

Resume of DNA synthesis in various passages  
expts 0-132 and N111, M14122

<u>Infecting virus</u>	HIRT DNA Total cpm in				Pfu/ml.	Total cpm pfu/ml	Days to full CPB	Pfu x 10 <sup>3</sup> homology 16
	I	II	I+II	I/II				
X 8	2.2x10 <sup>6</sup>	1.6x10 <sup>6</sup>	3.8x10 <sup>6</sup>	1.4				
0-132								
777 CV/1 B	2.8x10 <sup>6</sup>	2.5x10 <sup>6</sup>	5.3x10 <sup>6</sup>	1.1	1x10 <sup>9</sup>	5.3	0.3	4
777 CV/1 G	5.6x10 <sup>5</sup>	4.6x10 <sup>5</sup>	1.0x10 <sup>6</sup>	1.2	1x10 <sup>8</sup>	10	0.85	5
777 CV/1 H	7.8x10 <sup>5</sup>	1.4x10 <sup>6</sup>	2.2x10 <sup>6</sup>	0.56	3x10 <sup>6</sup>	730	0.72	>9
777 CV/1 I	6.8x10 <sup>5</sup>	5.8x10 <sup>5</sup>	1.3x10 <sup>6</sup>	1.2	2.1x10 <sup>5</sup>	6500	1.6	>11
0132								
776 CV/8 B	2.0x10 <sup>6</sup>	1.0x10 <sup>6</sup>	3.0x10 <sup>6</sup>	2.0	5x10 <sup>8</sup>	6	0.8	4
776 CV/8 F	5.1x10 <sup>5</sup>	1.2x10 <sup>6</sup>	1.7x10 <sup>6</sup>	0.43	5x10 <sup>6</sup>	340	1.4	>14
776 CV/8 G	8.6x10 <sup>4</sup>	4.3x10 <sup>5</sup>	2.2x10 <sup>5</sup>	0.66	2x10 <sup>6</sup>	110	3.7	8
776 CV/8 H	0.9x10 <sup>6</sup>	2.1x10 <sup>6</sup>	3.0x10 <sup>6</sup>	0.43	4x10 <sup>7</sup>	75	2.3	4
776 CV/8 J	7.2x10 <sup>4</sup>	1.2x10 <sup>5</sup>	1.9x10 <sup>5</sup>	0.60	0.9x10 <sup>6</sup>	210	15	n.d.
N-111								
777 NP <sub>2</sub>	3.3x10 <sup>6</sup>	7.2x10 <sup>5</sup>	4.0x10 <sup>6</sup>	4.6			3.3	4
777 NP <sub>4</sub>	1.3x10 <sup>6</sup>	1.5x10 <sup>6</sup>	2.8x10 <sup>6</sup>	0.9			25	>12
777 NP <sub>5</sub>	1.2x10 <sup>4</sup>	1.0x10 <sup>5</sup>	1.1x10 <sup>5</sup>	0.12			40	n.d.
N-111								
777 CV/B B	6.8x10 <sup>6</sup>	1.4x10 <sup>6</sup>	8.2x10 <sup>6</sup>	4.9	0.9x10 <sup>9</sup>		0.32	3
777 CV/B E	1.7x10 <sup>6</sup>	2.3x10 <sup>6</sup>	4x10 <sup>6</sup>	0.7	1.1x10 <sup>9</sup>		75	4
777 CV/B D					2x10 <sup>9</sup>		35.3	
777 CV/B E <sub>2</sub> (labeled)					5x10 <sup>8</sup> -1x10 <sup>9</sup>		81	
777 CV/B F					9x10 <sup>7</sup>		71	
777 CV/B G					4x10 <sup>6</sup>		78	

# Prep of BSC DNA

D 71.

See methods book.

7.27.71 Pellet of 12.11.71, BSC Flow 60 plates. Washed 2x with PBS.

Throw at rm temp. Add 1 ml / 5 plates = 12 ml of SDS - Saline (see methods) (S Lauer). ~~Vortex~~, Wash onto long t.t. Vortex. ~~#~~ Very viscous, not really homogeneous, see lumps. Add 6 ml ( $\frac{1}{2}$  SDS-saline) 1 M NaTCA, pH 7.4. Inc. 37° 30 min. Still some light tan lumps. Add 19 ml. Tris- $\text{G}(\text{OH})$  Shake on rotary, high speed, 30 min. Spin 15' @ 8,000 rpm Sorvall. Collect upper, aqueous. Difficult to get last bits, got only 10.8 ml. Therefore, decided to wash  $\bar{c}$  more ( $\sim$  ml/centrifuge) of 1x SSC, 0.05 M Tris pH 7.9.

Quick vortex. ~~#~~ Spin as above.

Collect total of 20 ml aqueous. Add 1 ml 20x SSC.

Add 42 ml cold 95% EtOH along side of rod al beaker. Stir with rod winding DNA onto rot.

Dip 3-4x in fresh EtOH tube. Press against ~~#~~ wall to drain. Dry 5 min at rm temp.

Dissolve in 3 ml 0.01 SSC 1 hr in temp.

Set overnight. refriger.

8.12.71 Add  $\frac{1}{20}$  vol. 20 SSC (0.15 ml) and 0.2 ml of heated 1 mg/ml RNase. 8:25 am. 37°. To 9 am.

Add equal volume (3.35 ml) 0.02M Tris. Vortex 15', "medium" in PYREX Sorvall tube. Centr. as above. Take off sup. Store several hrs. refriger. 32°

Add equal vol 99%  $\text{CHCl}_3$  1% isoamyl alc.

shake on hand vortex. Spin Clinical Centr.

$\sim$  10 min #5. (all in 12 ml conical). Remove sup. 2.8 ml. Only tiniest speck of protein at interface.

Place 2.8 ml in 40 ml conical tube. Add 1/20th vol, 0.14 ml 20×SSC. Add 2 vol aq to EtOH, cold as before, wind DNA, rinse in EtOH. Dry in air. Put to dissolve in 1 ml. 0.01 SSC. 30 min. on shaker. Refrig. overnight

9-12-71 Dialyze vs 1L 0.1 SSC start noon. Cold.

10-12-71 Change to 1L 0.01 SSC.

11-12-71 Change fluid.

12-12-71 Stop. 11:30 pm. 1.5 ml.

$$0.02 \text{ ml} + 1.00 \text{ } \frac{1}{100} \text{ SSC. } A_{260} = 0.210$$

$$\frac{.210(1.02)}{.02} = 10.7 A_{260}/\text{ml.}$$

$\frac{60 \text{ plates}}{15 \text{ ml}} \times 15 \text{ ml} = 60 \text{ ml}$   
 $\frac{60 \text{ ml}}{60} \times 0.25 \text{ mg/plate} = 0.25 \text{ mg/ml}$

$$\text{then, } 10.7 A_{260}/\text{ml} = \frac{530}{535} \text{ mg/ml} = 0.54 \text{ mg/ml.}$$

E 0.81 mg yield.

Store frozen

$$50 \mu\text{g/filter, } \frac{0.81}{50} = \frac{810 \mu\text{g}}{50} = \boxed{16 \text{ filters}}$$

mp & t

$$\frac{810}{40} = 20 \text{ filters}$$

20V .21

$$\frac{810}{35} = 23 \text{ filters}$$

HNO<sub>3</sub>

HF

310 V

$$\frac{810}{30} = 27 \text{ filters}$$

Growth of *H. influenzae* strain Rd.

D-9171

9.12.71  $\frac{90}{400 \text{ mg DPN}} \text{ in } \frac{9}{\text{ml H}_2\text{O}}, \frac{40^\circ}{10 \text{ mg/l NAO}}$

Biohringer Control # 6458441 (663) Free acid.  
free acid.  $\text{C}_{21}\text{H}_{27}\text{O}_{14}\text{N}_2\text{P}_2$

Stable cold neutral, (Kornberg, Meth Enz III p 878.)  
Very labile in alkali.

$$\textcircled{2} \quad \epsilon_{260} = 18 \times 10^6 \text{ cm}^2/\text{mole}$$

21 x 12	252	Add 5 ml H <sub>2</sub> O. Then add 1 N NaOH
27 x 1	27	to pH $\approx$ 6.8. Then H <sub>2</sub> O to 9 ml.
14 x 16	224	Took $\approx$ 0.2 ml 1 N NaOH. pH 6.0.
7 x 14	98	$1 \text{N} = 1 \text{ mmole/l}$ .
2 x 31	<u>62</u>	$\approx 0.27 \text{ ml}$ .

$$\frac{90 \text{ mg}}{663 \text{ mg/MW.}} = 0.136 \text{ mmoles.}$$

$$663 \text{ mg/mole} \quad 0.27 \text{ mmole.}$$

Sterilize in 2 lots thru Swinney into sterile tubes.  
Freeze.

\* Brain Difco  
12.12.71 Beef Heart Infusion: 37 gm dissolved in 1 liter  
deionized H<sub>2</sub>O. Disperse 50 ml/each  
of 6 250 ml flasks.

\* BHI-Agar:  $\frac{37}{4} \text{ gm } + \frac{15.02}{4} = 3.75 \text{ gm Difco agar}$ ,  
in 250 ml deionized H<sub>2</sub>O. After autoclave, cool  
to 55° in water bath. Add, 2.5 ml hemin, 0.05 ml DPN. Pour  
plates. Cool. Keep overnight 37°.

\* Nutrient Agar, Difco, 5.75 gm + 250 ml H<sub>2</sub>O. Heat to boil  
to dissolve. Autoclave. Cool to 55°. Pour plates.  
Cool. Keep overnight 37°.

\* Vials -  $\frac{1}{2}$  0.2 ml glycerol

\* Sterilize by autoclave, 15 min.

Equine hemin. 2x recyst. Mann 3054, Batch V3032.  
from J Gressel.

9.8 mg into 9.8 ml triethanolamine in 50 ml sterile  
Balenmeyer. Much dissolves ↗ WRONG  
15 min 65°. Store at 4°. see below

12.12. Received from Nathans. (sent B.12.7)

4 vials, stabs, incubated 37°. Growth visible  
4 vials, stabs. not incubated. No visible growth.

Inoculate 50 ml BHI-DPN-Hemin #1, 37° stab

Inoculate 50 ml BHI-DPN-Hemin #1, 25° stab

Shake - 37° - start 11 pm.

Put #2, 25° stab in 37° incubator. 11 pm  $\frac{1}{2}$

13.12. 25° #2 Stab - no signs of growth. 9 am. Remove  
from incubator.

37° #1, 25° #1, no signs of growth in water bath. 9 am  
noon see growth.

Media for these 2 and BHI plates made up 12-12 ~~for~~  
Wrong since hemin should have been in  
4% triethanolamine.

Make 2 ml triethanolamine  $\rightarrow$  50 ml in sterile water.

Filter thru millipore. Add 50 mg Hemin (E.K. from  
Nathans.) 15' at 67°.

14°  
20°  
25°  
30°

From 37° #1.  
Streak on BHI Choco Agar  
Streak on BHI (12.12 plate)  
Rest into liquid medium.

D-91  
P2

13 | 12.71 Cool incubated media 230 pm

Read Abs 650 vs H<sub>2</sub>O.

medium	0.015
25° #1	0.016 - Discard
37° #1	0.174.

Add Hemin + DPN to 50 ml media aliquots using new hemin solution.

Inoculate 2 flasks (50 ml media) with #1 0.50 and #2 1.00 ml  
of 37° #1 above. (1 and 2%). :- Zerotime of media = 0.069 = 41.5 pm.

Inoculate a third c stab from  $37^{\circ}$  #2 - stab of Nathan.

Store 37#1 in cold room  
Older from 37#1. (37°.) Street 4:15 pm.

BAT - Hemen - DPN

## Blood Algae

## Nutrient Algal

Choc Agar

$\frac{6}{1} \cdot 52$	0.097	0.110	0.072
$\frac{6}{1} \cdot 52$	0.028	0.041	0.003
-2010			
$10^{\circ} 2$	1.665	1.832	0.117
$\frac{11}{1} \cdot 780$	<del>4568</del>		0.045
-2010			
-2010			0.156
-2010			0.084

Step 1 to 2 at 10<sup>02</sup> pm, cool.

14.12 #1, #3 - Dense cells.

Control #4.

Medium menus items, menu DPN.  $A_{G50} = 0.035$ .

Add 1 ml of 37# <sup>note have opened this many times now.</sup> 1 at 10:30 pm. Put @ 37° Shake.

$A_{650} = 0.085$  at 3:30, 14.12. Replace at  $37^\circ$ :  $A_{650} = 0.085$  on 15.12.

O.K. Essentially No Growth.

From #1. 1 ml of #1 / vial containing 0.2 ml.  
glycerol.

Mark: Vial 1/091

~~to~~ 15 vials.      ~~Mark 1 → 15~~

From 37#1, original culture - of 12.12 & 12.13.

10 vials. mark 2/091. (Direct from stab)

mark 1-10

Store -20° overnight

Then next day put at -60°.

D-91

P3.

For antibiotic test

20-12-

Vial 1/Dai #2.

21-12-71.

Streak, 2 BHI plates  
1 chocolate agar. } 37° noon.

Store vial @ -20°.

22-12-71.

Good colonies on Choc. agar.

Nothing on BHI.

Take Choc Agar plate to lab.

951 224 86  
7djk 7"3  
4 1"Cejk 8"7  
5/21/77

D-211-71.

Trials for Best prelabeling conditions

21-12-71

large plates-  $d = 8.6 \text{ cm}$   $r = 4.3$ , area =  $\pi(4.3)^2 = 58 \text{ cm}^2$ .  
small plates  $d = 4.7 \text{ cm}$   $r = 2.35$  area =  $\pi(2.35)^2 = 17.4 \text{ cm}^2$   
ratio of area of large/small =  $58/17.4 = 3.33$ .

Small plates, seed  $2 \times 10^5$ , grow to  $10^6$  in 6 days.  
Usually 4 ml of medium. 8 ml in large plates.  
Seed 16 plates  $\bar{c}$  BSC today.

22-12- Sparsely cells.

23-12-

day 3 or 4 -  $\approx 50\%$  perhaps more  
Trials for long term labeling at low level of  $^3\text{H}$ -dT.  
4 plates: Replace medium  $\bar{c}$  MEM + 10% calf serum  
 $5 \times 10^{-5} \text{ M}$  in dT (disregard dT in hot).  
 $2 \mu\text{Ci}/\text{ml}$ , dT- $^3\text{H}$  in 2. 2 cold.  
 $1 \text{ mCi}/\text{ml}$ , 23.7 Ci/mmole,

D-211.1       $^3\text{H}$   
2       $^3\text{H}$   
3      no  $^3\text{H}$   
4      no  $^3\text{H}$ .

Plans #2 & #4, replace medium  $\bar{c}$  hot and cold, respectively,  
after 24 hr. Save medium to count.

For 20 ml medium,

18 ml MEM

1.8 ml calf serum.

0.25 ml. stock cold dT, 1 mg/ml,  $42 \times 10^{-4} \text{ M}$ . 4.2 micromoles/l ( $\frac{1 \text{ ml}}{4 \text{ ml MEM}}$ )

Add 4 ml to 3 x 4. Have 12 ml left, need 24  $\mu\text{Ci}$  add.  $0.025 \text{ ml } ^3\text{H}$  dT.  
Count medium A

D21171  
P2.

23-12 Save 0.1 ml of  $^3\text{H}$ dT + 4 N MEM, frozen sterile.

24-2 #2. #4. Remove medium. Replace ~~c~~ hot (2) + cold (4) fresh medium ~~ex~~ made exactly as 23-12, using dilution of  $^3\text{H}$ dT. made 23-12: ~~Medium E~~ (9 ml MEM + 1 ml CS + ~~0.5 ml~~ of 0.13 N cold stock dT. Add 4 ml to #4. Then add 0.5 ml of diluted  $^3\text{H}$ -dT, add 4 ml to #2. (Medium E).

~~Plan: at 60-70% confluence: 2 plates at 24°. 2 days.~~  
~~Then label at  $5 \times 10^{-5} \text{M}$  dT, 10  $\mu\text{Ci}/\text{ml}$ .~~  
~~12 hr (use warm medium).~~  
~~Chase 2 days on maintenance medium.~~

24-12

Plan at 60-70% confluence. Actually 75-80%.

Media for 5-10.

5  $10 \mu\text{Ci}/\text{ml}$   
 $23.7 \text{ Ci/mole}, 5.1 \times 10^{-7} \text{ M}$ .

6 - no label, no dT.

7  $10 \mu\text{Ci}/\text{ml}$   
 $2.37 \text{ Ci/mole}, 5.1 \times 10^{-6} \text{ M}$

8  ~~$5 \times 10^{-6}$~~  no label,  $5.1 \times 10^{-6} \text{ M}$  dT.

9  $10 \mu\text{Ci}/\text{ml}$   
 $0.237 \text{ Ci/mole}, 5.1 \times 10^{-5} \text{ M}$  dT

10 no label,  $5.1 \times 10^{-5} \text{ M}$  dT.

F: 14 ml MEM + 1.5 ml CS + 0.15 ml  $^3\text{H}$ dT ( $15 \mu\text{Ci}$ ) (plate 5)  $\frac{4}{4}$

G: Same but minus  $^3\text{H}$ dT to #6 (44).  $\frac{4}{4}$

C: 5 ml F + 0.2 ml  $1 \rightarrow 33$  dT (0.025  $\mu\text{mole}$ )  $\frac{4}{4}$

$^3\text{H}$ -C: 5 ml F + 0.2 ml  $1 \rightarrow 33$  dT  $\frac{4}{4}$

D: 5 ml G + 0.2 ml  $1 \rightarrow 3.3$  dT (0.25  $\mu\text{mole}$ )  $\frac{4}{4}$

$^3\text{H}$ D: 5 ml F + 0.2 ml  $1 \rightarrow 3.3$  dT.  $\frac{4}{4}$  to 9

(dilutions in MEM.)

25-12 Nbrs 1 → or at

10 all look like close to full monolayer.

D21171.

p3

~~12.20 - 12.21~~

26 12.21, All good monolayer. #7 may be slightly less dense.

26 12.21 Do chase on all plates #1 → 10.

Stocks are 1 mg/ml dT and dC. want 50 µg/dL dT, 10 µg/dL dC.

10 plates: wash  $\times \bar{c}$  <sup>4 ml</sup> warm chase medium, then add 4 ml for maintenance ( $\bar{c}$  to <sup>150</sup> cold serum).

Make ~~10~~ <sup>150</sup> ml medium

150 ml MEM

3 ml CS.

2.5 ml dT stock cold.

1.5 ml dC stock

28 12.21. (Tues) 10 am. #7 still looks not quite so good, also #5.

wash each plate  $\times \bar{c}$  4 ml PBS

$\times \bar{c}$  2 ml PBS.

Prior to wash,

Add 2 ml <sup>0.5</sup> NaOH. From labeled plates: 1, 2, 5, 7, 9, add ~~eth.~~ overlying medium (chase medium) to one of plates that was untreated. Remove immediately. Remove all media. Wash.

Add 2 ml 0.5 M NaOH. as per methods.

Remove 2 0.5 ml aliquots; to each, <sup>cold,</sup> 0.5 ml cold 20% TCA. <sup>a</sup>b.

c: count 0.1 ml by drying onto GF/C for total cpm.

a**b**, filter & wash a. per methods. Dry ~~1 hr~~.

Dry a, b, c, 1 hr 80°. Toluene - count.

For all plates, hot, cold, controls, take 0.1 ml to 1 ml for Abs. reading.

D21171

p4.

Count dry filters in follow.

Background ~~is~~ 20, included in blank: for C, total cpm =  $\times 20$ .  
for exp't. for other up to for ~~other~~, total cpm  $\times 4$ 

	CPM repeat	CPM - Bl	Total cpm/plate	A <sub>260</sub> 260	#	A <sub>260</sub>	
C	1 control	59	51		3	2.35	
	1	3181	3120	3065 61,300	2.06	4	2.34
	2 control	27	22		6	2.32	
	2	3271	3245	3233 64,660	2.01	8	2.37
	5 control	120	103		10	2.31	
	5	221,600	220,700	221,040 44,20,800	control	2.25	
	7 control	56	64		1 cont	2.34	
	7	73,360	74,631	73,304 14,66,080	2 cont	2.13	
	9 control	41	35		5 cont	2.07	
	9	25,320	25,080	25,280 505,600	7 cont	2.25	
a+b	1 control	194	191	173	9 cont	2.21	
	1 control	168	159	144			
	2	34	34	14			
	2	16	28	2			
	5	181	206	170			
	5	179	177	158			
	7	123	140	112			
	7	159	156	138			
	9	130	112	101			
	9	115	120	98			
	1	15,200	15,341	15,300 61,200	> 2.06		
	1	13,500	13,260	13,400 53,600			
	2	13,100	13,060	13,100 52,400	> 2.01		
	2	13,562	13,250	13,400 53,600			
	5	890,000	898,945	899,000 3,609,000	> 1.54		
	5	900,000	899,036	899,000 3,600,000	> 1.54		
	7	54,000	255,700	514,000 2,056,000			
	7	255,000	107,600	254,500 1,020,000	1.68		
	9	107,000	107,990	108,000 436,000			
	9	114,000	108,800	114,000 456,000	1.88		

Controls indicate  
less than 2%  
conformation by medium  
in all cases. Much less  
than that in some.

Ignore:  
Counts on hot media ( $\frac{1}{10}$  of total)

(cpm)

23-12-71 5100

"C" 47,000

"D" 46,000

"E" 4500

"F" 51,000

D21171

## Summary - Preflabel conditions.

P5

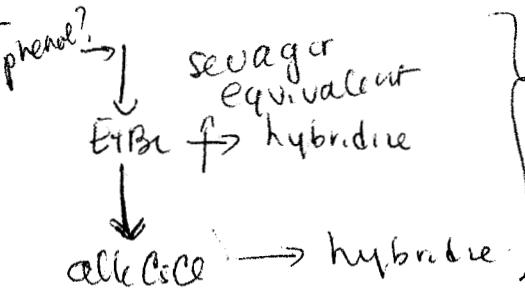
Experiment	$^{3}\text{H}$	Input			Determined			Acid insol cpm per Abs.	
		conc. $\text{dF}$	$\mu\text{Ci}/\text{ml}$	S.a.	Total cpm per plate	Total acid insol cpm per plate	Total Abs per plate		
		media. $\mu\text{Ci}/\text{mmole}$	media	cpm/ml	cpm per plate	cpm after chase per plate	cpm of chase - plate		
1	+	$5 \times 10^{-5}$	2		$5.1 \times 10^5$	$6.1 \times 10^4$	$5.7 \times 10^4$	$4.1$	$1.4 \times 10^4$
2	+	$5 \times 10^{-5}$	2 + 2		$5.1 \times 10^5$	$6.5 \times 10^4$	$5.3 \times 10^4$	$4.0$	$1.3 \times 10^4$
3	-	$5 \times 10^{-5}$						$4.7$	
4	-	$5 \times 10^{-5}$						$4.7$	
5	+	$5.1 \times 10^{-7}$	10	$2 \times 10^7$	$5.1 \times 10^6$	$4.4 \times 10^6$	$3.6 \times 10^6$	$3.1$	$1.2 \times 10^6$
6	-	$5.1 \times 10^{-7}$						$4.6$	
7	+	$5.1 \times 10^{-6}$	10	$2 \times 10^6$	$4.7 \times 10^6$	$1.5 \times 10^6$	$1.5 \times 10^6$	$3.3$	$4.6 \times 10^5$
8	-	$5.1 \times 10^{-6}$						$4.7$	
9	+	$5.1 \times 10^{-5}$	10	$2 \times 10^5$	$4.6 \times 10^6$	$0.5 \times 10^6$	$0.45 \times 10^6$	$3.8$	$1.2 \times 10^5$
10	-	$5.1 \times 10^{-5}$						$4.6$	

Compare 5 & 9 - s.a. of  $^{3}\text{H}$  dF decreases 100x from 5 → 7, but total counts incorporates decreased less than 10x.

Partly Also, have more cells in #9 - since Abs 3.8 compared to 3.1 in #5.

Use 5 or 9. Do infection in chase medium.

Use X8



MIRT - 5-10 plates pooled  
maximum

think about using vircors.  
20 plates - good banding

fragmentation.