

6/20/56
±

J. Ledeburg

Tues. Bealle Reis / Branger, Mitchell
Narney Rhodes

McCollum Pratt Seung.

Wed 9:00 ~~by~~ Nagia Alfrey Sol / Hamitt, Polkin, Zuns

Thurs. 9:00 Hartman Jacob Conat / Chergoff Markham (w)
row

Fri 9:30 AitK, Leha Cohen / Stunt, panel.

suggested date J.S.

1. Bill Mac asked me how after the Ford Eyes - 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 biased than may appear, and I also hope the copy won't read "indefensible" when I read "ridiculous" as happened at Detroit in discussing Dr. Hornitz's paper.

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

hardly
 (~~is~~ to talk about something is hard & about - 19 Feb)
 won't transmit until could study

but Lp^+ / Lp^s heterozygotes
 { H_{2}^{bal}
 Lp^{bal} linkage }

Mr. Achon program.

Beale

mediated... chromosomes. W/C suggests mechanism

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

can't find can
studies as
mut. vs. st. mut.

know nothing
about genes, not
mutates.

Confusion of classical test of allelism

① close linkage of related functions ② "complementary alleles"

propose to
leave out
word "gene"

define allele
of function to
complementarity

(General!) What
do we know
about
functions

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

Are complements always

found of homolog c/o
first. mutates?

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

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

Assumes c/o is related by requirement for "transmission"
proximity. Why not c/o as restriction phen? = (Lindgren) effect
seems to leave out this hypothesis = 2 haploids effect

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

oppose Beale's suggestions: gene = functional unit

what are means of
defining these!

hard

Mutational sites = sites
"heteroalleles" = "varalleles"
homoalleles = "alleles"

H₂O₂ x F⁻ tal should yield tal HSR
and doesn't

Varofsky E' see

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 circular)

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

But Thalassemia is not allelic & sickle cell!

quotes Ford Symp. on 1:1

[but. irrefragable]

"genes always allelic"

∴ behavior total spec. enzyme assoc. i single gene!

galactokinase

{ contra su/E

& evolution

eating vs breathing

cor. genes' cytoplasm. how do you tell.

End: primary genetic information in all living forms

by NA < DNA / RNA

it is in host cell & DNA, protein.

Why RNA carries information? Ditto for DNA

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

3 contents
C 13 content
is.

(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

Cytorchem results ① constancy of DNA - especially in relation to DNA repod. ② histone also constant ③ protein + RNA - more in active cells, variable.

EM - polytomographic analysis - 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 \approx degree of u. time at septation!

Mayia - units are $4000\text{\AA} / 200\text{\AA}$ - sites of r/o?
similar units of NP are intermediate causes of infection

multi-layers of sheath \rightarrow problems of A/uv DNA

Preparation of peripheral as the bridge of fibril to chromosome


Middle aspect:

Disc 11:30 AM Tues.

Aspect - [Maeshole - thinks too much systemic away.]


Pis - sequencing of Fe ligand trans total amount is small but ~ species.

Butter (Darsai + ...); nearly 1 base a / P. 2 Harris 3 trans protein restriction

Williams: - i as phase is  Fe



of Maeshole in relative sty. of NA, protein.

 empty or NA?

matter of fixation: if formalin non extraction?

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

applied!!!

automatis analogs belonged to Detroit commit that seems

unit of recombination: {existence < map vs env. physical distances

is a continuous series of units paralleled by experiments

evidence that there is a segment for cistron

is there a linear cistron?

trying to exhaust mutants in a segment.

n_I	n_{II}	n_{III}	phenotypes
$K-12$	$S = "B/B"$		
+	-	+	
+	+	+	
+	+	+	

The groups are unequal sizes.

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

A = 40%
B = 20%

Self

stable to insertion

can also be
cross-over suppressor
of asynapsis ✓

Exceptions: apparent deletions
are always linear & single segments?

mistake of exhaustive analysis —
had its origin possibly in Stahl's study

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

Concentrated as deletion class D. 11000 total

150 included & fitted in pairs

one locus most frequent; 2 phenotypes
(1 dominant)

mistake of secondary
locus alleles
complex loci only
confusing

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

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

no exceptions to linearity yet.

mutas + reton = or $\frac{1}{2}$ (multitalk not
excluded).

probably pairing
rather than
deletion w/ unit of
lac₁

S.B.

generate mutants from del x +? sum byproducts
c indicators (sum against + byproduct?)

H.H.

occurrence of -- as complements to ++.

heat shock ^{in part of} $<$ conversion, no effect as rec of
extension markers.

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

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

expects conversion in $-x$.
(combination of reciprocal crossover).

non-reciprocal crossing over?

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

got didymotera
Lindley

6
glossary
7/11/15

Warner - definition sus lat: prob. of specific
disrupting gametes or telomeres.

general homeostatic principle.

of mutation s.l. - any change S.S. - 119 sec, ^{stence} - ^{press.}
"hereditary modification" as broader.

cytoplasmic - extrachromosomal.

① meiotic 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 (Buggis, King, etc.). But: @ sexual @ cellular
in time @ regular predictable pattern @ integration ②
most data are ambiguous.

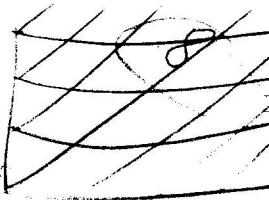
why not apply
study state concept to the
nucleus

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

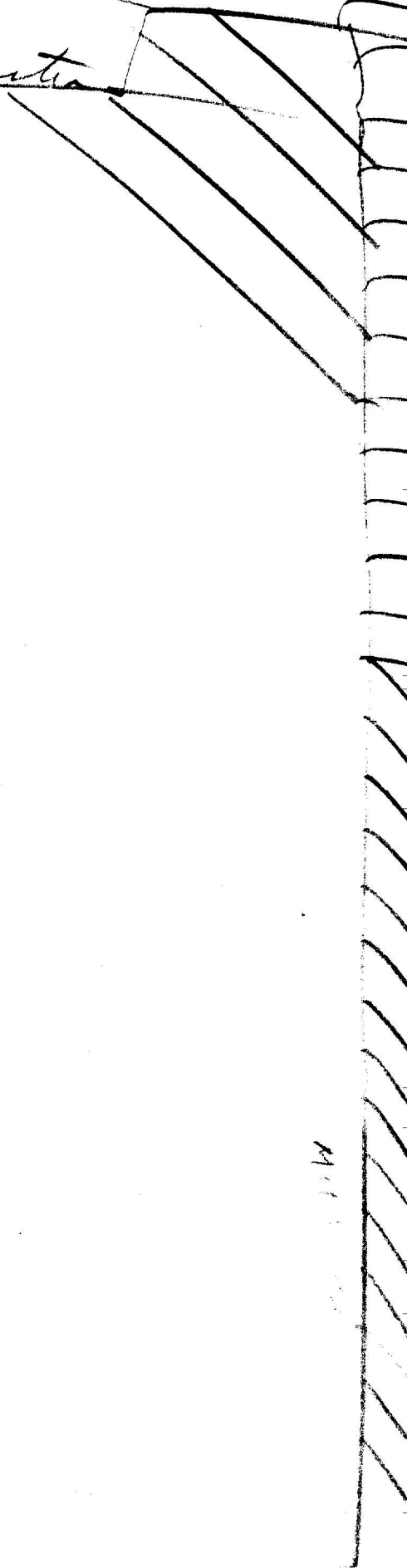
same familiar faces come back!
but correct the typo!
and a dictionary; what shows
glossary into paper

Send Buggis
Ford
up.

777



7 - Phonies - not guarantee - extra
why not go whole by - muscle take!



PHI 100

Mayia

spindles.

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

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

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

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

Mitochondrial mitosis - centers may influence orientation of chroms -
bonding system may be differing from them.

core extract + crystals of cytochrome gave 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-Berly: glycerol 4% + ATP:

elongation of spindles & shortening of chromosomal fibres.

minced nuclei - Alfrey
Hepner

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

25% centrifugation in whole cells. Study AA microtubules.

decarapherial, PPA had ^{little or} no effect on chromosome organization.

chromosomes had no effect on M - synthesis.

centers are abolished by colchicine!!!

5-b-Cl BzHib

anti plus D effects. inhib of synthesis effects minus.

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

Dan -
look for search
conjugates in
flagella?
implant
cytochrome in vivo
to measure center
cytochrome evidence
of -5H accumulation.

opt sucrose is .25M 10% deviation \rightarrow 30% inhibition!
secular variations from animal to animal (hormonal?)

F&T, heating, heating in blender destroy activity. Old nuclei do not
test overnight.

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

"not exchange" — not proven unless substantial
replacement had occurred! ~~not being supply~~ ^{induced by H₂O₂}

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

DNA removal with urea, then plateaus (very plateau)
cells are unaffected.

Radioautography is homogeneous — how closely carried
"fresh whole slide" — was pt/pt autography.

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

but 4x increase by "adding DNA"

Specificity of DNA?

all "denatured DNA" — just as effective

"acetic acid" had some activity.

deoxyribose phosphate

DNAse treated — core
deoxyribose

} both functions +

RNA also +

RNA mononucleotides etc. no. 9.

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

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

now? either DNA or RNA

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

azathymine?

in any synthesis

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

Chentremme: RNAs accelerates when catalase is induced

Vollmer - T2 cells - specific RNAs?

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

"I would have preferred to be injurious than congruous" (of Dale)
virulence no longer paying off.

cell parts expts:

Dale: - RNA fractured cells, restore by intact RNA. species -> p!

Sol's lab: mixed steps: activity 4-9. mixed as preliminary. + would like fossils to expect a water-soluble pupa to work. Thought to use protoplasts

Properties:

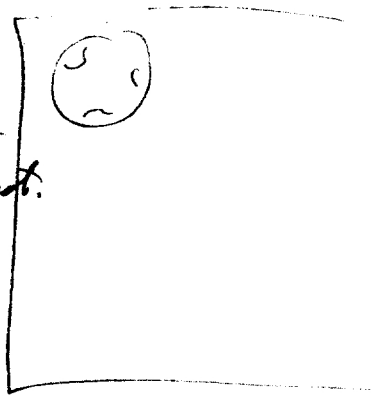
osmotic conditions

decided by 1, just; collected all DNA = parent RNA
Symbiotic prot.



what is membrane?
Symbiosis?

"fairly good" 1 per protoplast
quant recovery



Properties: 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 while making

(Do these in Ford paper?)

> 99% DNA removed. No effector enzyme with and same substrates.

∴ might make "strong statement".

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

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

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

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

Many pellets through pipette are disrupted.

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

resuspended loses 85% of DNA

RM = reading mixture. substrate substrates in a phosphate HDP lose DNA.

what are "substrates" really??

genes ^{read} RNA cannot factor fracturing 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 leg + linear synthesis.

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

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

~~RNA~~

nucleoside diphosphates.

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.

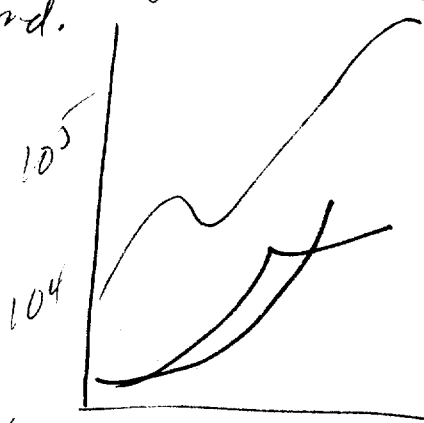
Harnett

Radiation effects on TP. Target theory may apply ($S^p \rightarrow S^s$).

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

Relationship of caught time to division cycle.

Thomas found k_{eff} depends on number of caught bacteria. Limiting reaction evidently bimolecular. Had to study kinetics at high DNA, plateaus are found.



log log plates

What are reasons for nonlinearity?
 heterogeneity of DNA? Extra DNA gene 2 gives early plateau, decreased by progress of unmarked particles. 1st plateau is of total DNA. Break due to inhibition. At high number, inhibition is partly wound, may be max. of effective synthesis.

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

[polymerase?]

Radiated DNA -

low bio assay at 3-4% survival.

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

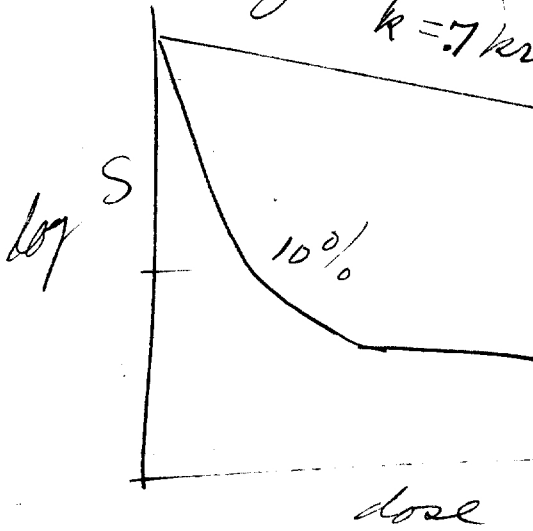
inactivation effects problem - mediated in frozen state, where protective substances have no effect, & serum volumes are smallest. i.e. sensitive to inactivation effects.

Kinetics of inactivation

($\approx 4 \times 10^5$)

$k = 7 \text{ hr}^{-1}$

direct eff.



Maumey + Fyfe
($2 \times 6 \times 10^6$)

only 1/10 app. chem. m.w.

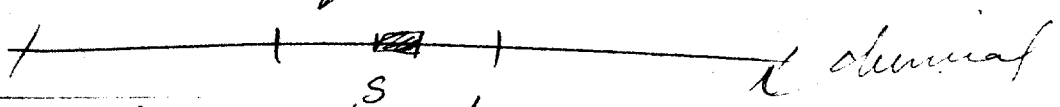
(radit) radar < center. ? Residual activity possibilities.

seeing DNA not heterogeneous as claimed.

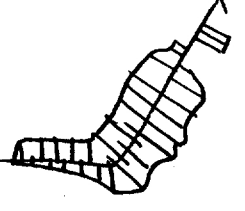
Distribution particles? Fractionate DNA $\left\{ \begin{array}{l} \text{high gravim high TP} \\ \text{low gravim low act.} \end{array} \right.$
 mean no effect no difference.

more sensitive units are present in same populations. why are small aggregates > sensitive?

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



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



Hotchkiss: assume \rightarrow hypothesis.

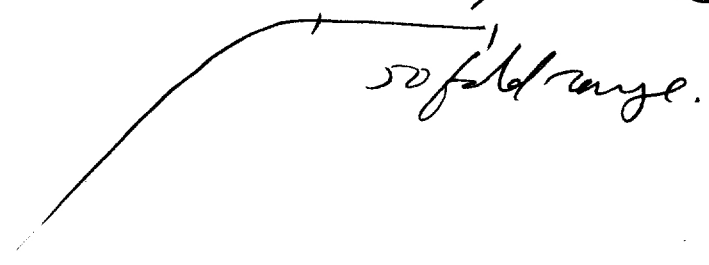
Quantitative factors

Capsule system - originally by deletion. Considerations: sediment too soon or too late n.g. momentum, entry must be specified.

If sediment is resuspended, may let reveal add. \rightarrow s. pres. matter of selective conditions. etc. So moved to dry environment.

Rollin doesn't see plateaus

level of plateau depends on proportion of S^R/S^S DNA.



Berndt - chromatography of DNA
anion exch.

put #1's on wood
DNA/cell
DNA/wall \rightarrow max.
"molecules" / cell.

Ectocellulose -
variety of fractions tested

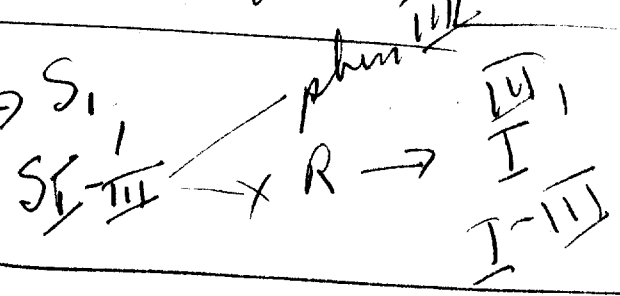
least eluable DNA is most active

2M NaCl MNH₃. \therefore heterogeneous

In cap. exps, can get intra-plateau?

sp. signature?

Australian: $S_I \rightarrow S_{II} \rightarrow S_{III}$



H. Taylor - Quant. assay.

ΔN_A is not exhausted!

What is being measured in essays?

\therefore not a stoichiometry + assumption of a time slice of a bimolecular reaction needs 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. "purplish" 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 taked to be altered.

③ Nature of heterogeneity?

④ Replication, so far.

⑤ an chain system.

-x is synthetic
medium?

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

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} v DNAse destroys " 100v/ml DNA in 20 minutes
during, dialysis also " cut & suppresses by prechopping.

assay methods
for these procedures

denaturation 5.2%, NH_2^-

didysis: loss of heat stability + denaturation.

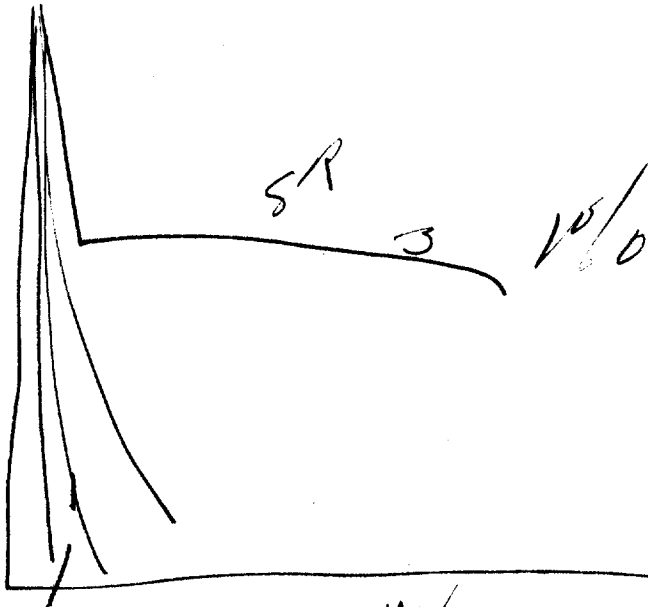
M. luteus:

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.

Mutant both + DNA competent but all ~~new~~ genes is a protein, making

pr. Her. coli? Neisseria

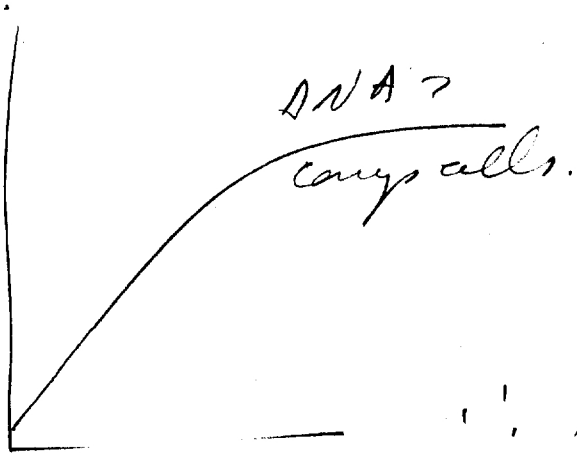
importance of

- 1 capsule e UV diffuses →
 - 3 = SR
 - 2 capsuled
 - 4. SR front type b.
- pr 2 similar

A to pr W. — Cell incorporated after cessation of growth & no more in DNA/cell. ∴ exchange without resynthesis of DNA

Goodgal: 4 DNA 300 mols/mutase ~ 1 cell's worth

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



Sd cells do not x
do not x DNA.

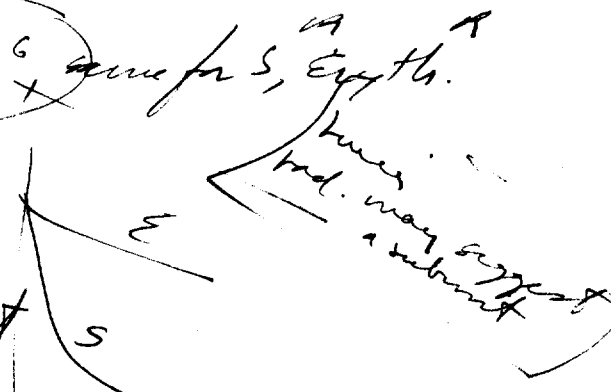
1. gene per cell.

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

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

No separation obtained at up to 4V

Sordgal coli caryome + ATP + Mg + light
restored UV'd TP from 3-30%



says Hemophilus, pneumo. to not photo-reactivate per se.

lung - ? as i phage. - no.

Benditt's Venulas early fractions have $M \approx 400,000$ even
ostenuledotals (pairing minimum)

Lidgy intrus periferi (a la Schaeffer)

	10^{-6}	10^{-7}	10^{-3}
influenza	x	x	xx
paramyxo	x	x	x
suic	x	x	x

SR → x
home ratio is
inter constant for
given p.p. trial
of strain of donor.

G. Salmonella
+ coli %
Lidgy's

"crossing over" —

13 seen in heterozygotes

4 steps in heterozygotes

v. occ. recurrence. isalleles for reversion.

also differs in recomb frequencies!

some mutants give confusing overlaps.
heterozygote.

Kalchauer

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
esp. 2-point tests

Non-aggression pact

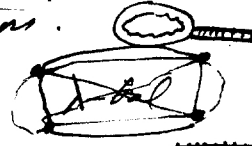
Herriott - general review of label marks in re DNA effects

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 lysogeny are
-x in Salmonella!
generalized.

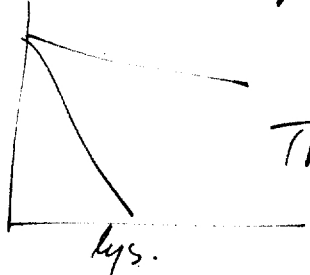
- Zinder E. coli.

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.
exog. \ll prophage.



This is Salmonella.

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

incorporation step - possibilities: "unilateral" $F/a^+ \rightarrow x F/a^-$

Jacob: eclipse to lysof. cyclic behavior
temperature quality.

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

expression: \rightarrow phage + immunity.

induction w/ \rightarrow phage.
etc.

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, marks in output
have input ratio. Induced bacteria superinfected behave as if 3
per bacterium. ≈ 3 units.

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

E. coli K-12 L+T. λ ML- λ^+ . λ cos λ^+ x λ^S .
Astrupts by L + E.W.

lamias "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, degen varies 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 of 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.

W+

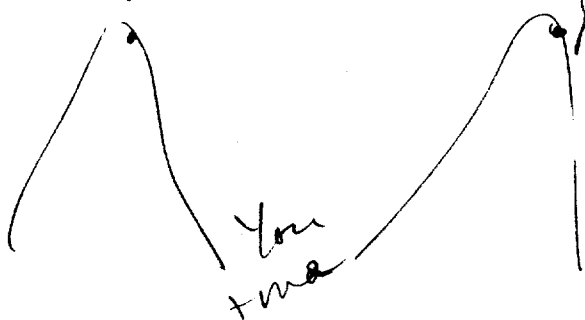


Schumann

Davidson

Davidson

f. H. mod.



host modification in
 chemical terms.
 any pure clones
 in TMV

Vollmar et al
 any effort to
 stabilize RNA
 with normal
 plant proteins
 essential viruses.

is the prophage bacterium
 hairy - an ensemble of
 prophages?



Buy
Ch+
Law.

Are there conservative
exchange reactions?

Doty - plan of mountains.

(also of section
is to show the
clouds.

Chargaff: language difficulties - dictionary lacking
 μ coil \neq mol helix

HC, AMC
N-Ho-Admme.

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

~~plant viruses~~ may be typed. mammal RNA close association /
 proteins.

This is the
fundamental
question
+ causes synthesis

"NA can carry information" $\Rightarrow \Rightarrow \Rightarrow$ 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 concluded that NA is huge cycle

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

Tb sea water

(6 amino = 6 keto)

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 terminator is.

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

~~Is~~ Is 5MC/C random. ~~but~~ occurs fractionally.

clearly not natural genetic content of T2?

PNA - no regularities, but no pure pyrimidines at first.

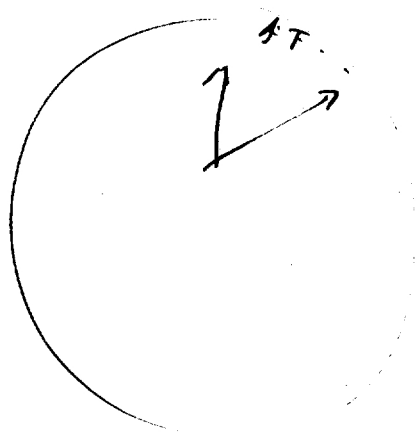
clearly nucleoproteins: the relation $A+C = G+U$ was preserved, i.e., but no others.

Smith - no good evidence on branching of RNA.

seq. some problems.

Whitfield - remove phosphate oxidize sugar

remove phosphorus... again need pure material.



Thayab
SBC

Choo
re print

Smith

aggly crystals in B. cereus cells.
not homogeneous. cyclic end turnover faster.

Crick 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 preferred not required. suggests repl. principle.

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

He suggests that XT replaces TT pairs but gives hemidoubles

(flexibility too much)

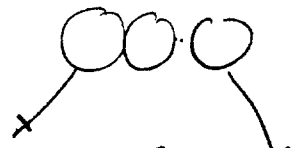
Chooff/complex

Touloume & Zam.

Choo - whole cell RNA ~ enzyme RNA.

Bonditch - recorded deviations from 1:1

Crick: 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.

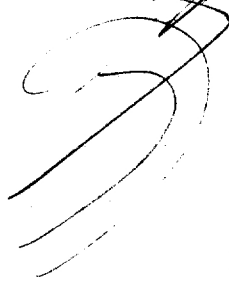
Not yet critical professional judgment on ϵ ray concordance
but no measure is yet.



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

nucleoprotein. prot fills in the "man" groove.

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



as 3d chain, base groups match by P.
non polar spacing top of 2 or more

\therefore non polar AA should occur in pairs.
much analytic detail fits that.

nucleohistone

JA

synth poly RNA, 4AUC. or AU

algae = bad photos

poly A - \rightarrow clear photo

card RNA \sim DNA specific base pairing
 \rightarrow except A:A

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

Large issues are

conc. org. organisms based on
feasibility of manipulative methods;
but things like eggs.

① Availability of test systems for DNA, RNA.

② Concentration of analytic effort on bio assay systems.

specificity absolutely essential.

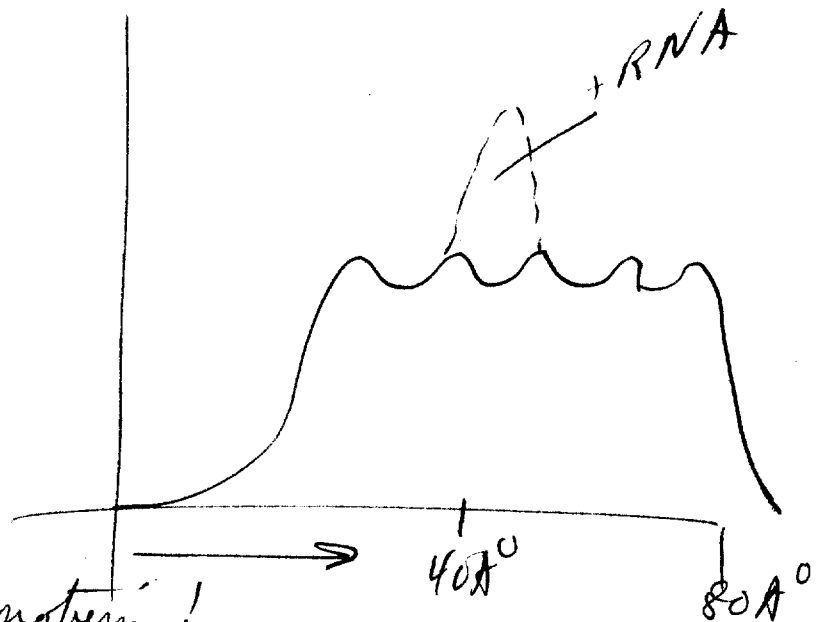
Phase
Transformation (was I like to
call them
DNA-mechanism)
x

Reels - 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 changes
of TMU
and TMU.



∴ not core but RNA is
wound into peripheral proteins!

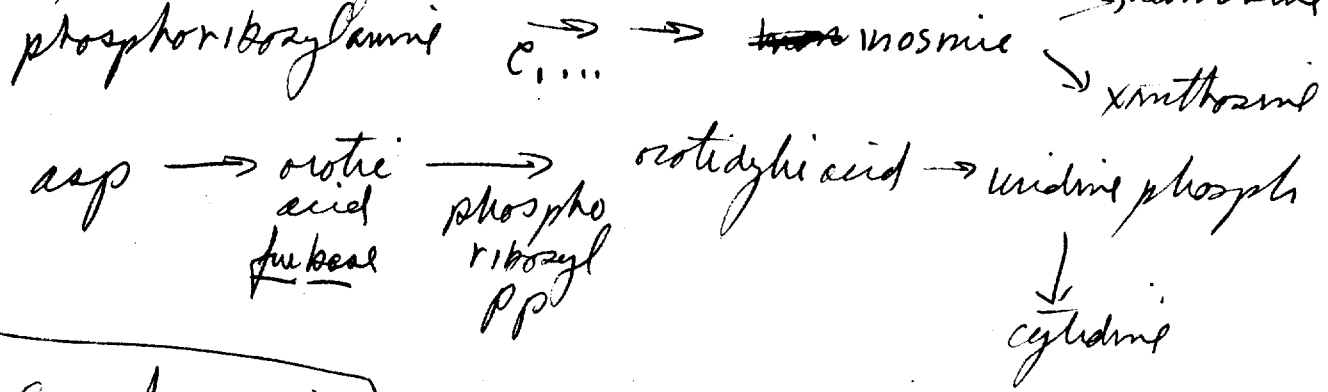
probably single strand if fully ordered, not em. RNA.

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

Art Kornberg: purine synthesis

- organically, brain, concentrated, urea

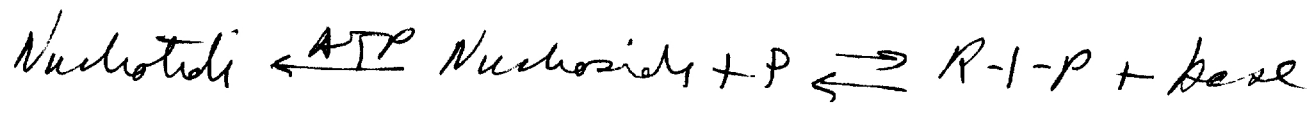
Enzymatic synthesis of nucleotides:



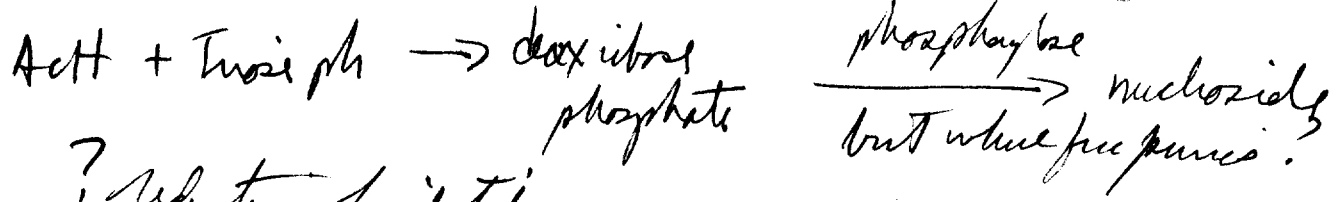
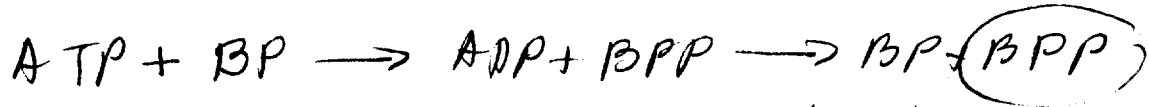
U.P. no free purines or pyrimidines

$C_1 \rightarrow$ uridine typical
 but various bases though uracil \rightarrow uridine phospho.

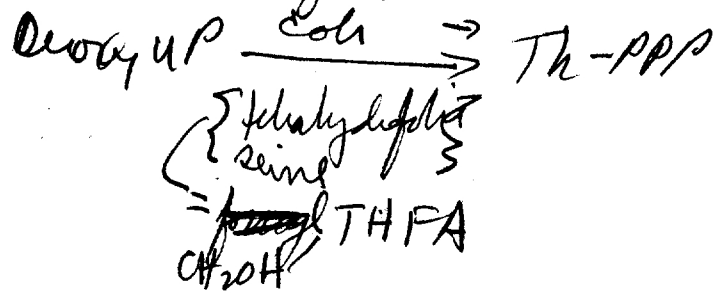
"salvage pathways" $HX, G, A, U \xrightleftharpoons{PRPP} \text{ nucleotide}$



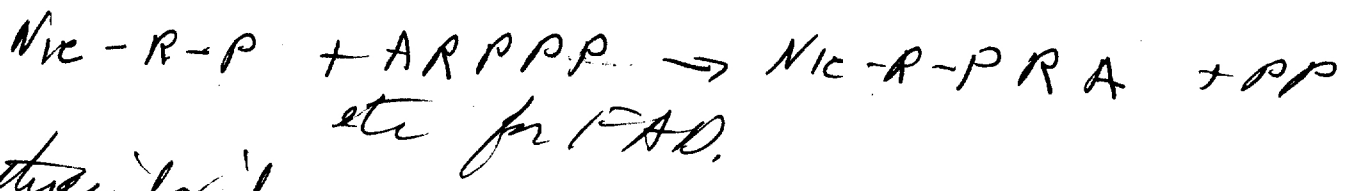
$\text{Nucleotide} \xrightarrow{-PP} \text{Nucleoside} \xrightarrow{-P} \text{Nucleotide}$ $\left\{ \begin{array}{l} \text{used for further synthesis} \\ \text{Krebs reaction} \end{array} \right.$



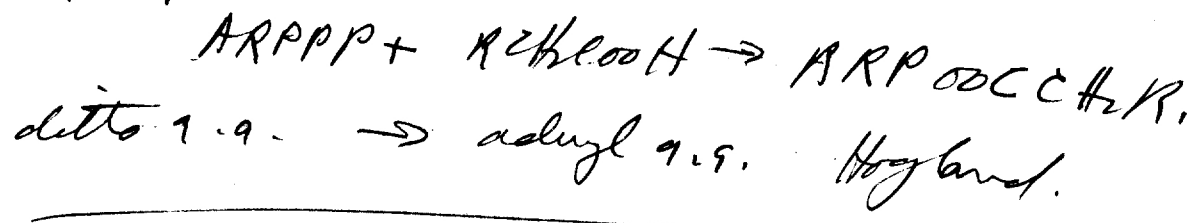
? reduction of ribotides.



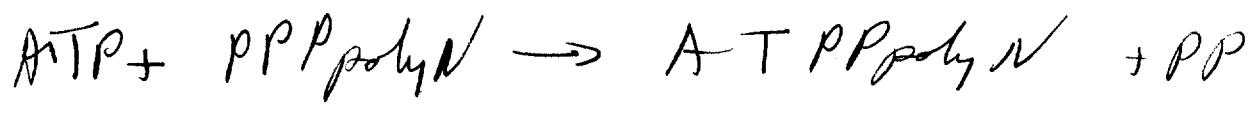
of coenzyme synthesis
methylol-enzyme reaction.



phthalic acid.

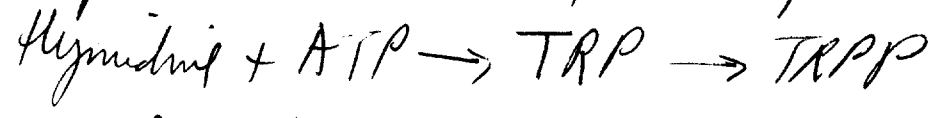
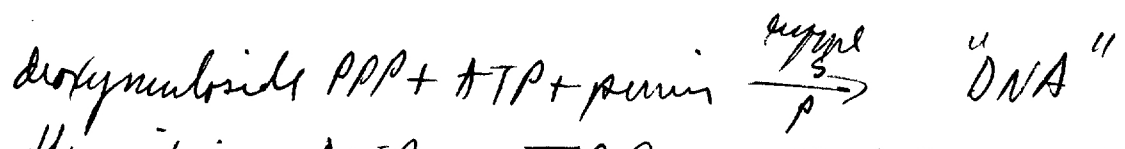


? polyphosphates



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

DNA:



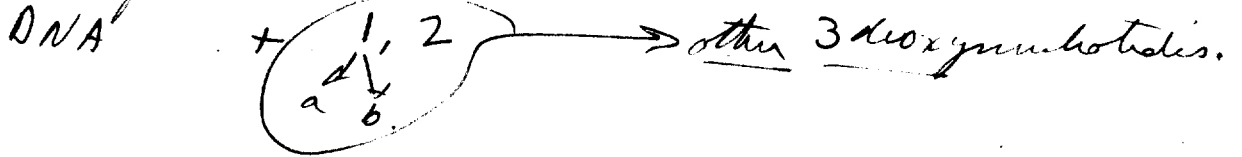
TDP is sufficient as TTP.

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

All 4 together do react

purin

DNA - purine heated, rich in
cyclic redifraction ↑, rich in DNA Recently purified. But see
come to nasidial fraction + dead.



now need all pur components

ATP not used. Suggests probably phosphorylase
not needed.

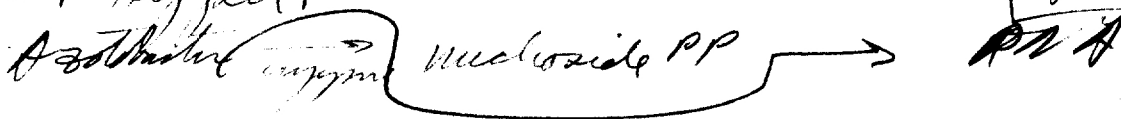
Now try net synthesis: 30-50% mass.

The RPPP is purine.

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

could bio activity increase
if pur DNA were used?

Lehwa + Heppel.



"polynucleotidyl phosphorylase"

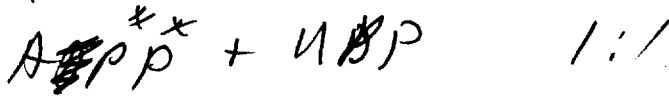
has made ADP → poly A, G, U, C etc. also

AGU, A G U C

Potter had shown the other phosphates in all.

? structure + nature of polynucleotide - structure w. is RNA (Pids)

poly AU.



11mg AU_n about 1:1

products of RNase. phosphodiester attacks pyr uncl. yields pyr 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^x trans from A to U at splitting. fairly interspersed A, U, in polymer.

? symm is "random".

Meaning of random.
random bias polymerase work.

? Intracellular route. Means of orientation? K_s is known very low. Km 10⁻²; Hill slope < interlink. Max # total sites of NPP formation. The enzyme is unspecific - e.g. polyHX. also unspecific in phosphodiester: also bacterial RNase. rates vary. # dihydrochloride O₂ protective as 5' 2' OH. also inactive on above 5-pp

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

try virus RNA

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

Needs NMPs start of chain? *no*

But AMP + UDP \rightarrow no A in end groups.

? one or several inhibitors.

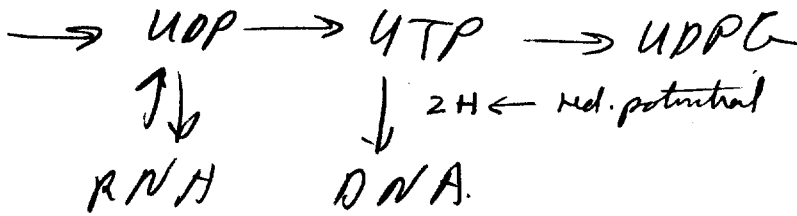
Is RNA specific
exc. in plant viruses
change of standard spec.

Potter: ATP \rightarrow RNA ADP \rightarrow RNA.

growth control. Thus ATP inhibits
RNA formation.

extra biosynthesis
of DNA.
reversibility of
Hornby ex
solubility.

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



Hornby extracts:

	<u>RNA</u> <u>prop</u>	<u>DNA</u>
Reversibility	+ inhibits	(-) but not inhib.
K_m	10^{-2}	10^{-5}
diversity of N	1	4
pH	- ?	+
extent	∞	$\neq 1/2$
specificity	"	ATP not usable.

Cohen - regulatable systems.

point strongly made.

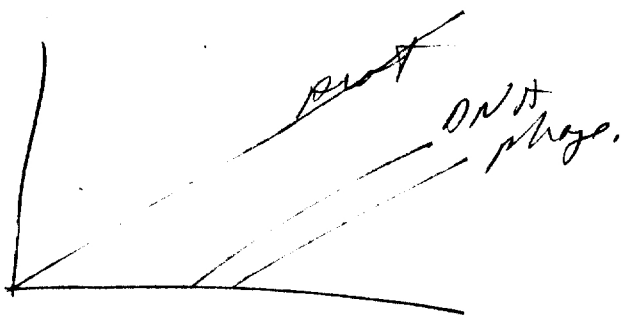
RNA polymerase why as
enzyme state?

reduces almost to E. coli and phage

Teven: HMC and G-HMC

1. may be unique to exphage
2. inf. cells stop new enzyme synth.
- 3.

do hosts?



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

early S35 does not merge in T4 strain.

? repair mechanism

SMT inhibits DNA synth. only if added at beginning.

if added later, → DNA s/ further protein synth.

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

? role of protein: inform or enzymatic support.

some small RNA is also turned over

to B growing, Hypermurine, Hypermurine \rightarrow DNA.

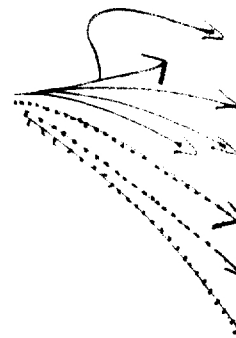
pres. Hypermuric acid.

defect B. Hypermurine \rightarrow DNA.

(consistent of Hamby's theme)

ribosyl nucleotides \rightarrow deoxyribosyl

Hypermurine analogues do not inhibit B.
do inhibit virus synthesis



Do Hypermurine in wild L15?

Amino acid \rightarrow methyl group missing.

Be Hamby insects itself

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

\therefore metabolic control of genetic material

Zamiatol Bell effect - "exchange in DNA"

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

Control of nucleotide pool a key to mutagenesis.

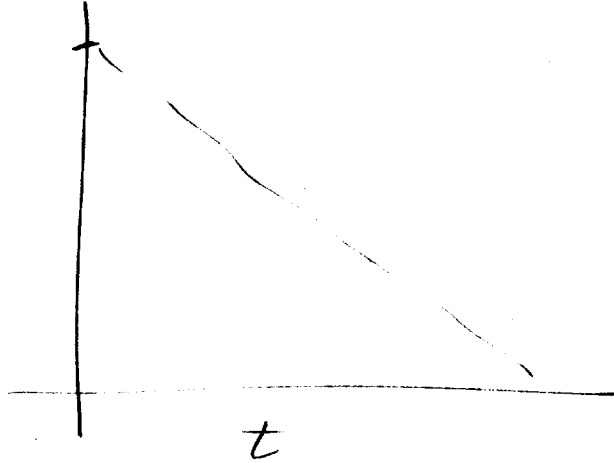
turnover is manifestly rather than whole RNA substitution is manifestly Volfain shape of pulse curve



Volfain specⁱⁿ granule fractions. In pulse exp showed RNA, not DNA turn

ad student
Seymour :

What's the suicide curve of T2 grows 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 effects dispersive depending on mechanism.

grow hot phase, $1 \rightarrow 200$
 $par_1 \rightarrow prog_1$
 $\rightarrow prog_2$

how does protamine dissociate?
c-o can still be a copy mechanism.

does this mean second step is still semi dispersive?
has 4 understand 1d expt.

~~W + M~~ W + M like 80% conservation.
mountain $prog_1 \rightarrow prog_2$ still 80%
Potter $\rightarrow prog_3$

Hershey thinks just inefficient.

cy: individual autoradiography

BS: P_{32} inactivation method $prog_1$ - inactivation assay, as ability to ~~maintain ability~~ to transfer P_{32} to progeny.

cy: big pieces $1/2$ is distributed to 10-20% into
 $1/2$ small pieces.

i.e., same dispersion of parental atoms.

of cy: Both methods as $prog_2$ \leftarrow another same fraction of big, small pieces

fractional number - hot cells at 1 division
assumes nucleus is unit all cells div.

15 nucleus one
unit
all cells 1 nucleus?

Thozzi

B.?
[Mapis]

degenerate material for replication:

eg: can measure 15 cpw worth.

star size does not change from $p_{12} \rightarrow p_{22}$
semiconservative replication: $2 \rightarrow 3$.

Is bij piecewise?

stars are in h^+ fraction & 90%.

(total activity?) \rightarrow

problem of bi helix \rightarrow may be 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 would like the new edition of the Soviet Encyclopedia on Stalin. ^{so many farm truss.}

"classical crossing over" - mythification of linkage model - possibility of epigenetic still relevant.

cf. Detroit Symp. - science as series of conferences punctuated by short recesses in the lab.

"not necessary to assume clo is ever intragenic.

Ris - emphasized binding from chromosome fibril to chromosomes.

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

tried rit, mit and pfrit, salt melt and pfelt.

Serious semantic problem & necessity of purifying germanic since rit 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 - ris is foldschmidt too close too?

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

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

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

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

@ host modification of fin

Chargaff - nucleic vs gene.

Call Dr. Gelman

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

Wednesday.

HAZIA

flagella is promising contractile units

is there cytochemical evidence of SH accumulation? dependent cysteine?

Alfrey

"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 mentroni?

Sol

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

Hammett - Pollin - Zamenhof

- relationship of competence waves to fission cycle

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

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

role of synapsis if, after all we're multistep 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	→	B	1	1 < 3 < 4
B _A	→	A	2	
B _A	→	B	3	
B	→	B	4	

3 > 1 may admit a more compatible symmetrical recombination best explanation.

Segregation only a transient condition.

Zamenhof may be implying a dis-integrated system of DNA molecules, - this is what Hammett had in mind getting genetic recombinations.

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

few papers had anything to do with the...
how to say what everyone does - elliptic approach.

... may not work...
here.

Most genetic discussions were matters of definition; Q. 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 tones have the obligations only of being true demerits; in our lab we had tried *trit*, *mit* and *pf1*. *muton* and *cistrons* 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 *Genetics* 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. — Bel. reliability of the *reconstruction* ^{Hartman}

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

On Dr. Rhoads' talk, ~~at~~ Dr. Spiegelman asked whether this was a semantic ~~issue~~ problem; of course Rhoads was asking the material question, are any nucleotide substitutions of nucleotides, which are the closest possible approximations to point mutations. — Sign of *Methionine* for pos. effect

Benzer to a point of Craygoff's query, which is really an old question on the relationship of mutons to genes. Undoubtedly a muton is a single nucleotide substitution, but it would be a meaningless case if deoxy acid were a transforming agent. The meaning of the code lies not in the letters or syllables, but the words and sentences. The specificity of transforming agents can best be visualized in terms of synthetic attributes; it would be worth knowing if deoxy nucleotides were mutagenic in the pneumococcus.

(return) →

Meeting reminiscent of the Detroit symposium - where Benzer would have been even more typical - seeing so many friends you feel that the real life these days is a linear sequence of conferences punctuated by short pf1 in the laboratory. Informational exchange = recons

End
somewhat platitudinous

Gales natural there and Dr. Spiegelman's line had emphasized the ease of this ingenious frontal attack (have to watch out for such words. at Detroit indefensible became indefensible.) 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 recon than chemical recon DNA. Genes this may not same recons 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 cytoplasmic should not be too hard to duplicate
in ~~in vivo~~ The RNA systems have turned out to be extremely hard~~

The transformists have upped energy in 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~~

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 genetic
material + physiological ~~of complexity~~. 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 ~~nucleic~~ demucronated
nuclei. Any scheme of DNA resynthesis here?