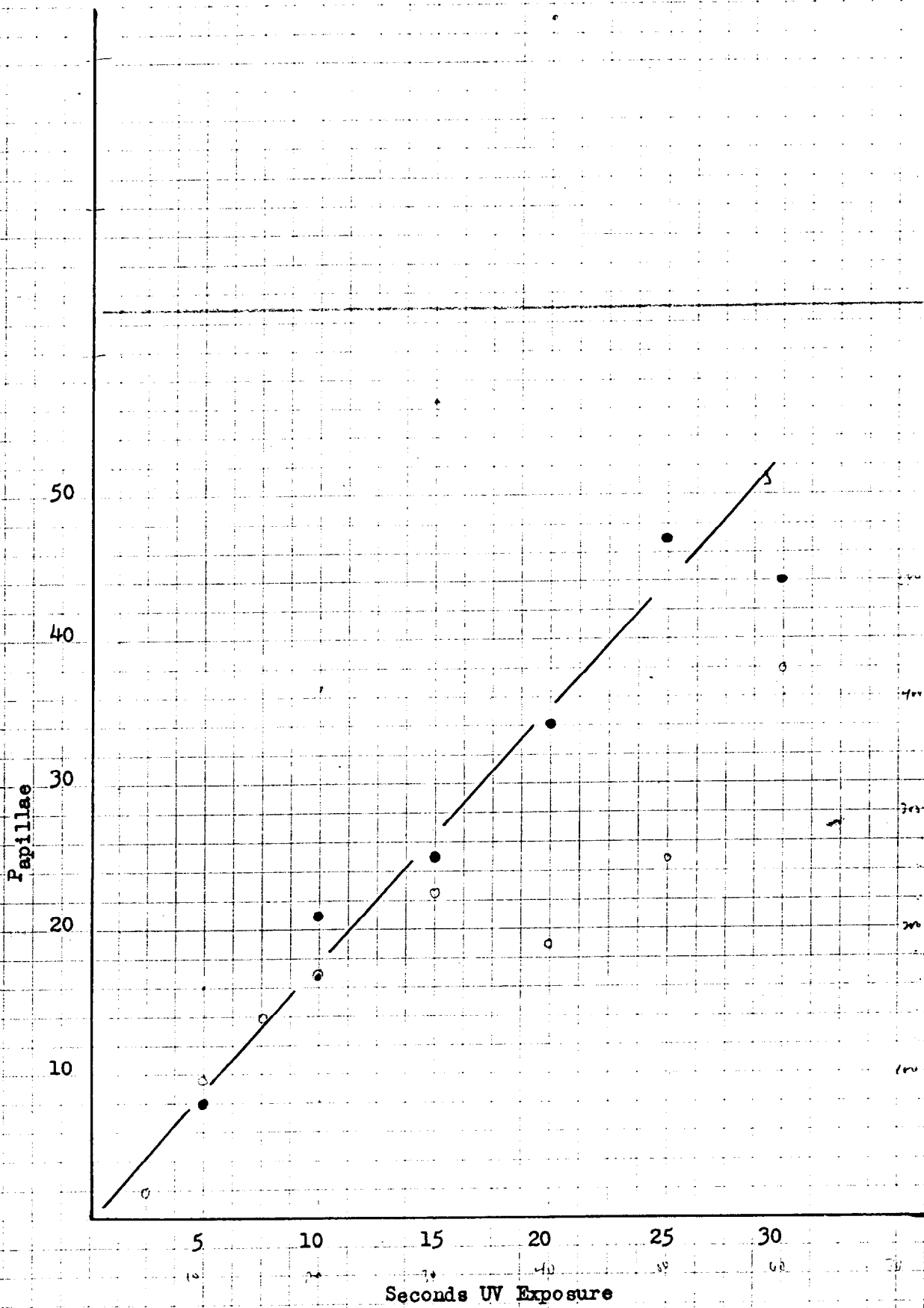
A second analysis (3/4/56) Using a +^R of 341-9 and comparing with a (-) segregant of this

Time	Galt colonies:		Gal - colonies:		In the HFT + culture, the population of +/- was:	
	Total	Fractn	Total	Fractn	1	2
0	325	1.0	151	1.0	2.15	0.47
5	296	0.9	152	1.0	1.95	0.52
10	176	0.54	124	0.82	1.42	0.7
15	179	0.55	92	0.61	1.95	0.52
20	108	0.33	72	0.48	1.5	0.67
25	90	0.28	45	0.30	2.0	0.50
30	72	0.22	22	0.15	3.1	0.32
35	53	0.16	20	0.12	2.7	0.37
40	14	0.04	8	0.05	1.75	0.57
50	13	0.04	2	0.01	6.5	0.16

23/31
19/3
42/4



1.500/20



Tst of conditions required to obtain "maximum" induction in B gel

$$0.58 = 2.6 \times 10^8 \text{ cells/ml} \times \frac{2.0}{15 \times 10^8} \times 0.64$$

A ① Overnight aerated 945 - diluted ca 1-20 incubated in aerated ca 3 hours, centrifuge, resuspend volume. Adjust density to 0.6

② Susp A in ad. 45 reads should yield count in 500 hand / ml

③ Test - read 45 counts, plate 0.05 + 30⁵⁰ 79

- ① 77
- ② 49
- ③ 63

④ Plate 0.05 ml of suspension on surface of B gel - incubate varying doses, add 0.1 ml of 30⁷⁹ resuspend

- 30"
- 45"
- 60"
- 75"

- ③ 99
- ④ 124
- ⑤ 115
- ⑥ 117
- ⑦ 86
- ⑧ 38
- ⑨ 57
- ⑩ 54

112
116
62
16
Average of these
continued?

⑤ Plate 0.05 ml of suspension on surface of B gel - incubate 2 hours - incubate 60", add 30⁷⁹ resuspend
Wash two plates dilute and assay
for total growth - add water 1.0 ml
plate, under water 10 min
once, 0.5 ml obtained, added to 10 ml
dil 10², 10³ → 0.05 ml

- ⑪ lost
- ⑫ ca 3000
- ⑬ ca 3000
- ⑭ ca 3000
- ⑮ —
- ⑯ —
- ⑰ 27
- ⑱ 34
- ⑲ 34
- ⑳ 46

Reagent
with respirator
preserved may
have been
contaminated

$$20 \times 10^4 \times \frac{1}{2} \times 20 \times 10^6 = 170 \times 10^8 = 1.7 \times 10^8 \text{ cells/plate}$$

$$1.6 \times 10^8 \text{ cells/plate}$$

21 = 30
22 = 40

B

Also broad	Dose	0	5	10	15	20	25	30	35
change	population	5	13	26	30	39	52	47	28
response	counts	0	8	21	25	34	47	47	22
v. x									
known									

C

② incubation
ca 60 hrs ca 1-10 } dose 45", + 0.1 ml 30⁷⁹ = 7
" 1-100 x 10⁷ } = 0 (no detect plaque)

DATE: 4/1/56

REF: 408

$$\frac{.12}{4.8 \times 10^7} \times 10^{-3} = 4.8 \times 10^{-11}$$

$$\frac{0.1 \times 10^8}{2} = 5 \times 10^6$$

945 for response to UV - linearity vs. suspended

~~Adapt~~ - Fresh culture - 3-4 hours post start of the bench several days
 Adjust density to 0.5×10^8 saline @ dilute 1.2×10^{10} and plate for count.
 response. (0.6 - at 0.15000 per) @ count $10^1 - 10^3$ for plaque-producing
 response - This last, used. ($1-20 \times 10^2$) and then add oil to 1.5×10^8 per.
 Inoculate samples about 30 min before plating.
 Inoculum = 3.0×10^9

10

Time	Mix 1	Count	Count	3. Inoculum 7 d	Plaque	Too
0	1	14	0	Time	10	0 15
5	2	28	14	given	11	143 158
10	3	26	12		12	349 364
15	4	24	12		13	549 564
20	5	28	14		14	816 831
25	6	25	11		15	0.1:25.87 2.1:12.67
30	7	43	29		16	1661
35	8	39	25		17	1654
40	9	120	106		18	too many ca 1000

20

30

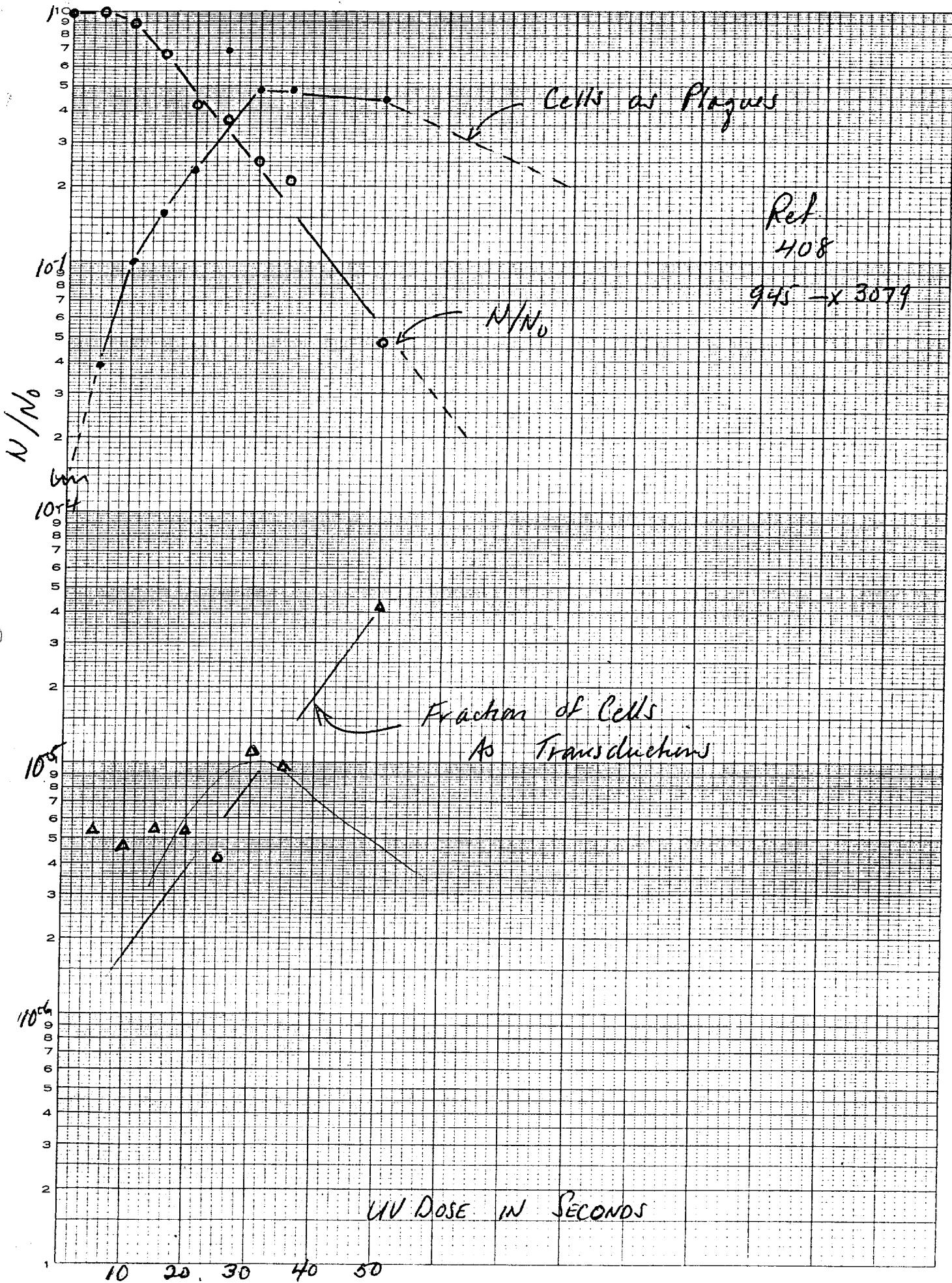
Transduction = $0.1 \times 10^8 \times \frac{11.2}{1.2} = 100 = 3.5 \times 10^4 \times \frac{7.5}{0.1} \times 0.2 = 3.5 \times 75 \times 10^8 = 2.6 \times 10^9$ cells/ml

Fraction of Cells co Transd.	Transd	FRACTION OF CELLS AS PLAQUES	Time	T. Plaque	N/N ₀	N/ml
0	0	0	0	0	1.0	3.5×10^4
5.4×10^{-6}	1.4×10^3	0.039	5	1.4×10^3	1.0	3.6×10^4
4.6×10^{-6}	1.2×10^3	0.099	10	3.5×10^3	0.9	3.2×10^4
5.4×10^{-6}	1.4×10^3	0.155	15	5.5×10^3	0.68	2.4×10^4
5.4×10^{-6}	1.4×10^3	0.230	20	8.2×10^3	0.42	1.5×10^4
4.2×10^{-6}	1.1×10^3	0.700	25	2.5×10^4	0.37	1.3×10^4
1.1×10^{-5}	2.9×10^3	0.48	30	1.8×10^4	0.25	9.0×10^3
9.6×10^{-6}	2.5×10^3	0.48	35	1.7×10^4	0.21	7.4×10^3
4.2×10^{-5}	1.1×10^4	0.45	50	1.6×10^4	0.048	1.7×10^3

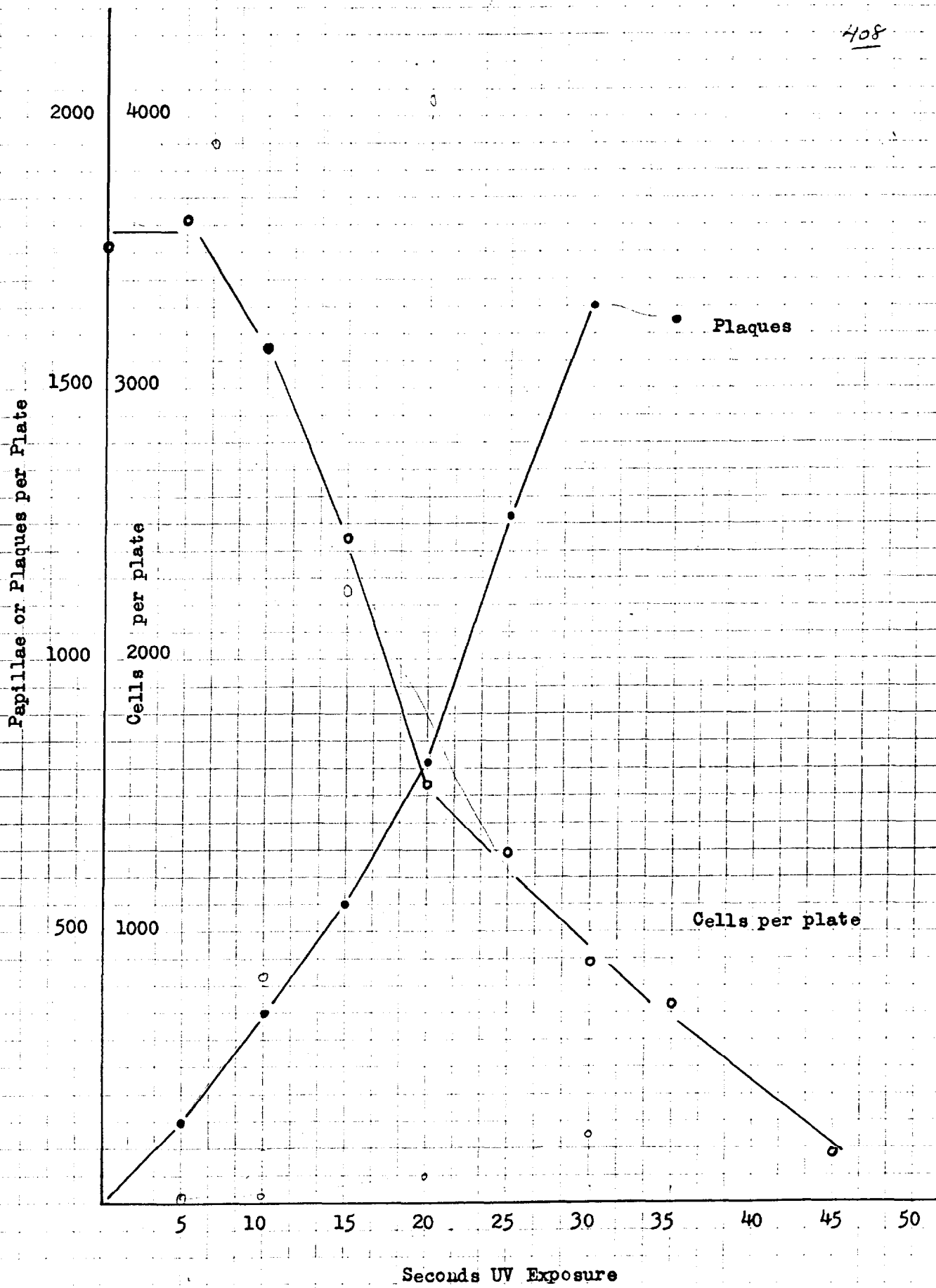
Fraction of
 cells yielding
 transducing
 cells =
 $\frac{2.9 \times 10^3}{2.6 \times 10^8}$

40

50



408



DATE: 4/3/56

REF: 409

Handwritten scribbles in the top right corner.

1 2 3 4 5 6 7 8 9 10

Examination of 4- x 7- heterocysts

1. 4- x 7- Overnight culture diluted 10² 0.1 ml + 0.2 ml lysate.
293-12 U2308 Incubated from temp for 10 minutes, diluted 10², 10⁴
lystate 0.1 ml plated 3 plates. P gel.

A

- 1. 31 (?) (1)
- 2. 37 (5) (0)
- 3. 29 (+) (0)

} 3 additional possibilities appeared second day - 3/5

10

3A

5/7/56 11 (-) seg obtained from the intermed " + " typed - 11/18 Galy -
12 (-) seg obtained from the intermed " + " typed - 12/12 Galy -

2. 7- x 4- No above - 1st row no population ~ semi (+) colonies observed.
2nd row - 2 presumptive observed - slight population in the centers of two colonies noted. Plated and showed total colonies ca 40, all phage contained.

20

C

30

40

50

DATE: 4/3/56

REF: 410

One step with W945, against 3077

1. From overnight culture, diluted 1:10 into 3 tubes
2. Adjust Salini mag. to 0.0. - 0.6, dilute 1:10 into 3 tubes
dilute 1:5; incubate 60 seconds, (Use 1-2 with broth (Per) 3+9)
3. Pre assay. 53

In a different step at this time approx. 300 colonies 945 tested, all found inducible

Cells

7. 38 } 7.4x10⁸ cells
 2. 38 } in OD=0.53 suspension
 3. 0 }
 7. 0 } no indicator?

10

2:15

20

30

40

50

TIME	1	2	3	4	5	6	7	8	9	10
2:15	5	0								
2:20	6	0								
2:25	13	20		15	7	14	17	0		
2:30	14	20		16			18			
2:40	17	23					21	0		
2:45	19	19		20	20		22			
2:50	25	33		26			27	0		
2:55	27	28		28			29	0		
3:00	29	25		31	13	13	32	0		
3:05	30	28		32			33	0		
3:10	35	29		37	7	14	37	0		
3:15	38	29		38			38			
3:20	41	25		40	1	5	41	1		
3:25	42	19		44	4		42	0		
3:30	47	75		47	4	13	51	2		
3:35	48	83		50	6		52	2		
3:40	53	130		55	2	4	57	8		
3:45	54	141		56	2		58	4		
3:50	59	238		61	9	21	63	5		
3:55	60	238		62	33	supra	64	9		
4:00	65	313		67	14	supra	69	10		
4:05	66	326		68	38	supra	70	20		

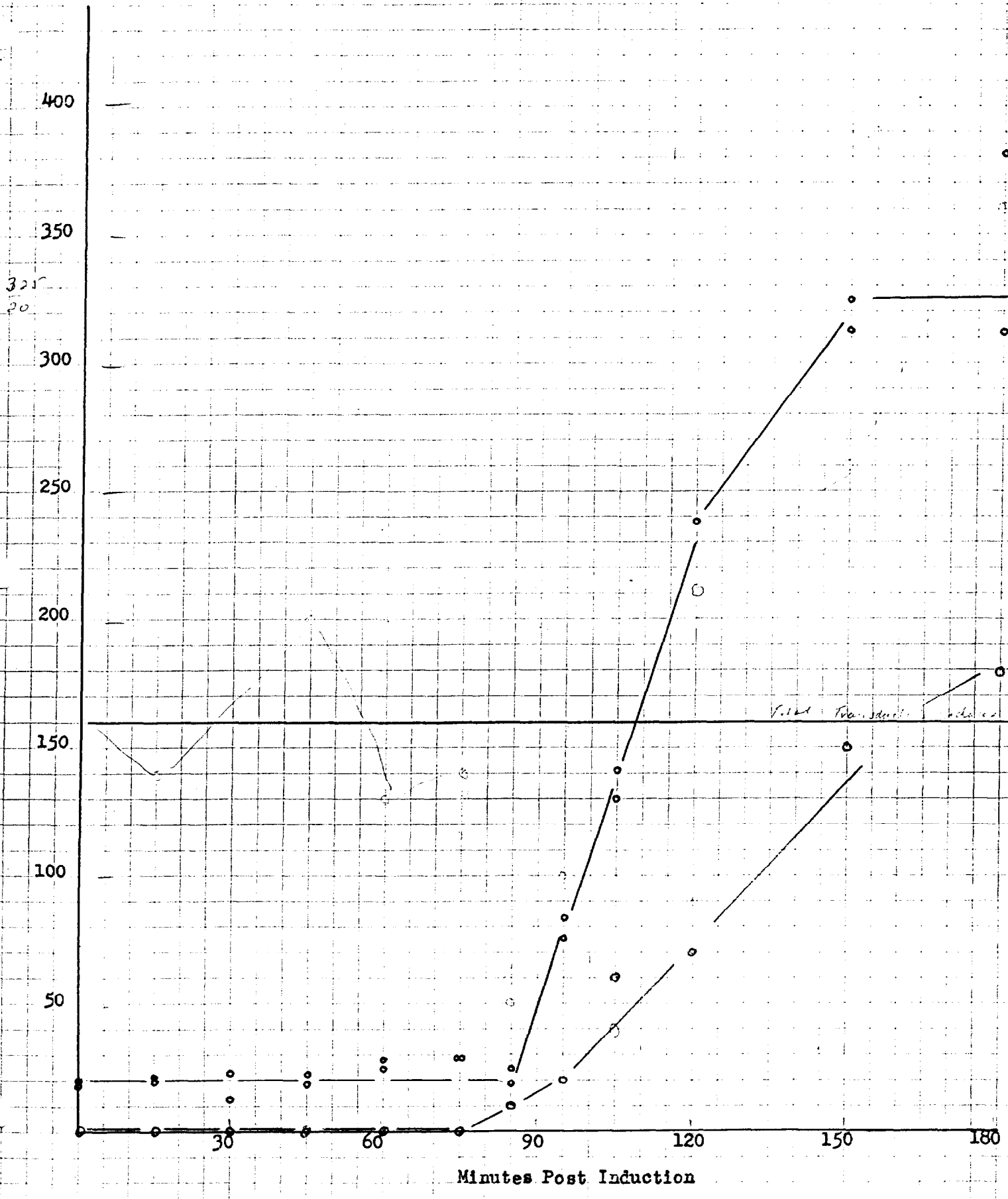
Franklin's cells yielding around 3x10⁸ = 7.4x10⁸

Meat table hand

16	16
17	7
18	20
19	13
20	7
21	4
22	5
23	2
24	21
25	26
26	38
27	157
28	14

These were all on force plate

100	71.38v	72	37	26
	72.3v	74	36	
		75	19	18
		76	17	



Gal + 4/5

Gal - 4/5

DATE:

REF: 411

UV. Resistance 4/5 heterogene 10 & 20.
 W286P taken - and a 4/5 seg of it. On taking against 1, 4p
 the segregant found 4/5 as well as the heterogene

Both cultures diluted ca 1-10 incubated 2 hours, diluted 50X 50X 50
 in saline. The heterogene just, 0.05 ml to 2 8 gal plates to obtain Gal+/Gal-
 ratios. No seg diluted in same tubes. The last tube incubated
 and samples taken (25) at the following times.

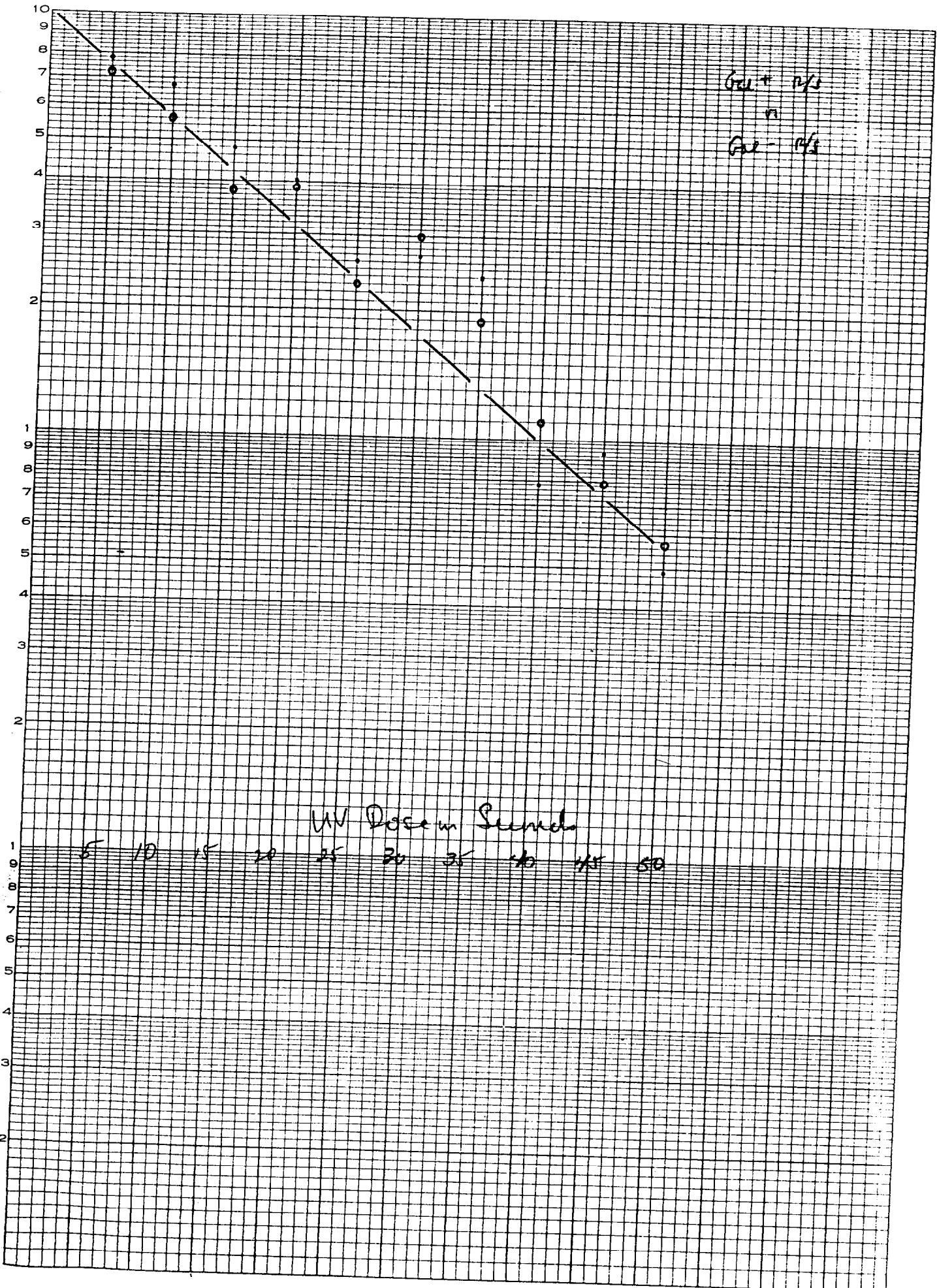
	Time	Gal+	Gal-	Fisher Gal+	Fisher Gal-	Ratio Gal+/Gal- in 286P	Gal+	Gal-
10	0	600	204	1.0	1.0	1.	229	10
	5	431	158	0.72	0.78	2.	259	1
	10	336	139	0.56	0.68		488	11
	15	228	99	0.38	0.49			
	20	236	81	0.34	0.40			
	25	138	54	0.23	0.26			
	30	178	55	0.30	0.27			
	35	115	48	0.19	0.24			
	40	66	16	0.11	0.078			
20	45	48	19	0.08	0.093			
	50	34	10	0.057	0.049			

30

40

50

Gen. 1943
n
Gen. 1943



UV Dose in Seconds
5 10 15 20 25 30 35 40 45 50

DATE: 3/10/56

REF: 412

One step with 945

Overnight culture diluted 1-5 incubated ca. 1 hour.

Centrifuge, resuspend in saline OP. 60

(~~Adapted~~), moderate 60 seconds, dilute 1-2 with 2x pen

Assay 1. Pre UV. Dilute $10^2, 10^4, 10^5, 10^{10}$

Cults
 (1) 163
 (2) 186

Bgal

1.75×10^6

2.50×10^6

2.5×10^8 cells/ml

OP 0.1

10

und

Plaque
 Found

(5) 0 ca 1000 plaques
 (7) 0 ca 1000 -

4.2×10^3

3.5×10^8 cells/ml

Post UV - dilute $10^2, 10^4, 10^5$

Per incubate

Unad. incubated

Bgal

Bgal SM

9:25

Optima

Cults
 (5) 0
 (6) 0

(9) 104 103
 (10) 101

(11) 1 ca 1000 plaques
 (12) 0 " " "

no cells washed
 to yield transd
 out = $210 \times 20 \times 2 =$
 0 = no growth
 + = growth

20

Ascetics
 (2) 42
 (8) 32

4:40

15, 13, 43
 (19) 40

(15) 85 11.7
 (16) 150

(17) 0 ca 1000 plaques
 (18) 0 ca 1000 plaques

Some preparation checked
 45 75

9:55

30, 21, 56
 (20) 56

(21) 72
 (22) 98

(23) 0 ca 1000 plaques
 (24) 0 " " "

30

45, 37, 47
 (25) 37
 (26) 47

(27) 113
 (28) 112

(29) 0 ca 1000 plaques
 (30) 0 ca 1000 plaques

10:25

60, 37, 39
 (31) 37
 (32) 39

(33) 93
 (34) 124
 AVE = 105.2

(35) 0
 (36) 0 (10^5 plaques)

10:45

75, 53, 57
 (37) 53
 (38) 57

(39) 88
 (40) 127

(41) 2 ca 1000 plaques
 (42) 5

10:55

90, 54, 64
 (43) 54
 (44) 64

(45) 80
 (46) 67

(47) 31
 (48) 24

40

100, 99, 143
 (49) 99
 (50) 143

(51) 71
 (52) 118

(53) 29
 (54) 46

11:15

100, 357, 448
 (55) 357
 (56) 448

(57) 59
 (58) 81

(59) 67
 (60) 78

11:25

120, 1024, 930
 (61) 1024
 (62) 930

(63) 87
 (64) 65

(65) 91
 (66) 124

11:55

150, 1442, 937
 (67) 1442
 (68) 937

(69) 181
 (70) 161

(71) 150
 (72) 149

12:25

180, 1422, 1196
 (73) 1422
 (74) 1196

(75) 203
 (76) 174

(77) 193
 (78) 208

412
945-X-3579

200

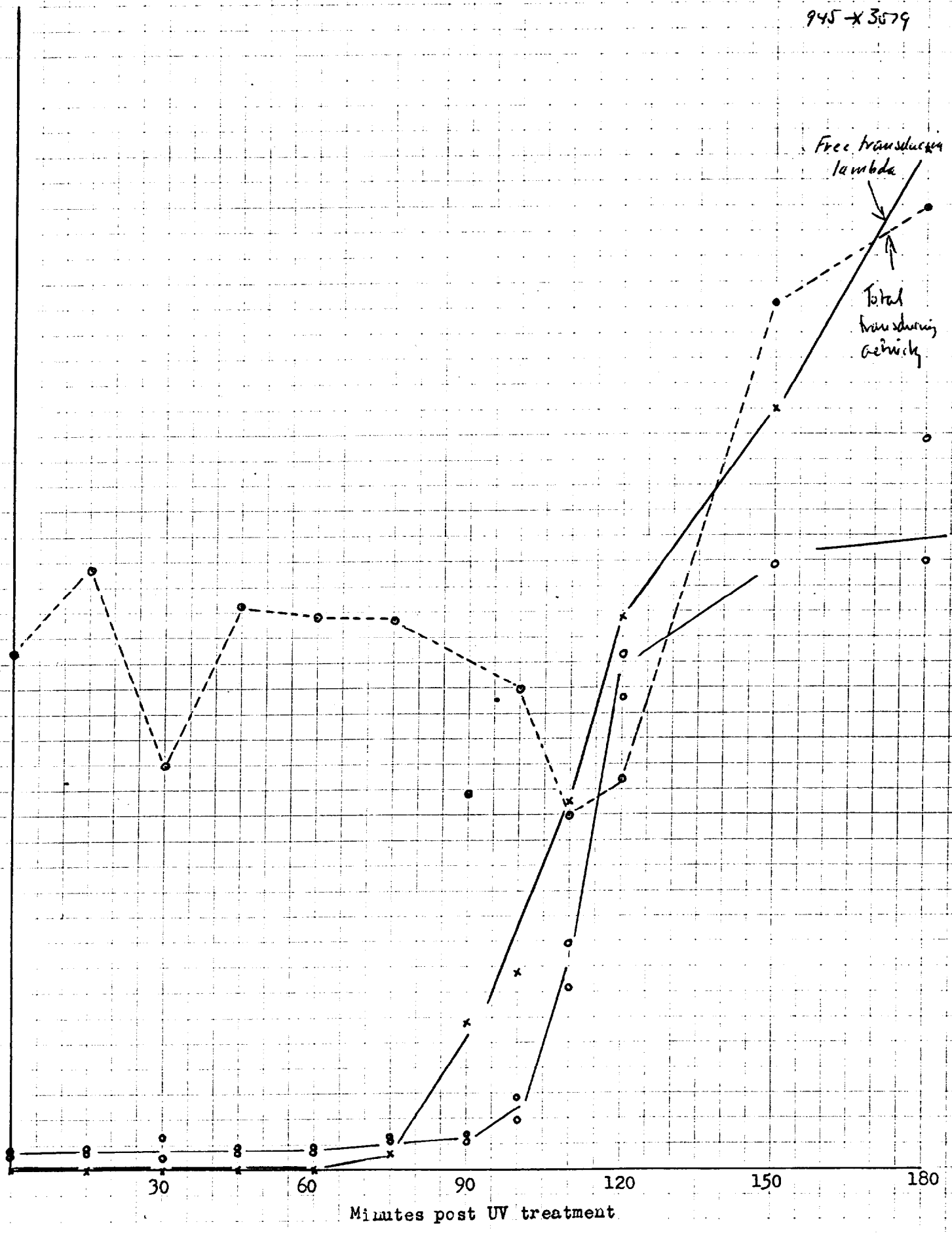
150

100

50

Free transducer
 λ

Total transducer
activity



Minutes post UV treatment

DATE: 4/14/56

REF: 413

Recheck on induction - as plaques - is it exponential?
 - as transmission - ... linear?

1. Plaque induction - 945 rotated 2.5 hours post overnight diluted 1-10. Cells frozen made up to D.O. 950 in saline - diluted 1-2, ^{10⁵} in adjusted CV standard dist, etc., assay for cells: 8 fold (pura method) ^{2.0 x 10⁸}
 for plaques: 2 fold + 3079, also pura method.

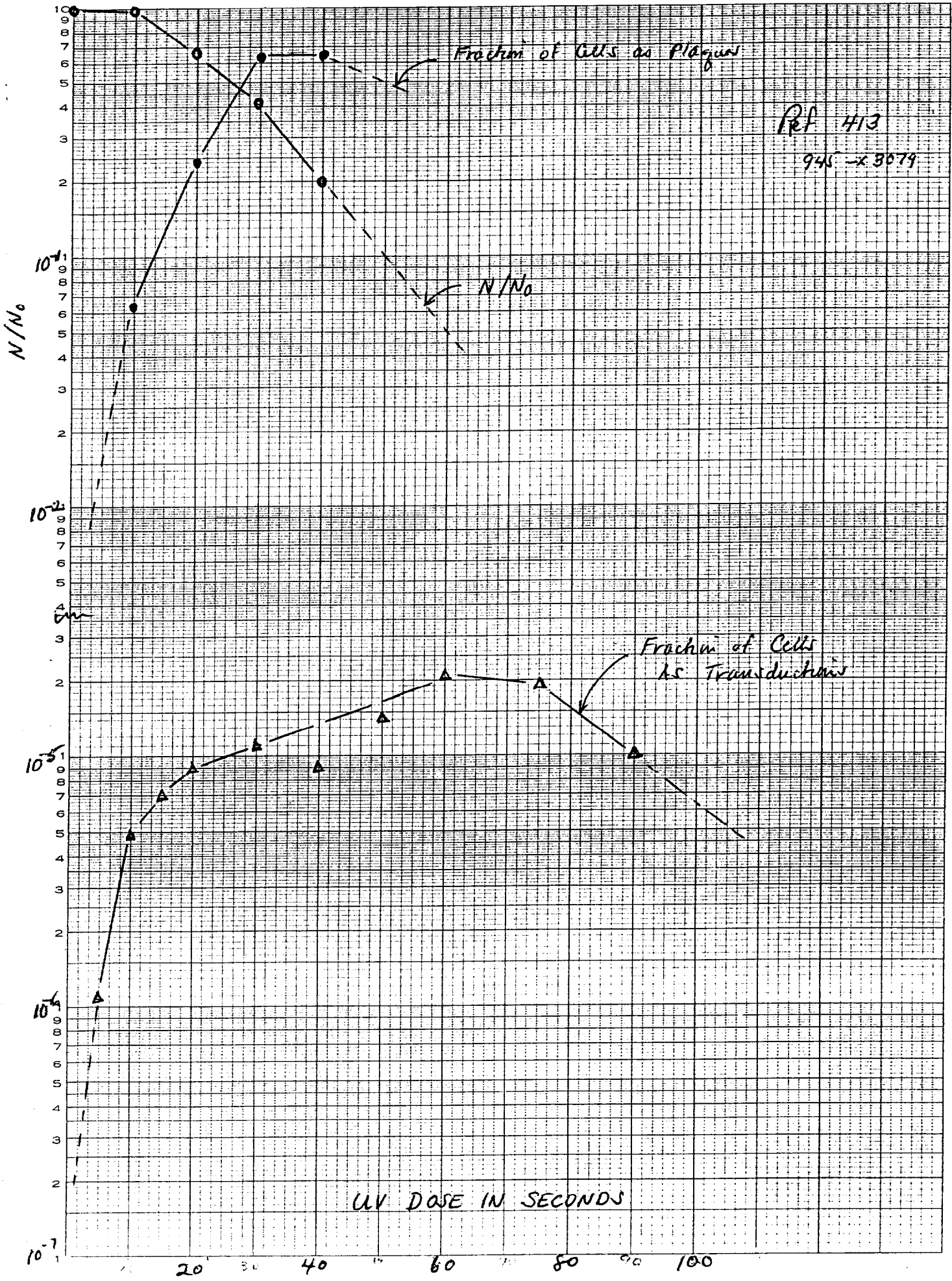
	Dose	Plaques	Cells	Total ^{out} _{average}	I/No	Plaques/wt	Friction Coef _{to Plaques}
10	0	0	198	198	1.0	0	0
	10	13	207	220	1.0	1.3 x 10 ⁷	0.064
	20	49	136	185	0.66	4.9 x 10 ⁷	0.24
	30	127	86	213	0.42	1.3 x 10 ⁸	0.64
	40	130 ^{many} _{large} lump.	42	172	0.20	1.3 x 10 ⁸	0.64

20 2. Transmission induction. ^{2.7 x 10³}
 hi 1-27 00 = 60

	Dose	Population	A	Transmission _{per}	Friction Coef _{yellowy trans.}
	0	23	0	0	0
	5	45	22	2.2 x 10 ²	1.1 x 10 ⁻⁶
	10	110	97	7.7 x 10 ²	4.9 x 10 ⁻⁶
	15	161	138	1.4 x 10 ³	7.0 x 10 ⁻⁶
	20	190	167	1.8 x 10 ³	9.0 x 10 ⁻⁶
	30	247	224	2.2 x 10 ³	1.1 x 10 ⁻⁵
	40	208	185	1.9 x 10 ³	9.0 x 10 ⁻⁶
30	50	292	267	2.7 x 10 ³	1.4 x 10 ⁻⁵
	60	444	421	4.2 x 10 ³	2.1 x 10 ⁻⁵
	75	411	388	3.9 x 10 ³	1.9 x 10 ⁻⁵
	90	235	212	2.1 x 10 ³	1.0 x 10 ⁻⁵

40

50



DATE:

1 2 3 4 5 6 7 8 9 10

Isolation of λ^k from 2307

1. Overnight culture - diluted and plated to give ca 100 colonies per plate
 2. 9 plates - two counted

1. 97 colonies } all colonies gold -
 2. 109 colonies }

3. Replicated to 3 plates spread with 3079, UV'd 20 seconds 50cm, etc.
 4. On the mini plates 3 colonies found not to give a lysogenic reaction (2 on one plate, one on another)

5. The colonies were streaked on λ , 3079 after suspicious found λ^k and non lysogenic

6. Lysozyme made (not as conc. as usual) - after 7 hours spotted on 3079 on B gal

1. ca. 20000 plaques
 2. ca. 17 " "
 3. ca. 9 " "

W2307 λ^k
 = W3172

$45/9 = 15 \times \text{at } 0.01 = 15000 \text{ per } 2 \times 10^8 = 15 \times 10^4$

About this time also: 2345 treated in the manner above.

1. Colony of ca. 300 (on 2 plates) found non lysogenic
 2. on subsequent test found λ^k

W2345
 against λ^k
 = W3173

21. 10 μ with 945 after 3 hr growth in Pen and plates 1-100 on B gal, red. on S gal

(3) Tested against HET 2, 1, + by 2, 0 by 1

(4) gave no plaque when treated as above in lysate

945- Are the ~~test~~ cells giving HET lambda ^{in LFT culture} equivalent? Ca 2×10^5 945 colonies per plate replicated to 3079 ^{spread plate} and UV'd 20 sec. Incubated 3 days gold + looked for HET. Several suggestions that some are UV induced reversions of 945 (as was the case previously with 1210) 15 plates in all representing ca 40,000 colonies. Areas on the 945 plate ~~located~~ corresponding to the Gold on 3079 located and picked and restreaked by spotting on 3080 and rechecking - This to find a clone of 945 cells giving HET lambda. Gold area picked from 3079 plate to confirm ~~handwritten~~ 12 945 area picked per plate.

Results

1. first 5 945 areas - 60 total found LFT. None of the Gold areas on 3079 corresponding to these showed any clone.

2. to distinguish between gold area to reversions of 945 plates replicated to B gal etc, since 3079 is Sp. From the Gold on Gold + SM reference made for the original plates to obtain reference to original clone. (2) the ^{UV} spread plate to see

to
418A

DATE: 3/27/56

REF: 413A

if HFT + clone induced by UV.

Z range = regions 445 plate - on UV'd
 A " = 3079 + 945 UV'd
 S " A same replicated to B gal SM

3. The SM Col plate show G₁T, indicating non-division or reversion of the doubly - - (or possibly more complicated).
 Examples ~~XXXXXXXXXXXXXXXXXXXX~~
~~XXXXXXXXXXXXXXXXXXXX~~

	1	2	3	4	5	6	7	8	9	10
10			Z3	19		Z12	7			
			A3	24		A12	33			
			S3	11 4%		S12	18			
			Z5	9		Z4	11			
			A5	ca 100		A4	40			
			S5	58 3/4 growth normal 15% 15% plate		S4	18 3/4			

Theoretically
 Z + S = A, however
 many many G₁T
 on plates and may
 but replicate well

4. From Z3, Z12, Z5, Z4, Z6

- ① 7 areas corresponding to the areas of transduction ~ 3079 (as determined by replication to SM) picked and spotted on B gal. G₁T to spot size (24 hrs); then from on-clone spots ~~with~~ growth picked, tested for HFT via UV on 3080.
- ② Results of the S5 area picked and tested, none found HFT + after 48 hours.

Repeat this experiment adding HFT + 1 to obtain lower limits of resolution of the method. Suggestion is that cell productivity (HFT) is not directly related.

3/2/56
 Preparing for Kallahan experiments

Obs that i, l, T are all dependent for 2173 on II (2062 transposon?)
 Prepare gal₂ -, gal₄ - stocks also 2734 (= non-transposable gal -)

1. Because 3082 variable, a new stock made by HFT₂ x 3010
 A. a reg. clone obtained.

2. 2734 crossed with 1895 - growth in both, plate M gal -
 A gal - colony picked purified once, & checked on D(10)
 and tested against HFT₂ (+)

A. 3/6 colonies give D(10) (labeled 413A-1) = W3142
 B. 3/6 colonies not trans. to + by HFT (+) - all must be λ.

DATE: 4/20/56

REF: 414

1. Recheck - 269-1 by side.

	1	2	3	4	5	6	7	8	9	10
			269-1	10 ²	10 ²	10 ⁴				
		3079								
		3080								
		945								
	(#13)	343-8								

10

↑
Resistant
above
pH

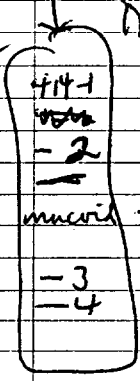
very tiny
plaques

2. Examination of the rest of 1.

After two purification streaks - 2 from each culture

		269-1 (1-100)	hand	1	1-2
3079	(1)	R	normal	S	S
	(2)	R	"	S	S
3080	(1)	R ?	"	R	S
	(2)	R	"	R	S
945	(1)	R	slight mixed	R	R
	(2)	S	normal	—	—
343-8	(1)	R	"	S	S
	(2)	R	"	S (mixed)	S

Stoelks



These tests in
B-jel - all
cultures Δ -

- is contaminated?

← streaks
discarded
3/9/59

20

30

341-9 Lysozyme 4/22/56 - To show that fragment size is constant - tubes
of 2" x 1-4", 4", 1" are the same

Array

lit 10⁴ → 0.05 ml (surface plating)

- ① Preliminary - 13080 (1-4) 101 hand.
- 13102 (1-1) 4 hand, ca 500 plaques.
- 13104 (4) 4 hand, ca 500 plaques
- ② Ppt. same divisions
- 13094 5 hand.
- 13091 8 hand.
- 13080 96 hand.

40

50

DATE: 5/1/56

REF: ~~415~~ 415

307-1C
 307-1A
 307-1B
 307-1D
 307-1E
 307-1F
 307-1G
 307-1H
 307-1I
 307-1J
 307-1K
 307-1L
 307-1M
 307-1N
 307-1O
 307-1P
 307-1Q
 307-1R
 307-1S
 307-1T
 307-1U
 307-1V
 307-1W
 307-1X
 307-1Y
 307-1Z

	1	2	3	4	5	6	7	8	9	10
	307-1C Segregation			1 - X 7 -		see 307A, 307				
	1. Overnight culture from a single colony B gal.									
	2. dil $10^2 - 10^4 - 10^6 - 10^8 \rightarrow$ 0.05ml samples spotted on B gal, 6 per plate. (8 plates)									
	Inoculated 4.0 hours, 0.05ml - 0.1ml 100% added and the plate area									
	Versed.									
	3. Parent sup. $\frac{100}{100} = 1.00$ $\frac{95}{100} = 0.95$ $e^{-1.1} = 0.338$									
10	Calculated			Gal +	Gal - pop	Gal -	Total			
	3 samples	1. 5?		10	209	10	229			
	get a + cell			10	212	9	231			
	3 samples	4. Samples								
	get a + cell	Plate 1		Gal +	Gal - pop	Gal -	Total			
	16 samples	1		220	0	0	220			
	set no cells	2		0	222	0	222			
		3		0	6	0	6			
		4		0	201	1	212			
20		5		58	ca 100	ca 100	-			
		6		0	9	0	9			
		Plate 2								
		1		0	265	1	266			
		2		0	0	0	0			NG
		3		0	185	0	185			
		4		0	209	0	209			
		5		3	309	3	312		0.602	$N \log N$
		6		0	2	0	2		1.8	7.78×10^{-2}
		Plate 3								
		1		2	192	0	192		1.2	4.38×10^{-2}
30		2		0	307	1	310			
		3		2	164	0	164			
		4		0	148	0	148			
		5		0	17	119	136			
		6		0	381	0	381			
		Plate 4								
		1		65	116	2	183			
		2		0	0	0	0			NG
		3		0	272	3	275			
		4		47	1	0	48			
40		5		0	183	0	183			
		6		0	0	0	0			NG
		Plate 5								
		1		3	252	4	259			
		2		0	233	0	233			
		3		0	77	0	77			
		4		0	0	0	0			NG
		5		0	260	105	365			
		6		0	21	0	21			
		Plate 6								
50		1		0	158	3	162			
		2		0	38	0	38			
		3		0	245	1	246			
		4		0	129	0	129			
		5		0	464	1	467			
		6		0	223	10	233			

$230/100 \times 2 = \frac{2.3}{2} = 1.15$ / sample

$\frac{E.C.}{N \log N} = a$
 $1.8 \times 7.78 \times 10^{-2} = 2.3 \times 10^{-3}$

4.38×10^{-2} 3.6×10^{-3}

DATE: 5/1/56

REF: 415A

	1	2	3	4	5	6	7	8	9	10
	Continuation -									
		Gal+	Gal-py	Gal-	Total					
	Plate 7	0	0	0	0	NG				
	1.	0	153	0	153					This det. is 1/2 (307-1C)
	3.	0	68	0	68					
	4.	1.	ca 50?		ca 500					For comparison with Lot
	5.	0	450	1	451					389 for 307-1A
	6.	125 ^{CAF}	25	0	150					
10	Plate 8	0	223	1	224					
	1.	0	0	0	0	NG				
	3.	0	72	0	257 72					
	4.	0	-	ca 100	467 -					
	5.	0	272	5	277					
	6.	0	223	2	225					

II. Single Gal- py obtained from as many clones as possible - 17 obtained from 17 clones.

20
 6- Lp^d = 8
 7- Lp^d = 8
 7- Lp^{M/S} = 1 = 415A-1

30

40

50
 3057 - in search of Lp^R
 1. ca 2000 colonies tested by replication / 2730 seq (Gal₂-) - None found with Lp^d

DATE: 5/3/56

REF: 416

1 Redo a 945 experiment to see if "mutants" are responsible for + strand activity in LFT cultures.

1. Technique - nearly confluent colony formation of 945, to which was added a few HFT⁺ cells as a check - replicated to 307² (Coul + ¹⁰⁰ (F₂-purified) m)

AMSD

This plate was replicated to M Col

10 Example of best after 2 days
 1/2 plate 307
 Bgal replica 300
 Bgal SM 10
 Mgal replica 300

2. Aerial 945 diluted 10²-10⁶
 Overnight 389-1

10⁶ → 0.05 ml
 1.0 ml
 10²-10⁴-10⁶
 ↓
 0.05 ml = 108
 65
 173

2. 118
 231
 0.1 ml / 10 plates
 This contains 231 x 2/9 x 100 = 4620 = 512
 19 945 cells
 230²
 25 col LFT
 11 col LFT
 173 x 1/12

6 original plates (w/ UV) ca 10

20 This experiment abandoned - Replications to Bgal SM showed ca. 10 Col + per irradiated plate which presumably are non-duplicates of 945 x 307. The irradiated plates (and Bgal replica of them) showed at best about 200-300 Col + clones - of these replications to M Col showed at least 200 were prototrophic and derived presumed

30

W 2872 (12102 - 81A)
 in search of HFT⁺
 1. 17 seg tubes against B⁺ 3094 - all LFT but 8⁻ appears to have a selective advantage of 4⁻ as it grows through G₄ background.
 2. 21 seg tubes / 3094 - all LFT
 15 seg tubes / 3094 - all LFT

40

W 7580 x W 7584
 no add 14
 0.1 ml by 54 (from 4000)
 50 9. 13 seg tubes - 12 Col⁺, 1 Col⁻ G₄⁻

DATE: 5/11/56

REF: 417

Segregation in 323-2
as in previous ofpts

Overnight cult 10^7 10^4 10^6 $10^8 \rightarrow 0.05$ ml sample B pet - 10 spread 4 lines.

↓
copy 0.1 ml -

10

	+	-	pop -
1.	18	30	99
2.	22	24	116

Total

2

Respread
Clones

Colony

#s 1-11

+

-

pop -

no growth

20

#s	+	-	pop -
12.	21	0	0
13.	0	0	22
14.	0	33	0
15.	0	0	92
16.	0	0	30
17.	0	0	7
18.	0	1	0
19.	0	13	2
20.	0	0	42
21.	0	0	29
22.	0	0	81
23.	0	0	3
24.	0	0	16
25.	0	6	0
26.	0	0	4
27.	3	4	37
28.	20	0	22
29.	0	0	28
30.	0	0	43
31.	0	8	12
32.	0	0	18

30

40

50

DATE: 5/13/56

REF: 418

	1	2	(2 ³ -x7-)	4	5	6	7	8	9	10
	Segregation from		W287F	2-7	heterozygote - It is full Gal ⁺ , both Gal ⁻ alleles?					
	1.		12-	17-	Number					
			+	0	14					
			0	+	4					
		2.	14/14	2- tubes for HFT ⁺ / 3091						
				all LFT						
10			4/4	7- tubes / 3092						
				all LFT						

} both alleles present
but segregation
pattern reversed

At this time W2308 x HFT⁺ } to compare reciprocal cross
W2915 x HFT⁻ }

made in the "standard" way. 0.1 ml of 1-yr old overgrown culture + 0.4 ml HFT
lysozyme - diluted to obtain ca 100-200 colonies / 13 gal plate (150 ± 10%)
Results not recorded but ca 200 colonies / plate. 75% show contact with
phage (partially lysed)

FOR
SEG.
ANALYSIS
SEE
420

W2308 - 2 apparent transductant clones apparent [418-1, 418-2]
W2915 - 1 trans. clone obtained - not as "x" as the clone [418-3]

A. The mixed plates of these experiments replicated to HFT⁺ + 3091 and incubated
with UV 30 minutes. This, to see if any homozygous clone formed by transduction
(one transductant clone from here been selected on the basis of plus phenotype, in
the case of (-) recipient (-) or minus phenotype in (+) recipient. The evidence
of HFT (lay gal⁺ clone on HFT⁻) indicating no Gal⁻ homozygote formed.
The extent of the Gal⁻: ca. 500 gal⁻ colonies (W2915) } planned to replicate
in 300 gal⁻ colonies }
to Gal⁻ indicator
generated by reversal of
recipient

5/14/56
1210 x - 308-5 }
8- x - 1-6- } for stock for reanalysis

no odd - 0
0.1 ml 308-5 84 (small) ① 15/15 found 8- 15 tubes / 3091 and found LFT
lysozyme (original, perhaps 2 years old) ② 7/7 found 8- 7 tubes / 3091 MVT found LFT

1210 x - 344-6
8- x - 6-7
no odd - 0
(old lysate) 0.1 ml 153 (small) ① 19/19 found 8- 18 tubes / 3091 and found LFT
344-6 ② 11/11 found 8- 10 tubes / 3091 MVT and found LFT
③ 16/16

50 2580 x - 346-4
no odd - 2
0.1 ml 346-4 64 ① 20/20 found 19 Gal⁻ } See 347
② 10/10 found 1 Gal⁰ }

DATE: 5/16/56

REF: 419

	1	2	3	4	5	6	7	8	9	10
	<p>1. In the experiments to see if HFT⁺ cells are distributed as "mutants" in LFT clones (such as on 4/6) many self clones possibly transductions, reversions or something else ^{observed on the 1st passage} appear to be present. Both cultures used in these later experiments were F⁻ and no use crossing is prohibited. Are beta-caryons being found.</p>									
	<p>2. Experiment -</p> <p>a. To avoid lambda transduction by ¹⁰ cultures used.</p> <p>W2915 F⁻ T⁺ B⁻ = Gal⁻ Lac⁻ Ara⁻ Xgt⁻ MHI⁻ P⁻ Lp⁺</p> <p>W3079 F⁻ M⁻ Gal⁻ Lac⁻ Ara⁻ Xgt⁻ R⁺ SR Lp⁺</p>									
10		<p>populas</p> <p>W2915 UV 9</p> <p>W2915 11</p>		<p>} which of these UV'd not known, plates not labeled</p>						
		<p>W3079 0</p> <p>W3079 0</p>		<p>} which of these UV'd not known - both labeled as such</p>						
		<p>W2915 } UV</p> <p>W3079 }</p>		<p>Two classes of populas large = ca 30 - Gal⁺ or res⁺</p> <p>small = ca 500 - most of these just faintly + or Gal⁺ or res⁺</p>						
20	<p>6. Test some of the populas for markers.</p> <p>① 6 small populas listed in Bera - 1 Gal⁺, ara⁻</p> <p>4 Gal⁺, ara⁻</p> <p>1 Gal⁺, ara⁺</p> <p>6 large populas listed in Bera - 6 Gal⁺, ara⁻</p>									
30										
40										
50	<p>3551 X K1⁺ (new lysate) made 5/14/56</p> <p>no odd = 0</p> <p>o.l.w = 14</p> <p>1. 15 reqs and found 7-7-</p>									
	<p>346.4 X K12 (new source) this culture very rough.</p> <p>no odd = 0</p> <p>o.l.w = 19</p> <p>7. 21 req found 4-6-</p>									

DATE: 5/23/56

REF: 420

	1	2	3	4	5	6	7	8	9	10
	Say analysis		418-1	2-17				This cleavage		
		Number	12	17						
		16	+	0	= 7-	all 4 ^s				
		2	0	+	= 2-	3rd 4 ^s 1 ² , probably 4 ^s				

10	Say analysis		418-3	7-12						
		Number	12	17						
		4	+	0	= 7-	all 1 ² , probably 4 ^s				
		8	0	+	= 2-	all 4 ^s				
		1	0	0	= 2-7?	appears 4 ^s				

20	2880X-2851		2-14							
		Number	11	12	14					
		19	+	0	+	= 6 ² -	— hand			
		2	0	+	+	= 6 ² -	we saw 9			
							0.1 me 2817.50			

30	Heterogeneity for Photographic Illustration									
	Lysate of		K-12							
	Cult.		meat		0.1 me					
		3091	0	45						
		3092	4	64						
40		3094	0	63						
		3096	2	80						
		3097	0	44						
		2857	0	88						
		2854	0	55						

50

DATE: 5/20/56

REF: 421

Segregation from some heterozygotes to check them before use in photogrophy

1	2	3	4	5
2580 X-308-5				
Number	1-	2-	16-	Genotype
16	+	0	+	Gal ₂ -
2	0	+	0	Gal ₆ -Gal ₁ -

10

1	2	3	4	5
2580 X-344-6				
Number	16-	12-	17-	Genotype
19	+	0	+	Gal ₂ -
1	0	+	0	Gal ₆ -Gal ₇ -

20

30

40

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