

7/5/45.

[commonplace notes. See refs to sex, crossing. 2/48]

X a. In medical Microbes, now in Fungi imperfecti, test for sex by crossing different "species". What is a sp. in an organism that cannot be hybridized? Consider range of genetic variation in *Dm*, *Nc* or *Hs*!!

[*Drosophila*, *Neurospora*, *Homo*]

✓ b. For variability in *V. crassa* adaptants, plate out microconidia in "Fluffy" strains.

[Lysine being used] ✓ c. For limiting growth ~~in~~ to produce "resting" *V. c.*, use a deficiency mutant, e.g. pantothenic acid in a medium containing limiting amounts.

d. For stats. adaptation, adapt leucineless by keeping varying lengths of light imp. then being to low where adaptation to *V. crassa*.

Pab¹ inhibitor
??

e. Feature of solid media accelerating ^{in pab¹} adaptation.

X f. Quick means of sexing *Neurospora*?

nutritional acid.

g. Yellow pigment produced by 10575 (*tryptophanless*) when adapted.

Kohn et al. - Induction of a methionine requirement.
Is this selection of a mutant? Does selection require
both sulfanamide + methionine? A mixed culture of
methionineless and parent *E. coli* of Kohn's strain
should be purifiable by such selection. Speeded by
single cell isolation. Are both factors necessary
both for mutation + selection. Alternate transfer
in normal + methionine - sulfanamide mediums.

In *N. tetrasperma*, a deficiency mutant can be in-
troduced by crossing to *crassa* + backcrossing
successively. The regular bicarya are produced.
Purification by selection would be indicated by a
failure to produce perithecia spontaneously.

Sex det'd by 2 genes? in basidiomycetes?

Relationships between color and pig-ment?

Methionides in *Neurospora*.

Valine - Isoleucine, separate adaptation?

adaptation
not clear yet

✓
healthier
If adaptation can be prevented, lipase in bacteria can be selected for by using two different mutant strains of *E. coli* and growing in continuously renewed minimal medium. In general, a sexual process could be demonstrated by plating out mixed cultures + finding a wild so all strains adapt? Transformation?

Does gene determine type specificity in pneumococcus alleles? Meaning of alleles in bacteria.

✓
labat
Hirsch
impractical
Production of serogenic mutations - use *E. coli* strains, absorbing lysed cells over deficiency anti-sera, and immunizing against purified antigen. These antisera should then induce mutation.

Die
ec. lys
apt.
nutritional
selection.
Does adaptation occur & selection? To a 1633-5531 mutant make part limiting. When growth is complete to an intermediate level, wash mycelium by removal of Ca^{2+} with pipette + replace Ca^{2+} with Mg^{2+} . After various periods, add Ca^{2+} part steadily. If adaptation has taken place, it should grow immediately. i.e., if a plate level that does not cause de-adaptation can be found, washing is unnecessary.

one can also transfer + wash 1633 mycelium in buffer
+ sucrose + debrun adapt. by adding nutrient.

In bacteria, a selection hypothesis for dissociation would
always be tested by recovery of the dissociant from
a mixed culture under the conditions of dissociation, in a
serial shorter time than normal dissociation. In *Neuro-
spora* one can make the adapted component, combine
it another marked, and attempt recovery.

Emerson 45 describes an adaptation which looks
much like mine - heterozygote. It is genetic, he
seems to regard prot production as a factor! Also selection
Why is this heterozygote not selected for??

Assay for prot: anti ~~to~~ sulfathiazole activity
in presence of excess methionine, etc. (v. Sull)

Ask Beadle
in his recurrence
of 1633

Alleles in *N. crassa*: irradiate 1633 to non-adaptability.
Possible alleles: non-adapted; partially adapted;
stable + unstable. (Compare a , a^s , a^{bs} , a^p ... A
series in Mary's interacting c. DT.) Here,
quantitative. Hue a 1633^p would gradually increase

Compare mutability of 8829, 33757. at 25, 30. The former is probably higher.

PAB-synthesis is a limiting reaction in 1633⁺ strains.
Temperature studies? Compare 1633⁺ = 1633 + enough
pab to give same growth rate at 25°. ~~Penetration - temperature?~~
~~Proteolysis + temperature~~
~~compare 1117~~
Growth tubes + liquid: 30° - 25° & !!

Review types of pabless adaptations!

As bacterial recessions, estimate micropopulation size
by sectioning + staining agar.

Examine Bradles genetic data on 16117. Note:

21 asci = 168 spores from 16117 x 15300a.

all asci were 1:1 but a total of 517 spores, including
these and random isolates were 273 (-): 244 (+).

Comparing random isolate data = expectation of 1:1 $\chi^2 = 2$.
that they are the same; comparing The 42 chromated re-
combinations = an expectation of 5% computed from
random isolate data, $\chi^2 = 2$. They certainly therefore
have not conclusively shown that 16117 is a single gene.

The best way to refute them is to obtain monogenic stocks by mutation!

✓ Compare 1633⁺ and 1633[°] E⁺ as limiting sulfide mutants.

in rate as plasmagins accumulated!!!

Growth factor requirements of tissue cells, cancer?

Maintenance of aneuploid cells in man. Synplasm + Hetero-
caryosis. (Bacterial associations?)

"Adaptation" to biotin-requirements in Neurospora. Mass iso-
cultivation. (Caryover?)

Transformation in heterocarya. $a+b \times a-b$. find
++. Either mutation or transformation or sexual process.
(Presence of $a-b$ in $a+b$? demonstrate easily. would
certainly be selected against.) $a+b$ killed b .

Cytoplasmic "inheritance": $a+b$, monoconidial
isolation - if still grows, = how genetically (a) Would prove
plasmagone hypothesis [a lethal??].

Multiple mutants in bacteria by Sulfamethoxazole
selection and plating out of various heterokaryotic
strains simultaneously, SA-resistance is obtained

v. plasmids

Amie by
Tatum v. 7.

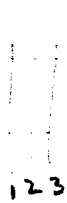
Beale
misses
point.

↳ Lys-arg.

See Tatum

Fabrics (braids) + permutation groups.

e.g. $n=3$. $(32) (12) = (123)^n$

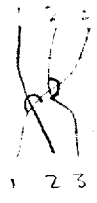


123



123

12



1 2 3

32



12

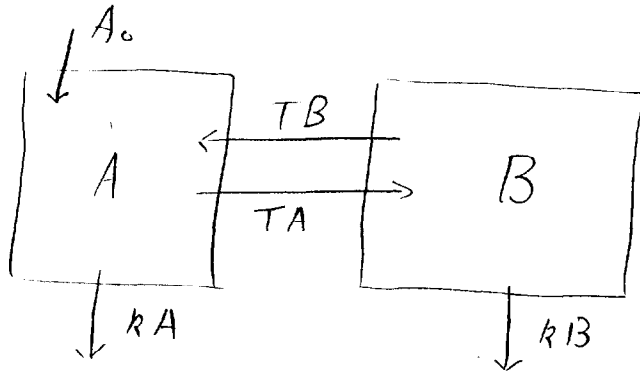


32 etc.

Mathematical simulation of gene distribution in n is required. Mathematical analogy for n genes is of ~~$f_1(a) \times f_2(a)$~~ $f_1(a) \times f_2(a)$. (Vector analysis?)

9/15/45.

Plasma decay disappearance curves in parabolic.



\$T\$ = transfer coefficient
 \$k\$ = disappearance

$$u = -(k+T)$$

$$\dot{A} = -(k+T)A + TB = uA + TB$$

$$\dot{B} = uB + TA$$

$$\ddot{B} - 2u\dot{B} + (u^2 - T^2)B = 0$$

~~$B_{max} \cdot uB = TA$~~

~~$\frac{B_{max}}{A} = \frac{T}{k+T}$~~ $\frac{B(+)}{A(+)} = \frac{T}{k+T}$

$$A = \frac{A_0 e^{-kt}}{2} (3 - e^{-2Tt})$$

$$B = \frac{3A_0 e^{-kt}}{2} (1 - e^{-2Tt})$$

$$A/B = \frac{3 - e^{-2Tt}}{3 - 3e^{-2Tt}}$$

$$B/A = 3 \cdot \frac{1 - e^{-2Tt}}{3 - e^{-2Tt}}$$

Using the Ryan-Ledeburg method of determining mutation rate one can do studies on influence of such factors as temperature, hot-cold shocks, radiation, antibodies, etc. [incl. genetic background] on a large scale. One could also, if controlling genes were discovered, examine their effect in heterocarya. This is a powerful tool! [for studying gene structure. E.G. influence of adding extracts of *Uromyces* spora [transformation experiment]

One can also calculate the composition of heterocaryotic mycelium using color marked *le* stocks, labeled = different test genes.

E.g. ~~prot~~ *prot* *prot-le* - + and *lys-le* 15300 The proportion of + adapted to 15300 adapted cultures is a measure of their proportions in the mycelium.

Expt.

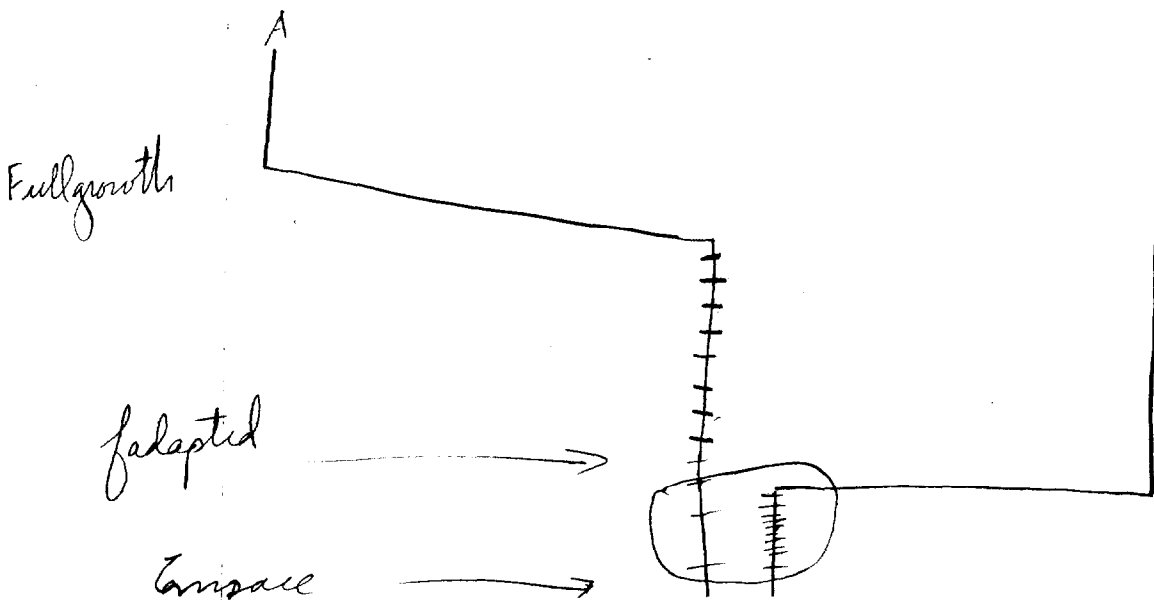
1. ~~Use~~ ~~from~~ ~~Fruc~~ $\bar{c} = .25$, 50mg leucine. Stationary at 3-4 days. Watch for adaptations on stationary mycelium by noting increased growth [show that leucine was limiting.]

2. Lys-leuc stocks. Use limiting ~~leucine~~ lysine, excess leucine. After 3-4 days, take R_2 and wash mycelium, transfer to excess lysine, limiting leucine, and watch for adaptations. When the half-life is reached [should be calculable from (1) data], transfer the other $\frac{1}{2}$ of series in same way [show that lysine was limiting]

Lysine limiting
Excess leucine.

Leucine -
lysine +

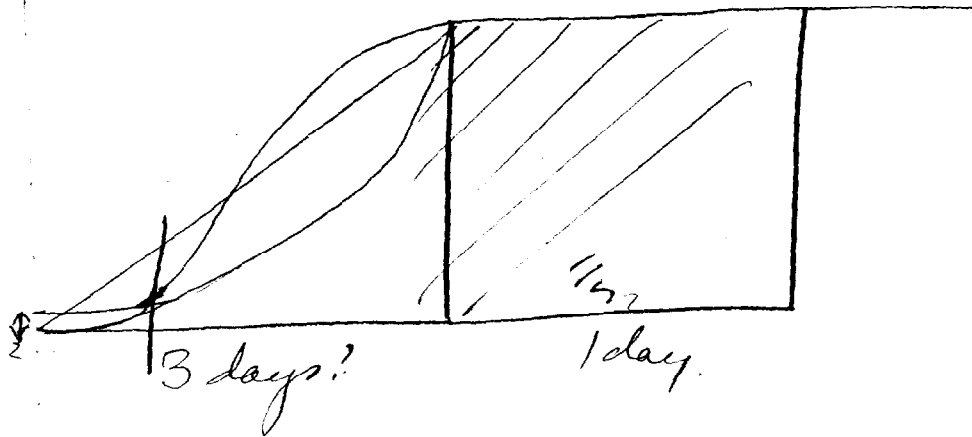
Lys +
Leuc -



This will determine whether, in a stationary culture, leucineless mutations occur whether leucine is limiting, or leucine is in excess and lysine is limiting. Therefore the mutations that occur in a stationary culture in which leucine is limiting do not occur in response to the absence of leucine, but due to an unrelated cause, with a uniform probability per unit mass of mycelium.

In order to calculate the adaptations expected during growth, plot area under curve. Use an empirical approximation.





$$A_m = A_0 e^{kt}$$

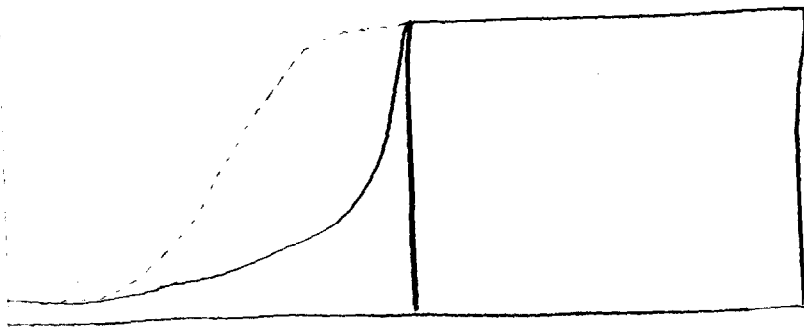
$$\int_0^m A_0 e^{kt} = A_0 \left[\frac{e^{kt}}{k} \right]_0^m$$

$$= \frac{A_0}{k} (e^{km} - 1)$$

negligible

$$rca = k A_0$$

$$rca = ca k A_m$$



See: Beutner & Snyder Pr. 19: 370-380 (1933)
Boell & Taylor JCCP 3: 355-383 (1933)
Kernd Biol Bull 60: 245-268 (1931)
Mensch Pr. 11: 447-474 (1930)
Mensch PSEBM: 29: 666 (1932)
Lund & Moorman JET 60: 249-267
Tang, P.S. QRB 8: 260-274 (1933)