

6/20/56
±

J. Lederberg

Tues. Bealle Ris / Bangs, Mitchell
Warney Rhodes

McCollum Pratt Symp.

Wed 9:00 ~~Sydney~~ Mayia Alfrey Sol / Hammett, Robbins, Zimm

Thurs. 9:00 Hartman Jacob Conrat / Chargoff Paulshaus (100)
1000

Fri 9:30 Art K, Echoa Cohen / Street, panel.

suggested date J.J.

1. Bill Mac asked me how after the Ford Eye - where my function was to resist strenuously the idea that mutants were making any progress at all. If I assume that my function here is to be a "wower", I hope this won't be taken as a balanced presentation of my own views. It is more ^{of a} pieceable than may appear, and I also hope the copy won't read "indefensible" when I read "ridiculous" as happened at Detroit in discussing Dr. Hammit's paper.

alt - moderator: don't mention flux & try it from doing anything
 fed back problem of moderator = disc leader.

hardly
 (~~is~~ to talk about something is hard & about - is, feed)
 won't transmit until could study

but Lp^+ / Lp^s heterozygotes)
 { H^s bal
 Lp bal linkage }

Mr. Achor from you.

Beale

mediated... chromosomes. W/C suggests mechanism

can't find can
mechanism or
mutls vs stx

c/o does not cut through genes because they came out
in general different functions if separate and discrete
Anotes pairs heterozygotes (cf. D's; heterochrom effects!)

know nothing
about genes, only
mutates.

Confusion of classical test of allelism
① close linkage related functions ② "complementary alleles"

propose to
leave out
word "gene"
define allele
differs to
complementarity

(ground!) What
do we know
about
functions

"M Mitchell phenomenon" $a+/-+b \rightarrow$ tetrad c/follows
 $a+-b \quad a++b \quad A-+B \quad A-+B. \therefore$ noncomplements.

Are complements always

found if homolog c/o
first. mutates?

explanations 2:0 mis-copying
often not acc. by c/o but there
is a significant correlation.

6-strand
-over?
can explain
mutates not
c/o.
effectively close
doubles.

Assumes c/o is related by requirement for "heteromutation"
proximity. Why not c/o as restriction phen? = Evolutionary effect
[cannot leave out this hypothesis] = 2 homologous effect

"Heteromutation" may be distinct from randomness of c/o.

oppose Beale's suggestions: gene = functional unit

what are means of
defining these!

Mutational sites = sites
"heteroalleles" = nonalleles
nonalleles = "alleles"

hard

if F^- F^- should yield F^- HSR
and doesn't

Yarofsky E' su

B. hyp.

most E

see may inhibit defect

↓
↓
man E

many questions involved in synth each enzyme is direct consequence

"unable to suppose that gene acts via RNA as template" not yet v good evidence. + don't know whether enzyme "is modified or is absent" : (argument is unclear)

also quotes blood Hb's evidence on allelism not good. but accepts it!

But Thalessemia is not allelic & sickle cell!

quotes Ford Symp. on 1:1

[but. in indefeatable

"genes always allelic"

∴ behavior total gene suppression assoc i single gene!

galactokinase

{ contra su/E

eatig vs breathin

cor. genes cytoplasm. how do you tell.

Emil: primary genetic information in all living forms

By NA < DNA RNA

it is in host cell & DNA, protein.

Why RNA cause of point? Ditto for DNA

not necessary to assume c/o is intra-genic in higher org.

3 contains
< 15 content
no.

(not necessary to assume it is over exogenous)

But only when observations are available can one detect the exceptions! May result either in exchange or non exchange

Res

Cytosol results ① constancy of DNA - synthesis only in relation to DNA rep. ② histone also constant ③ protein + RNA - more in active cells, variable.

EM - polyphosphorylation - fibrils $\begin{matrix} 500\text{\AA} \\ | \\ \text{OO} \\ | \\ 200\text{\AA} \end{matrix}$ which are hollow.

no other structure. pres exists of NP. of TMV

fibrils not affected by N/Ase.

At periphery, fibrils move apart

long differentiation $\frac{2}{\approx}$ degree of u. time at septation!

Maxia - units are $4000\text{\AA} / 200\text{\AA}$ - sites of 2/0?
similar units of N/P are intermediate causes of infection

multi-shoulders of chromosome \rightarrow problems of all new DNA

Preparation of periphery as the bridge of fibril to chromosome



Middle aspect:

Disc 11: 30 Apr Tues.


Aspect - [Maeshale - thinks too much reformed away]

Pis - sequencing of Feigenbaum total amount is small but ~ species.

Butter (Pavani + ...) ; nearly 1 base a 1 P. 2 bases 3 bases protein restriction

Williams: — i as phase is  Fe 

of Maeshale in relative sty. of NA, protein.


empty or NA?

matter of fixation: i formalism non-contraction?

Bermyer — "gene" is missing — gene-enzyme as a definition.
function: matter of taste position test.

explained!!!

automotive analogy belonged to Detroit comment that seems

unit of recombination: {existence} < map in cm. physical distances

is a continuous series of maps generated by experiment.

evidence that there is a symmet for cistron
is there a linear cistron?

trying to exhaust mutants in a symmet.

r_{I-2}	$S = "BB"$
r_{II}	r
r_{III}	+
	+
phenotypes	
+	+

The groups are unequal sizes.

r_{II} in two cistrons + no exceptions to square.

A = 40%

B = 20%

stable to insertion
Exceptions: apparent deletions
are always linear single elements?

Self
can also be
crossover suppressor
of asynapsis ✓

Importance of exhaustive analysis —
had its origins possibly in Stahl's study

prolo Saxon equiv.
unit }
ret } bit
fit } bit

Concentration as deletion class 3. 11000 total

150 included + listed in pairs
one locus most frequent; 2 phenotypes
(1 deletion)

many singles, w/ 0's.
1: not yet "run into the ground".
about 12 sites so far.

Importance of identifying
linear alleles
complex loci
confusing

overlapping deletes found in some cases. Can be used to
subdivide.

nonexceptional to linearity yet,
mutas + reton = or $\frac{1}{2}$ (multitalk not
excluded).

probably pairing
rather than
deletion w/ w/ of
lac,

S.P.

generate mutants from del x +? same byproducts
c indicators (same against + byproducting?)

H.H.

occurrence of -- as complements to ++.

heat shock ^{infects} < conversion, no effect as rec of
reduced markers.

" of mutants
> conversion; negligible effect as c/o.
to as high as 4%.

~~ABC~~
~~abc~~
~~ABC~~
~~abc~~

expects conversion in -X.
(conclusion of reciprocal crossover).

non-reciprocal crossover?

Roman confirmed Lindgren on + --- asci
and identity of the (-).

got didmenton
Lindgren

6
glossary
7/11/15

Waddington - definition: subset: prof. of specific
disorderly gametes or telomeres.

general "homeostatic principle".

↓ mutation: s.l. - any change S.S. - v19 sec, change -
"hereditary modification" as broader.

cytoplasmic = extrachromosomal transmission.

① metastable concept
② study state concept
most observations on sexual heredity fit ①

but not satisf. for differentiation he says. But gross partitioning
not held up. ∴ suggested that cytoplasm > cellular heredity.

then new work (Buzgari, King, etc.). But: @ sexual @ distinct
in time @ regular predictable pattern @ integration ②
most data are embryonic.

why not apply
study state concept to the
nucleus

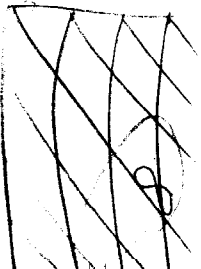
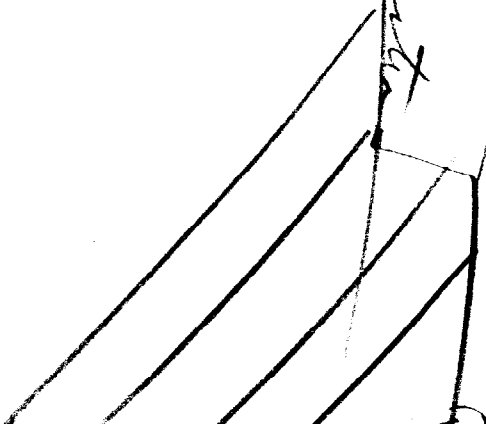
Rhoades - point mutations
extragenic: minute def; rearrangements/PE; PE i do
c/o of components of a "compound locus".
intragenic: "true gene mutations"

Some families have some defect!
But correct the typo!
and a dictionary; with them
glossaries, with paper

Send Buzgari
Ford paper.

1 - Planes - not separate
why not? why by - multiple (etc)

with



M11

Mayia

spindles.

Digitonin as solubilizer - sea urchin eggs.
did not work in KCl solution!

Dan -
look for search
conjugates in
flagella?
1 mg/ml
cysteine 14 DVO
to reduce water
cytosol evidence
of -SH accuml.

ATP - 4/100 (natural level = 5×10^{-3})

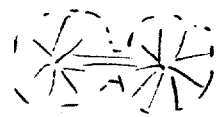
98% protein: but not ATP as formerly thought.
- 10% of total cell protein

nucleotide of spindle form not RNA - no pyrimidines. May be
ADP or ATP conjugate, i.e. nucleotide-protein (= ester)
back to Raphael.

Mitochondrial nitosis - centers may influence orientation of chromis -
bonding agent may be differing from them.

core extract + crystals of cysteine give model asters and spindles.

Where does this protein come from?



soluble proteins in sea urchin eggs by u/e. techniques
- a component disappears during meiosis.

Hoffman-Berby: glycerol 4% + ATP:

elongation of spindle & shortening of chromosomal fibres.

mirrored nuclei - Alfrey
Hepner

25M sucrose $CaCl_2$ - low speed blender + diff. cent.

25% centrifugation in whole cells. Study AA microtubules.

decarboxymethyl, PPA had no effect on ^{little or} chromis microtubules.

chromis had no effect on M - tubule.

center one abolished tubule!!!

5-b-Cl BzPhib

anti plus D effects. inhib of tubule effects minus.

Na^+ is required contra K^+ !!!

opt. source is .25M 10% deviation → 30% inhibition!
secular variations from annual to annual (hormonal?)

F&T, history, analysis in blunder activity. Old nuclei do not
rest overnight.

- Specificity:
- a) of d, l - alanine* ~~not~~ flooded with unlabeled d, l. etc.
 - b) ~~labeled~~ unlabeled nuclei do not show*.

"not exchange" — not proven unless substantial
replacement had occurred! ~~not~~ ^{replaced by Nasas.} ~~replaced~~ ^{not} ~~replaced~~ ^{replaced}

30mg nuclei .6x alanine* / hour.
∴ replacement is 8x NP/hour

DNA removal with uracil, then platinum (very platinum)
cells are unaffected.

Radioautography is homogenous — low electron carrying
"from whole slide" — was pt/pt autography.

Supplementary DNA. (DNAse not removed at first)
but this not so successful

but 4x more by "adding DNA"

Specificity of DNA?

all "denatured DNA" — just as effective

"spermin acid" had some activity.

deoxyribose phosphate

DNAse treated — core
deoxyribose

} both features +

RNA also +

RNA mononucleotides etc. nr. 9.

dinucleotides: AA, AB, AC, GC, 4-9.

Sol) Template hypothesis - "modern theology" → DNA, RNA, protein

now? either DNA or RNA

azathymine?

Whole RNA synthesis 4U, azathymine, thymine analogs - no effect

on any synthesis

Whole RNA synthesis - mutant variants; analogues. inhibits enzyme synthesis.

Chentremme: RNAs accelerates when catalase is induced

Vollmer - T23 cells - spec RNAs?

? Is RNA a template? Need a sub-st. rept.

"I would have preferred to be injurious than convergent" (of Cole) injurious no longer pays off.

cell parts expts:

Cole: - RNA fractured cells, restore by intact RNA, species - sp!

Sol's lab: mixed steps: activity 9-9. mixed as preliminary. + would like fossils to expect a water-soluble pupa to work. Thought to use protoplasts

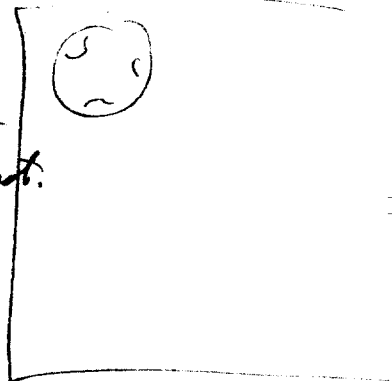
Protoplasts:

osmotic conditions: dissolved by 1, just; collected all DNA = prot RNA + membrane prot.



what is membrane? semipermeable?

"fairly good" quant recovery



Protoplasts: enzyme synthesis under right conditions.

Enzymes had little effect as a rule, but a few pups did show effects. Age matters. All protoplasts are unresolvable → 30 mins 4-9. Resolve whole making

(Do these in Ford paper?)

> 99% DNA removed. No effect on enzyme synth
and same structures.

∴ inject make "strong statement".

But acid soluble DNA still present (60% remains!)
System was also very labile. ∴ studied broken, protected.

Many methods tried: importance of initial splitting medium.
sucrose, phosphate n.g.; used a K, succinate system
then osmotic shock.

1:3 substrate/volume
1:5 " " control!

900 uM / mg P / hour.
what is enzyme activity?

Homog pellets through pipette are centrifuged.

dil ~~was~~ c 4.2 - 4.7-fold. (from what?)

resuspended loses 85% of DNA

RM = reading mixture. substrate & substrate in a 2.2 phosphate
lose DNA.

what are "substrates"
Nucleosidically??

genes ^{read} RNA common factor fractionating
acid soluble NA now absent.

∴ Substrates s/DNA make Enzyme
- RNA do not " " .

Is RNA being synth. in induction medium?
from 4-118 r/10ml.
c1 - 44.

40m log + linear synthesis.

What is "DNA" - acid pptable Dische reaction
hydrolysis \rightarrow UV absorpt. exp.

Examined other capitates & get consistently
synthesis of ~~the~~ RNA, DNA + some proteins
synthesis. 5-6 x more in all capitates
make more DNA than RNA, prot.
10x. at test es by culture.
but RNase-treated are inactive.

~~RNA~~

nucleoside triphosphates.

Put back RNA. — Ochoa substitutes restore EF ability.
enzyme stops effort.

not specificity.

Did you say - RNA cells can
be restored by nucleoside triph.
do they make enzyme?

Ochoa: ~~RNA~~

if so, is
the template only
protein
"what is DNA" —

poss. equilibrium of limits?

What are products morphol?

relate to Alfrey -
restriction by NA small fragments
of low specificity.

but not all RNA
was removed.

> 50% removal n.g.

how much tall dephosph.

DNA is nonsense DNA.

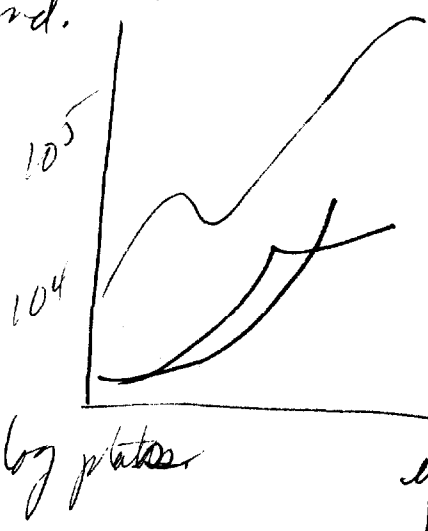
Harratt

Radiation effects on TP. Target theory negligible ($5^8 \times 5^5$).

1. Quantitation problems. Reactants a x b. Disruptant only about 15 mins. a is impure, containing unknown infective molecules. Synchronous development of capsid time in exptl. cultures. Level from 0 to 10^5 to $100 / 10^6$ bacteria. Later waves may also occur. ^{M.H.H. 2.} Can add DNA from start and get higher assays. Linear assay possible with ③.

Relationship of capsid time to division cycle.

There is a final \times eff. depends on number of capsid bacteria. Limiting reaction evidently bimolecular. Had to study kinetics at high DNA, plateaus are formed.



What are reasons for nonlinearity? heterogeneity of DNA? Extra DNA gave 2nd rise early plateau, decreased by progress of unmarked particles. 1st plateau is of total DNA. Break due to inhibition. At high multi, inhibition is partly washed, may be max. of efficient systems.

1st plateau is region where doubles begin to occur detectably.

? - gene imbalance? - Transformed cell may be killed or not transformable.

must be sensitive + resistant particles

extra DNA in phage may have a similar function

[polygeny?]

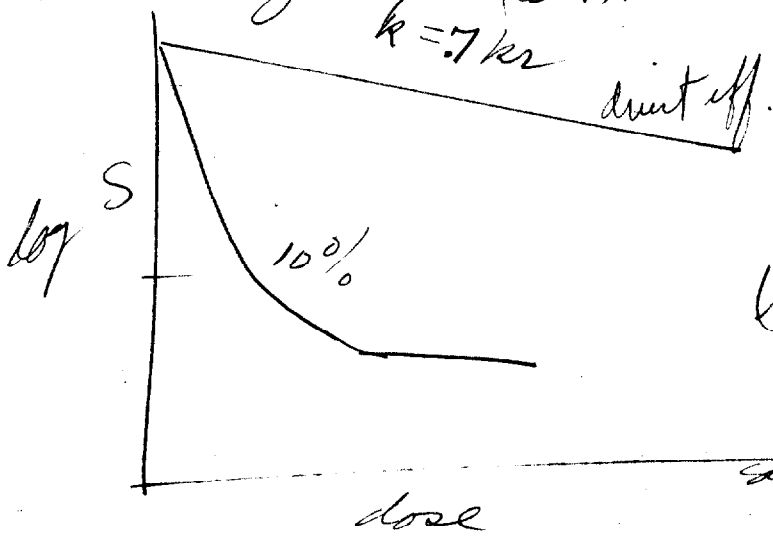
Radiated DNA -

low bio assay at 3-4% survival.

~~radiated DNA~~ mass of DNA determines bio activity.

inherent effects problem - radiated in frozen state, where protective substances have no effect, & serum volumes are smallest. i.e. susceptible to inherent effects.

Kinetics of inactivation ($\approx 4 \times 10^5$).



Mann + Fibre
($2 \times 6 \times 10^6$)
only 1/10 app. obs. m.w.

(radit) radar & center. ? Residual activity possibilities.

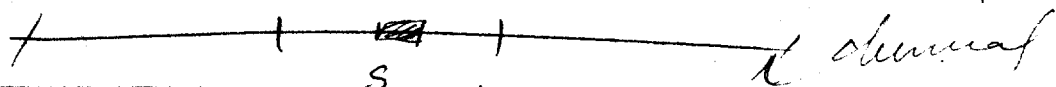
seeing DNA not heterogeneous as clones.

Distribution particles? Fractionate DNA $\left\{ \begin{array}{l} \text{high gravim high TP} \\ \text{low gravim low act.} \end{array} \right.$
 near no effect no difference.

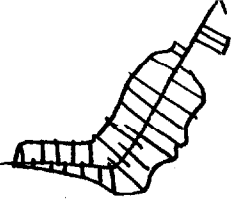
more sensitive units are present in same populations.

Why are small aggregates > sensitive?

postulates a sensitive region for S^R locus (guess 3×10^4 nucl.)
 target prevents incorporation.



Aggregates: ? energy transfer or cross-links which inactivate.



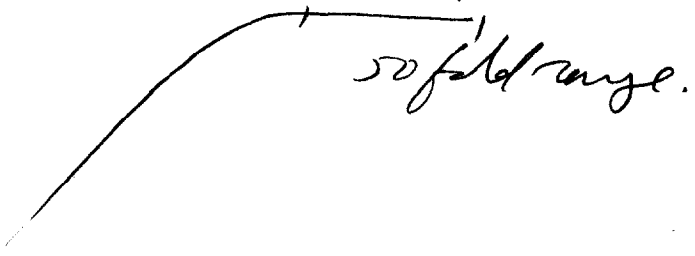
Hotchkiss: assume \times hypothesis.

Quantitative factors

Capsule system - originally by delution. \times calculations: sediment too soon or too late n.g. \times incubation, entirely must be specified.

If sediment is resuspended, may let several add. \times s. pres. matter of relative conditions. \times s. So moved to dry existence.

Rollin doesn't see plateaus



level of plateau depends on proportion of S^a/S^s DNA.

Pendick - chromatography of DNA \times mean \times ch.

put #'s on word DNA/cell DNA/molecule \times max. "molecules" / cell.

Ectocolla cellulose -

Variety of fractions tested

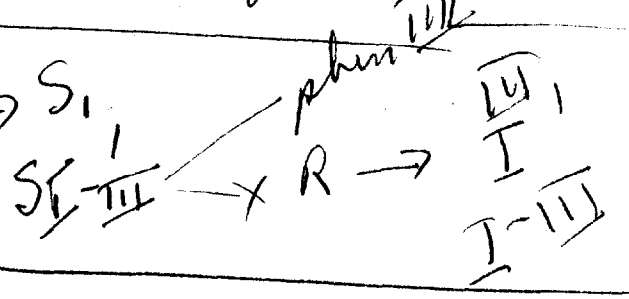
least eluable DNA is most active

2M NaCl MNH₃. \therefore heterogeneous

In comp. exps, can get intra-plateau?

sp. signature?

Acetabian: $S_I \rightarrow \times S_{II} \rightarrow S_{III}$



H. Taylor - Quant. assay.

DNA is not exhausted!

What is being measured in essays?

∴ not a stoichiometry + assumption of a time slice of a bimolecular reaction + kinetic justification

Zamenhof: The transforming principle.

① test for activity of DNA, as in physical analysis.

② composition - modified by Beltharil

Effect of BU uptake: sp. mut. rate not changed, cells still alive
But do get two colony types, incl. "purpoint colonies", but these → large by mutation. + can establish pure lines.
~ 10% of such mutations.

large colonies also gave 2 colony types. "Unstabilized" the strains, every colony's tend to be altered.

③ Nature of heterogeneity?

④ Replication so far.

⑤ auxilary system.

-x is synthetic medium?

$$2 \times 10^{-9} \text{ r/cell per molecule} = 5 \times \text{content of cell.}$$

1 cell has 25^0 molecules. @ ? groups of characters not DNA @ 1 mol
> 1 determinant (i.e. linkage).

attempt to decrease size of DNA → mut.

synthesis problem

10^{-4} r DNA destroys " 100 r/ml DNA in 20 minutes
duping, dialysis also " cut & suppress by prechopping.

assay methods for these procedures

denaturation 5.2% NH_2^-

didysis: loss of heat stability + denaturation.

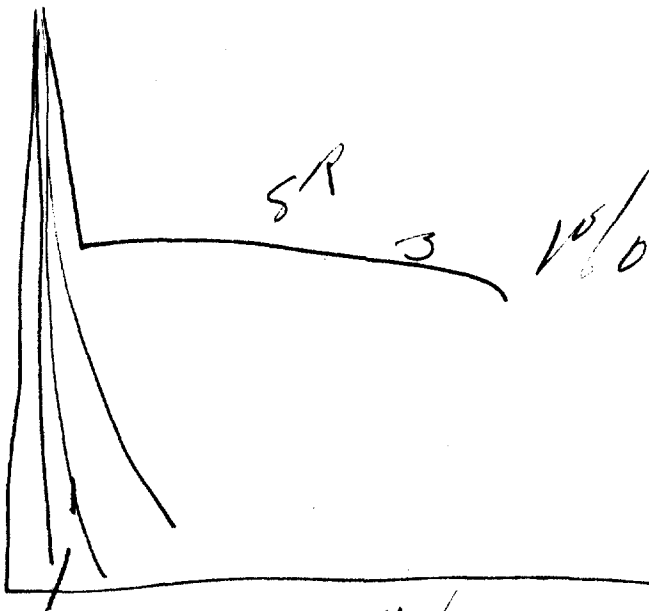
M. tetraepus:

diaminobut Me. 204 mutation is guanine. also ester for PCY^3 .

diaminobut - much?

∴ DNA is target of strong mutagens

UV treated, heat treated all destabilized.



assay systems for DNA

Bring up again.

X mutant both + DNA constant but all ~~mut~~ genes is a protein, making pu. Kern. coli? Neisser

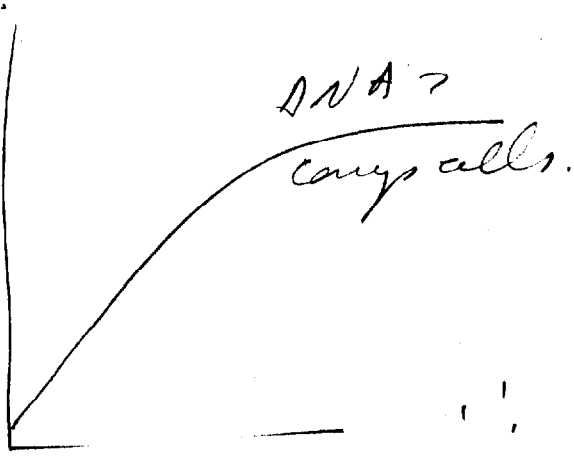
importance of

- 1 capsule e UV differences → X
- 3 = SR
- 2 capsuled V 2 similar
4. SR from type 2.

A to pms W. — Cell incorporated after cessation of growth + no increase in DNA/cell. ∴ exchange without resynthesis of DNA

Goodgal: 4 DNA 300 mols/watson → cell's with

is uptake homogeneous? Yes! from 0 - 90% range



Sd cells do not x
do not x DNA.

1. gene per cell.

Assumed $M = 6 \times 10^6$ (i.e., $10^{-17} N_A$)

count - cont - diff values = 5.5×10^6 , same for S, Eryth.

No separation obtained at up to 4V

Sordal coli cymmet + ATP + Mg + light
restored UV & TP from 3-30%

Linear
had. may suggest
a substrate

says Hemophilus, pneumoc. do not photo-reactivate prese.

Luria - ? as phage. - no.

Bond's Venues early fractions have $M \approx 400,000$ even
(pairing minimum)
ostenditels

Lidgy intrus periferi (a la Schaeffer)

	10^{-6}	10^{-7}	10^{-3}
infrayae	x	x	xx
paramfrayae	x	x	
suio			

SR → x
homo ratio is
hetero constant pro
gives p.p. + viral
of strain of donor.

E. Salmonella
+ coli %
Luria

"crossing over" —

13 seen in heterozygotes

4 steps in heterozygotes

v. occ. recurrence. isalleles for reversion:

also differs in recomb. frequency!

some mutants give confusing overlaps.
heterozygote.

Kalchkar

Only 2-point in

only comment is that
detailed data have yet to
be published & therefore
actual comments not yet
available. There are
difficulties in analysis
chp. 2-point tests

Non-aggression pact

Hewitt - general review of label work in re DNA effects

Hartman: - classifies recombination by vectors

DNA, phage, cell conjugation:

a. DNA b. Transductions; lysogeny c. cell conjugation.

Types: generalized vs. prophage-like.
stable vs. abortive



S^R - not transduced in T4D?

Induced lysogenies are
- x in Salmonella!
generalized.

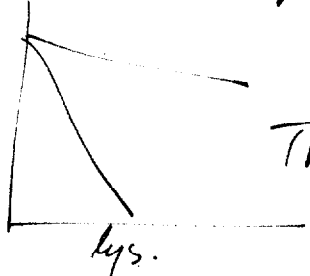
- Zinder etc. Expts.

Protein hyp. - attachment specificity?

Jacob: gal by heterologous phage \rightarrow h^+ : induction can be lethal
Metabolic state of recipients: need phenomic expression: or effect
of host - background on eff. \rightarrow

Chemistry of exogenotes - analogy;
p32 decay:

no direct evidence.
evog \ll prophage.



This is Salmonella.

at least several genes involved. - "20-30 bp long?"
(not sites?)

incorporation step - possibilities: "unilinear" $F\lambda^+ \rightarrow x F\lambda^-$

Jacob: eclipse to lysof. cyclic behavior
temporal quality.

prophage = phage mat in noninfectious state.
synthesis coordinated w/ bact. div.

expression: → phage + immunity.

induction w/ λ → → phage.

immunity usually specific for related

superinf. phage does not interact genetically except on induction.

? chemistry of prophage - Hershey & Chase
injected DNA, but no data on whether info is transferred.

How many prophage/cell. Double infection, manifests in output
have input ratio. Induced bacteria superinfected behave as if 3
per bacterium. ≈ 3 units.

Genob - also competent between related prophages. usually no mixed lysogenies for +, mutants of same phage.

E. coli K-12 L+T. λ ML- λ^+ . can cross $\lambda^+ \times \lambda^S$.
Astrupts by L + E.W.

lamis "never visits Xyl, Mal, etc!"

14 phages of cross-immunity. 7 are inducible A
7 are non-inducible B

A are ~~delet~~ to λ . better Gal and R
all show zygotic induction, degenerate in same order as time of entrance.

B: all are delet to R - most are linked to

I L H₂ T₁ Lac Gal X. Xyl Mal S

no zygotic induction in these phages.
 each phage has fixed site. two general regions.

one of the B's →

3 prophages can be → by no other.

phage crosses among A's.
 3 c mutants



specific minority site = localization-determining sites, near C1-2

caricatures: depth by in.

Myer terms.

F-locat

8/15 been mixed from (mixed RNA) proteins.
 none mixed from mixed virus.

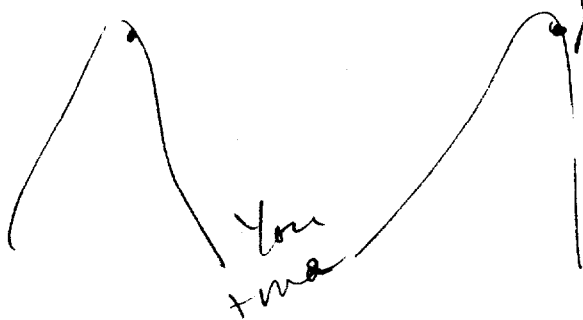
Schramm

↳ L. P. ...



Davidson

f. H. mod.



most modification in
 chemical terms.
 any pure clones
 1st TAV
 Volkmann et al
 any effort to
 stabilize RNA
 with normal
 plant proteins
 essential course.

is the prophage factor?
 hairy - an ensemble of
 prophages?



Buy
Ch+
Law. How true conservative
exchange reactions?

Doty - plan of mountains.

as ev. of section
is to show the
clouds.

Chargaff: language difficulties - dictionary lacking
in coil ≠ mol helix

MC, AMC
N-He-Adenosine.

base composition: 4 bases in PNA - AGUC
4 " DNA AGCT

plant viruses may be typed. In animal PNA close association /
protein.

This is the
fundamental
question
+ answers suggests
represented more

"NA can carry information" → → → Is protein important?
pres. as nucleotide eq. Any finite sequence represented more
than once. Is matter carried many times?

maybe other than 3-5 linkages; i.e. as folds.
not included that NA is huge cycle

$$\frac{A+T}{G+C} \text{ varies from } 0.4 - 2.0$$

Tb sea water

6 genes = 6 bits

But $A \approx T$ $\frac{A+B}{G+C} \rightarrow 1.$
 $G \approx C$

Meaning of regularities.

-Chayoff - unitary components well preserved.

now? sequential analysis.

Clearly G-C relations in terminated tubes.

can fractionate to small extent RNA's for different G/A ratios.

~~Is~~ Is G/C / C/G random. ~~but~~ occurs fractionally.

clearly not natural genetic content of T2?

PNA - no regularities, but no precise prediction at first.

code nucleoproteins: the skeleton. $A+C = G+U$ was

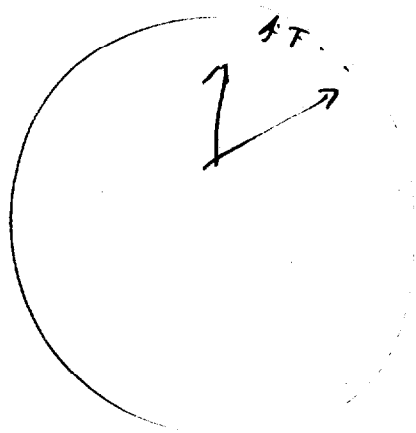
preserved, ~~i.e.~~, but no others.

Smith - no good evidence on branching of RNA.

see some problems.

Whitfield - remove phosphates
oxidise sugar

remove phosphates... experiments and pure material.



Thayab
SBZ

Oct 22
Oct 23
re print

Smith

aggr. nptatae in B. cereus cells.
not homogeneous. cyclic and turnover faster.

Crile SMC does not go in at random. Agree that naive repl. numbers is ruled out & maybe unit of agglomeration as not nucleotide but analog nucleotide unit.

SMC is purified must to guanine. suggests repl. principle.

~~analog~~ - ^N-methyladenine, replaces thymine.

He suggests that XT replaces TT pairs. but gives hemidores

(flexibility too much)

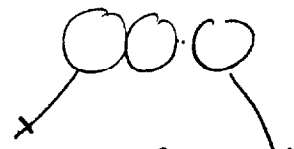
stiff/compliance

Touloune & Zam.

Oct 22 - whole cell RNA ~ enzyme RNA.

Burdick - recorded deviations from 1:1

Crile: must have Pur: Pyr (large: small)



X-ray picks out the regularities. and for given room, G-C A-T fit best.

maybe some unpaired chains in some pyrs.

If all 4 bases are in any chain, only w/c model so far fits the X-ray.

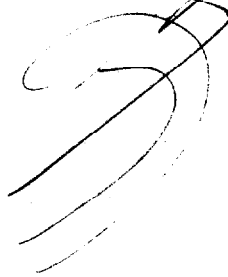
Not yet critical professional judgment on key concordance
but no measure is yet.



edit is paranemic not pleonemic. No state of model yet, at
least (Why read SF if you can read PNAS!)

nucleoprotein. prot fills in the "main" groove.

1 side ~~chain~~ per $3\frac{1}{2} \text{ \AA}$. not stay lit up.



as 3d chain, basic groups mark high P.
non polar spacing logs of 2 or more

∴ non polar AA should occur in pairs.
much analytical detail fits that.

nucleohistone

JA

synth poly RNA, 4A6UC. \ or AU

algae = bad photos

poly A - ~~is~~ clear photo

card RNA ~ DNA specific base pairing
except A:A

Kelebas what he's
publishing
Med concentrations of
analytical work on
biol significant
material!
Substantive factory?

Largest issues are

can. org. organisms based on
feasibility of manipulation methods;
but ~~through~~ like eggs.

① Availability of test systems for DNA, RNA.

② Concentration of analytic effort on bio assay systems.

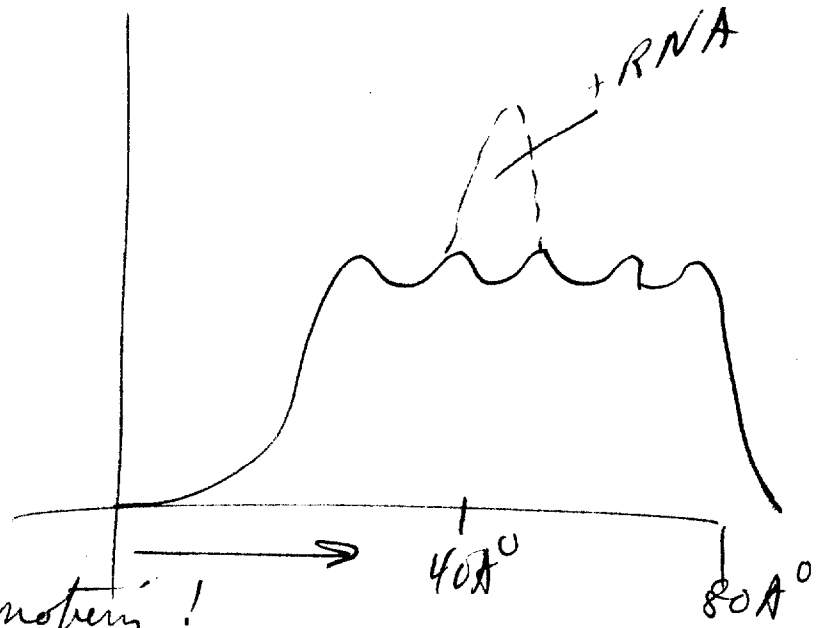
specificity absolutely essential.

Phase
Transformation (near 100% to
cellular
DNA-mediated)

Beck - mixing poly A, poly U interact at 1:1 ratio
 as measured by specific absorption. as well as sedimentation.
 possibility of base pairs as well as other.

Franklin

density diagram
 of TMU
 and TMU.



∴ not core but RNA is
 wound into peripheral proteins!

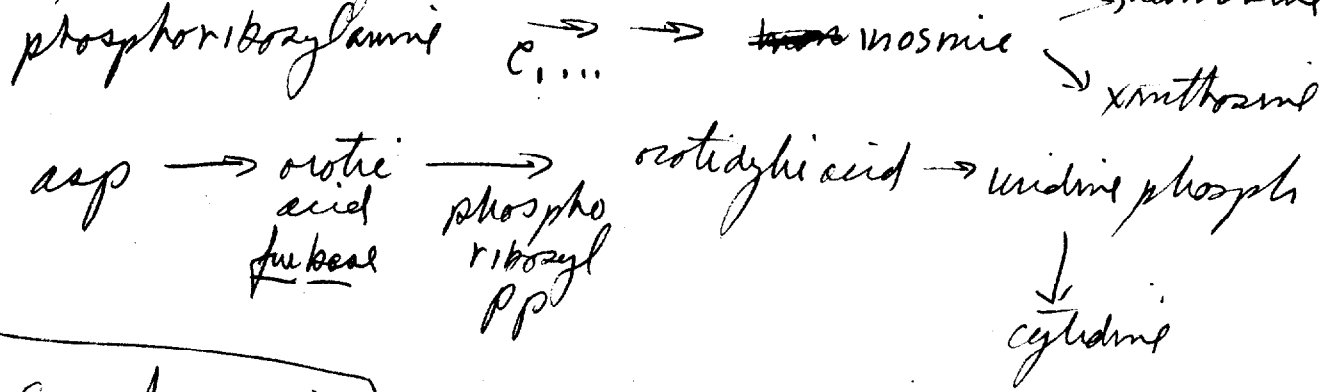
- probably single strand if fully ordered, not em. RNA.

low: ^{many helices.} globular proteins - any here?

Art Kornberg: purine synthesis

- amino, basic, conjugate, valuable

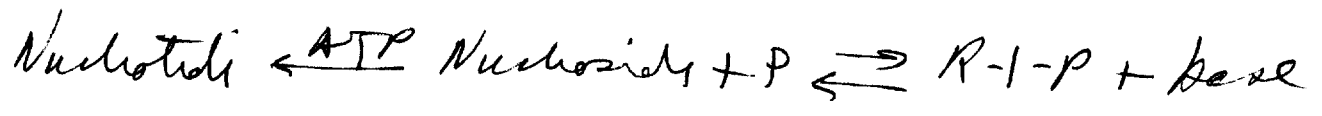
Enzymatic synthesis of nucleotides:



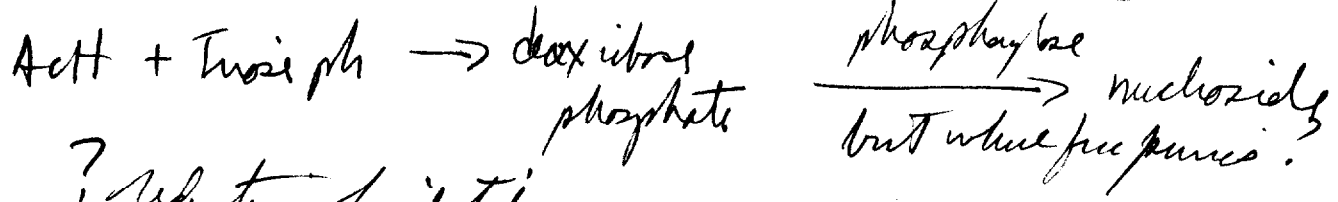
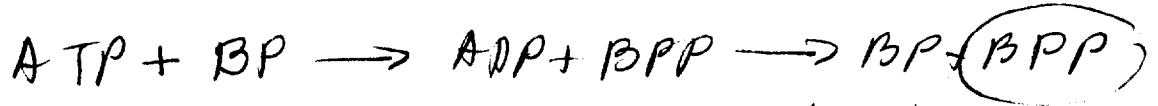
I.P. no free purines or pyrimidines

C \rightarrow uridine typical
 L. acabinous water though uracil \rightarrow uridine phospho.

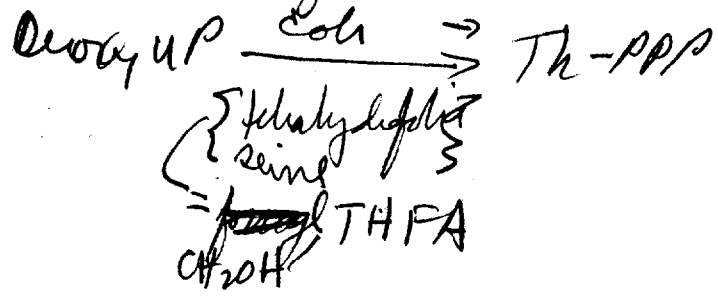
"salvage pathways" HX, G, A or U $\xrightleftharpoons{PRPP}$ nucleotide



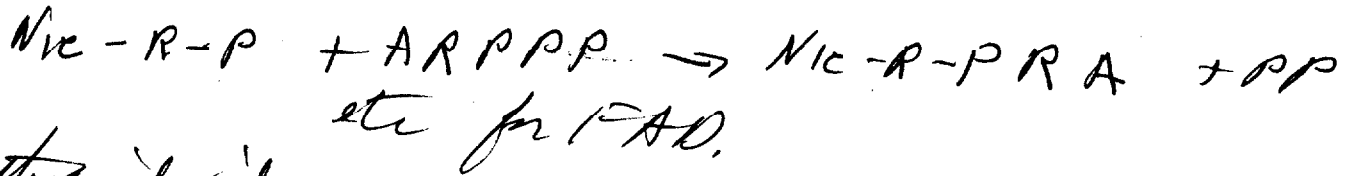
Nucleotide $\begin{matrix} -P \\ -PP \end{matrix}$ $\left\{ \right.$ would be further synthesis Kalcher wanted.



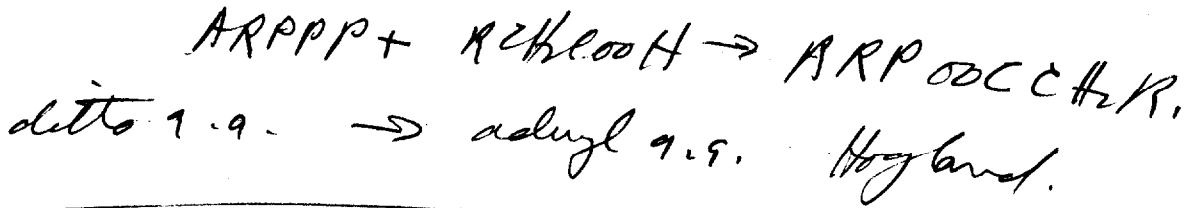
? reduction of ribotide.



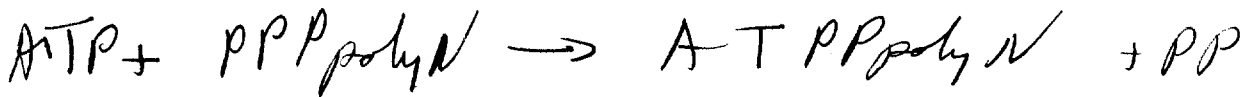
of coenzyme synthesis
nucleotidyl-enzyme reaction.



phthalic acid.

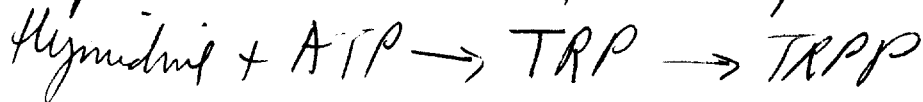
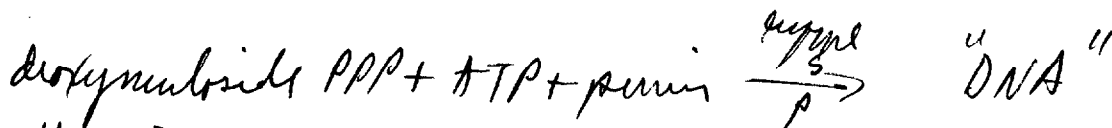


? polynucleotides



C. 4 ATP \rightarrow NA. Then ^{Aristobacter} ~~Actin~~ / Littauer purified a coli enzyme
made NRPP \rightarrow "RNA".

DNA:



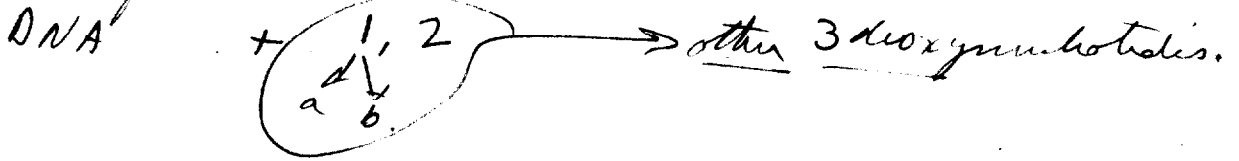
TDP is sufficient as TTP.

mixtures of these TP more active than single
amounts of added purin (about 1%).

All 4 together do react

purin

DNA - purin heated, rich in
cyclic ribofructan ↑, rich in DNA Recently purified. Better
come to nucleic fraction + deal.



now need all pur components

ATP not used. Suggests probably phosphorylase
not needed.

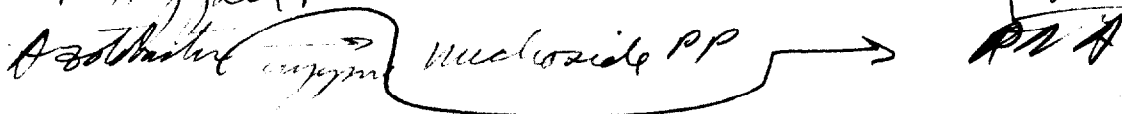
Now try net synthesis: 30-50% mass.

The RPPP is purin.

charged/linked: coli { + Hepes { T2 1/2	with specificity of purin DNA <u>is purin</u>
--	---

could bioactivity increase if pur DNA were used?

Lehman + Heygel.



"polynucleotidyl phosphorylase"

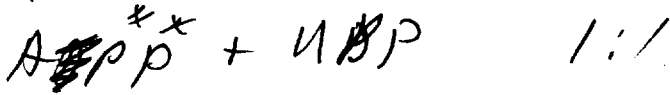
has made ADP → polyA, G, U, C etc. also

Ad, AUC

rather had shown the other phosphates in all.

? structure + nature of polynucleotides - structure w. is RNA (Rids)

poly AU.



11mg AU_n about 1:1

products of RNase. phosphodiester attacks poly U and yields poly U ends.

U 38 AU .35 AAU .20

AAAU 4.9 AAAAU 2.1

50% of #X in UX symm is A,

takes advantage of p* trans from A to U at synthesis. fairly interspersed A, U, in polymer.

? symm is "random".

Meaning of random.

random bias phylogenetic book.

? Intracellular route. Means of orientation? It is known very low. Km 10⁻²; Hill slope < intermolec. Maximal local sites of NPP formation. The enzymes are unspecific - e.g. polyHX. also unspecific in phosphodextrin: also bacterial NPP, unspecific. rates vary. dihydrochloride of protective as 5' 2' OH PP. also inactive as above 5-pp

Now general path is clear, but specificity is still dubious.

try virus RNA

Mechanism: primer? Most polys may have some nucleotide. possibly not.

~~Needs~~ NMP as start of chain? ~~ok~~

But AMP + UDP → no A in end groups.

? one or several suppresses.

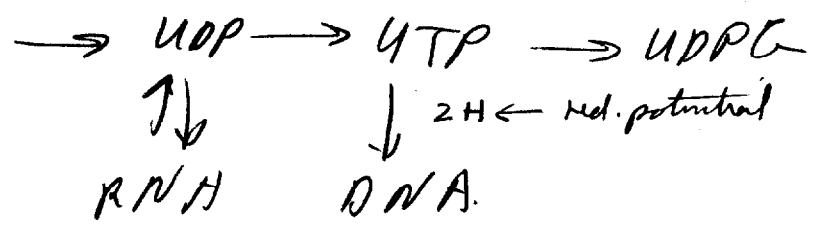
is RNA specific
etc. in plant viruses
change of standard spec.

Potter: ATP → RNA ADP → RNA.

growth control. Thus ATP inhibits RNA formation.

extra biostability of DNA.
possibility of Hornby ex standard.

homogeneous labelled RNA does exchange activity } no net change
" " " does not }



Hornby contrasts:

	RNA growth	DNA
Reversibility	+ inhibits	(-) but not inhib.
K _m	10 ⁻²	10 ⁻⁵
diversity of N	1	4
promin	- ?	+
extent	∞	7 1/2
specificity	""	ATP not usable.

Cohen - correlated systems.

point strongly made.

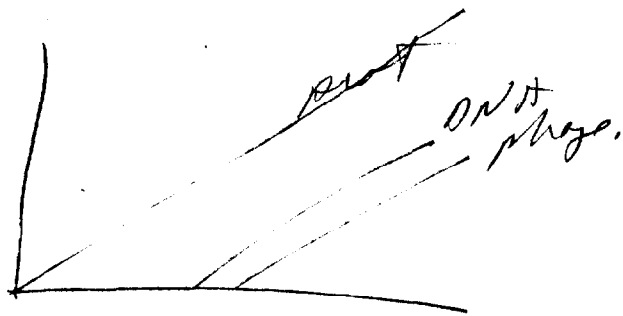
RNA apparently enters as energy source?

reduces almost to E. coli and phage

Tevens: HMC and GMMC

1. may be necessary to ex. hypothesis
2. inf. cells stop new enzyme synth.
- 3.

do of hosts?



- Start: transfer DNA inf to protein as 1/several poss.

early S35 does not major in T4 strain.

? repair mechanisms

5MT inhibits DNA synth. only if added at 1/2 lag phase.

if added later, \rightarrow DNA s/ further protein synth.

Rate of DNA synth \propto time of add. Ciba.

? role of protein: inform or logistic support.

Some small RNA is also turned over

Do B growing, Hypermur, Hypermur \rightarrow DNA.

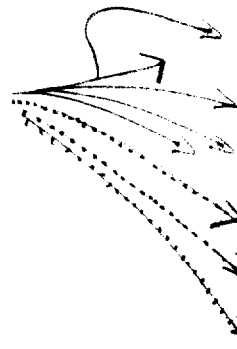
pres. Hypermuric acid.

Defected B. Hypermur \rightarrow DNA.

(cause of Bombay shunt)

ribos nucleotides \rightarrow deoxy nucleotides

Hypermur analogues don't inhibit B.
do inhibit virus synthesis



Do Hypermuric. in wild L15?

Amino acid \rightarrow methylating groups.

Be hail insects itself

Now + Sz. mut rate indgt; de novo vs. admiss.

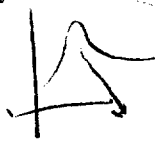
\therefore metabolic control of genetic material

Zamiatol Bell effect - "exchange is DNA"

Adelby $T^+h^- \rightarrow T^-h^+$ 100x adtest in defunct medium.

Control of nucleotide pool a key to mutagenesis.

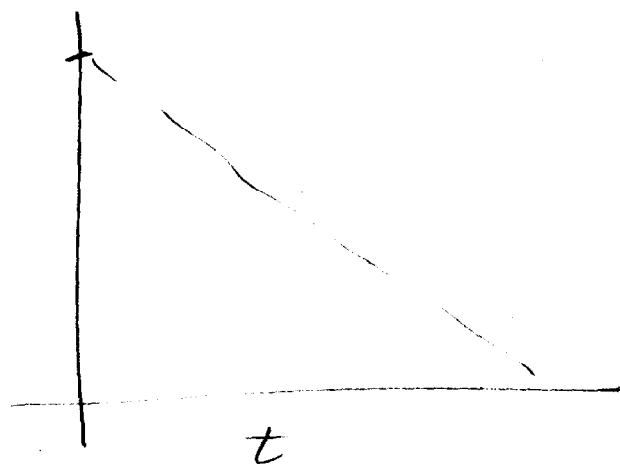
turnover is manifestly rather than whole RNA
mutagenesis was manifestly
Vollstein
shape of pulse
evils



Vollstein spec. ⁱⁿ granule fraction. In pulse exp't showed RNA, not DNA turnover

~~abstract~~ Suppose :

what's the suicide curve of T2 grown on hot bacteria, cold medium or reverse. (i.e., what is distribution of the bacterial DNA in progeny phage. ?)



Stent:

rotas -
unwinding by breaks + reconnections about every 5th.

separation c/ replication → dispersive replication:

other schemes are conservative. hyp base pair.

c/o affects dispersal depending on mechanism.

grow hot phase, $par_1 \rightarrow prog_1$
 $par_2 \rightarrow prog_2$

How does protein disassociate?
C-O can still be a copy mechanism.
Does this mean second step is still semi-dispersive?
has 4 understand I'd expect.

~~W+M make~~ W+M make
mountains
Potter
P/O conservation.
 $prog_1 \rightarrow prog_2$ still 50%
→ $prog_1$

Hesley thinks just inefficient.

cy: individual autoradiography

BS: P₃₂ inactivation method $prog_1$ - inactivation assay, as ability to ~~maintain ability~~ to transfer P₃₂ to progeny.

cy: } big pieces Bernth $\left\{ \begin{array}{l} 1/2 \text{ is distributed to } 10-20\% \text{ into} \\ 1/2 \text{ small pieces.} \end{array} \right.$

i.e., same dispersion of parental atoms.

of cy: Both methods as $prog_2$ < Bernth same fraction of $prog_1$ small pieces

bacterial nuclei - hot cells at 1 division
assumes nucleus is unit all cells div.

15 nucleosome
unit
all cells / nucleos.

Thozzi

B.?
[Maple]

obeying matrix for replication:

eg: can measure 15 cp/worth.

star size does not change from prog. \rightarrow ρ_{j2}
semi conservative replication: $2 \rightarrow 3$.

do they perceive matrix?

stars are in h^+ fraction & 90%.

(total activity?) \rightarrow

problem of bi helix \rightarrow maybe segments of open loops.

Baltimore Symposium Notes.

Bradley genes as units of CO; no CO in genes.

(Private discussion: are there discontinuities e.g. in *Drosophila* data).
Review Muller & Raffel.

Benzer made most of the necessary replies.

"Mary Mitchell phenomenon" -

glad someone mentioned Lindzen; for I dislike it sounded like the new edition of the Soviet Encyclopedia on Stalin. ^{so many farm truss}
G. Defont Sympos. - science as series of confusions perpetrated by short recodes in the lab.

"classical classroom" - mythicism of linkage model - possibility of epistasis still relevant.

"not necessary to assume clo is ever intragenic.

Ris - emphasized binding from chromosome fibril to chromosomes.

Benzer - "Advanced mutants from an elementary standard" - main objection as a manytimes demand neologist: simple & complex terms are preferable: around our lab.

'fried nit, nit and pfrit, salt melt and pfelt.

Serious semantic problem & necessity of purifying germanic since nit defno. too often abandoned.

automotive analogy belonged to Detroit.

meaning of cit, e.g. in McChintock's material.

Comment on content: possible relative reliability of asymptotic enumeration and mapping. discontinuity in *Drosophila* data? of Muller & Raffel.

Warney - steady state concept in nucleus. - B+K; L+1.

Rhoades - why not go whole hog - or is Foldschmidt too close too?

not a semantic question: only way to satisfy "point mutations" is nucleotide substitution.

Hartman Reliability of 2-point data? 3-pt. better but few results yet reviewed critically by geneticists. Questions both on the cogenes & on mapping.

Jacob - no comment now would take all day. Benzer's ud. ad absurdum: whole genome as ensemble of prophages

F. Court - @ no pure clones in TMV @ stabilize RNA c plant proteins - course of Volkmann?

@ host modification of flu

Chargaff - nucleic vs gene.

Call Dr. Gelman

10 AM Sat.
Barry - 6:50 PM
2-#29

Wednesday.

HAZIA

flagella is promising contractile units

is there cytochemical evidence of SH accumulation? dependent cysteine?

Allkey

"Exchange reaction" - whatever that means: is there any real evidence for this, if so this might be more interesting than synthesis. Did he answer Mason's objection?

DNA removal inhibited uptake - but pieces of RNA, DNA worked. Any evidence here of DNA re-synthesis à la Sol?

RNA removal studied? ditto mention?

Sol

Review specificity of RNA removal morphology of debris, mixture of substrates?

Harriet - Pollin - Zamenhof

- relationship of competence waves to fission cycle

importance of clarifying quantitative aspects for evaluating Dr. Bendich results e.g.

discrepancy in kinetics illustrates problem. What are units of activity.

role of synopsis if, after all uni nucleotide subst. is source of mutations.

auxiliary conditions in Hemophilus? Clarify growth in defined media.

Species interrelations - have some examples in Salmonella, which may be connected with phase adaptation. The datum is usually:

A	x B	1
B _A	x A	2
B _A	x B	3
B	x B	4

$1 < 3 < 4$
 $3 > 1$ may select a more compatible segment recombination best explanation.

Segregation only a transient condition.

Zamenhof may be implying a dis-integrated system of DNA molecules, - this is what Harriet had in mind citing genetic recombination.

Biol. function of "DNA" as emphasized here. Review other situations < Doivin Demerec

few papers had anything to do of course w/ heredity.
how to say what everyone does - elliptic approach.

A small group of us may not understand
his

Most genetic discussions were matters of definition; Dr. Benzer's remarks were especially refreshing. We are looking for discrete entities, either of the molecular order which we cannot assume as the ultimate chemical unit if DNA is accepted as the basic hereditary material. This is not the occasion to review the evidence for this thesis; several authors have cautioned that it may ultimately be not the whole story but at the present time, nothing else is in the picture (except RNA/VIRUS).

Re Benzer - his euphoric terms have the objection only of being Greek derivatives; in our lab we had tried Brit, nit and pfrit. Mutation and reversion are paid for a longer life. At this point would adopt his terminology's suggestions we should have absolutely no discussion of terminology, but in a mathematical sense should insist that each author start by defining his terms. Our next paper in *Genetic Control* starts with a glossary & until we seriously adopt a really useful dictionary, this may not be so outlandish and idea.

Of course we may also need existence theorems. One of Benzer's contributions is that cistrons do exist as recon segments. ~~Del. reliability of bacteriophage~~
Halterman

Drosophila has not necessarily been pushed to extremes: Benzer's rebase two analysis of a segment had its ancestry in Muller & Ruffel's analysis of the scute-achete region, a job that still needs repetition. Del. reliability of exhaustive enumeration + 2-point both in any material.

On Dr. Rhoads' talk, Dr. Spiegelman asked whether this was a semantic ~~idea~~ problem; of course Rhoads was asking the material question, are any mutations substitutions of nucleotides, which are the closest possible approximation to point mutations. - Sign of McChintala for pos. effect

Benzer to a point of Crayoff's query, which is really an old question in the relationship of mutants to genes. Conceivably a mutant is a single nucleotide substitution, but it would be a meaningless case if adenine acid were a transforming agent. The meaning of the code is not in the letters or syllables, but the words and sentences. The specificity of transforming agents can best be visualized in terms of synaptic activities known if ~~any~~ nucleotides were mutagenic in the pneumococcus.

(Benzer) →

Meeting reminiscent of the Detroit symposium - where Benzer would have been even more typical - seeing so many friends in it is me that the sc life these days is a linear sequence of conferences punctuated by short profits in the laboratory.
& informational exchange = recon

comment: plethorism

Gales natural there and Dr. Spiegelman's here had emphasized the lack of this ingenious frontal attack (have to watch out for such words. at Detroit infeasible became inderferable.) What does this attack mean?

Q We biologists need all the help we can get from the chemists & physicists? disappointing that the most advantage is not taken of biologically active systems e.g. in the physics chemistry of pneumococcal recombination. Benzer this may not same tactics but would be strict

② We biologists have to be more courageous about doing the impossible
in a essay — Dr Spizman was not denying he had been making gold.

~~Dr Majia's model system is cytoplasm should not be too hard to duplicate
in ~~in vivo~~ The RNA systems have turned out to be extremely hard~~

The transformists have reported encouraging progress in the
quantitation of the two presently available systems; ~~though the~~
~~discrepancies are somewhat disturbing, concerns in the catalytic~~
~~activity of DNA which is not essential. → I would at least call~~
~~it a success~~

There are good reasons for wanting to find simpler systems, though
no doubt Dr. Spizman could make protoplasts in *Serratia* or
Pneumococcus if he put his brain at it. The two systems now in use
are relatively unconvivial both from the viewpoint of availability of suitable
media + physical requirements. Reports of DNA-mediated transference in
have appeared from time to time but no reproducible recipes are now in
print. As the crucial problem may well be the penetration of DNA, we
may be more optimistic about the use of protoplasts, such as Dr. Spizman
has described. Unfortunately, our trials so far with penicillin-induced protoplasts
of *E. coli* have been entirely unconvivial.

The ~~most~~ direct approach at the replacement of cellular organelles also
has its place in the justification of models; e.g., Dr. Majia assures me it would
be feasible to implant cytoplasm in vivo to try to set up unsexually mitotic centers.

I would like to take the liberty of asking some questions to clear up
some points on which I was confused.

Would Dr. Sp. clarify the specificity of RNA substitution in his protoplast
system? This was reminiscent of Dr. Delfuy's ~~method~~ demucronated
media. Any scheme of DNA resynthesis here?