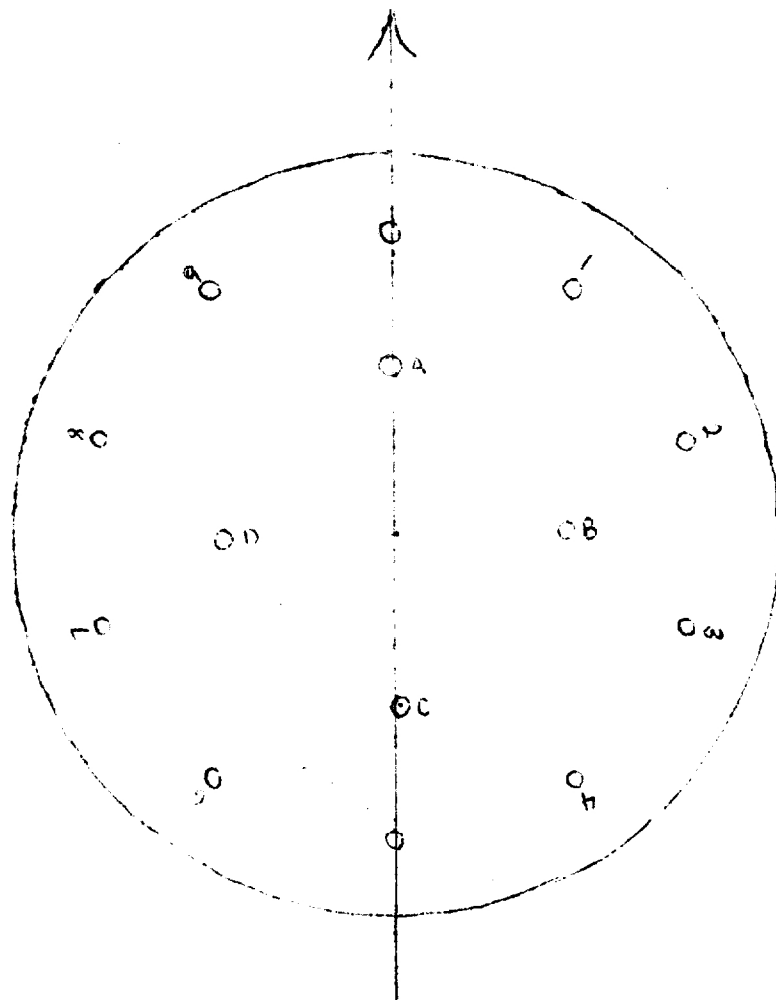


YALE UNIVERSITY
OSBORN BOTANICAL LABORATORY
NEW HAVEN, CONNECTICUT

EXPERIMENTS IN THE GENETICS OF BACTERIA

1946- 1947.

Joshua Lederberg.



AUXANOGRAM STENCIL -

Joshua Lederberg
 Osborn Botanical Labs.,
 Yale University
 New Haven, Conn.

Abbreviations:

F(x) = Fries supplemented \bar{x} . (Neurospora minimal).
 CM = complete medium (essentially aminoac + vits.)

Ca = HC = Hydrolyzed Casein.
 Vit = Vitamins B + hydrolyzed YNA
 YNA = yeast nucleic acid
 EAA = EA "essential" amino acids =
 NEAA = NA non-EAA.

⊙ = minimal
 ⊕ = complete
 T(x) = coll ⊙ = Tatum + Gray's + med. suppl.

Amino acids.

- | | | |
|---------------------|---------------|---------------------|
| 1 leucine | 9. methionine | 17 alanine |
| 2 isoleucine | 10 histidine | 18 cysteine |
| 3 valine | 11 threonine | 6 tyrosine |
| 4 phen. | 12 aspart | |
| 5 trypt. | 13 glutam | Ind = Indole |
| 6 lysine | 14 proline | Orth = orthocaulin. |
| 7 lysine | 15 HO- | |
| 8 arginine | 16 glycine | |

- Vits. V
- | | |
|--------------|------------|
| 1 Thiamin | 8 choline |
| 2 Riboflavin | 9 inositol |
| 3 PAB | 10 Biotin |
| 4 Nic | |
| 5 folicae | |
| 6 pyridoxin | |
| 7 pant | |

Protein + Pyr.

3/23/46. -3/24

Y 160.

Test 156 - Strains. Dil. surface growth in H₂O + test as indic.

broz 9P23. 33°
1st Reading: 12h.

2nd 24h.

#	0		C-M		HC		VLT		9A24. Eaa		Naa	
1	-	-	++	+++	++	+++	-	-	+	++	-	-
3	-	-	++	+++	++	+++	-	-				
4	-	-	++	+++	++	+++	-	-				
7	+	+	++	+++	++	+++	+	+				
9	±	++	++	+++	++	+++	-	+				
10.	+	+	++	+++	++	+++	+	+++				

(This agrees with the previous interpretation that 7-10 may readily back-mutate. 1, 3, 4 should be tested further. of (1).

Note poor growth in e.g. 9. Evidently, strong inhibition is present. Ws. is inhibitory??

broz: dil. from HC
9P24. 36h.

1: T()

lypt	-					isol.	-	++
thr	-							
lys	-							
arg	-							
dal	-							
meth	-							
v-i	++	+++						
leuc	-	+						
hist	-							
HC	+++							

Repeat 9: broz dil from CM. 24h. 1st. 36h 2

T (VLT)	-	+	T (0)	++	+++	T (HC)	+++
vlt	-	+	0	++	+++		
vlt	+	++	0	++	++		

Test 160 [156-1] for adaptation.

161.

9P26.

37°.

162

	broz. into 0.	12h.	24h.				
1	HC	0	-	Therefore these responses are key way of growth rather than adaptation.			
2	v-iso.	0	-				
3	lc	0	-				
4	cool.	0	-		1100A27	1120A18	
5	from HC				330	330	
6	" vit	into	CM slants for testing		+	+	
7	" 0.				-	-	
			dilute + test				
			2P27.	5	+	+	
				6	-	-	
				7	-	-	

Test 160 for growth vs. adaptation - 9P26. into T(0)

		12h.	24h.	48h.	920A29	10A30	test no	A29	10P29	10A30
1.	from HC	-	-	-	++				±	++
2	" v-i	-	-	-	2P27					
3	lc	-	-	-	-					
4	cool	-	-	-	-				±	

Test 156-9 substrans.

	a. to CM slants	5	from HC	6	from vit	from 0	0A29	4P29	10A30	
			1120A18	330P18	8P28	A29	Test 3	0		V
	to vit + 2P27	HC	{ 5 - vit	-	±	++			±	++
	min.		0	+	++	+++			+	++
		v	{ 6 - vit	-	±	+				
			0	-	+	+++				
		0	{ 7 - vit	-	⊕	±				
			0	-	+	+++				

This behavior is remarkable. What is inhibitory? What is the type of genetic modification?

See 164.

Mutant identification.
"Auxanography"

June 7 1978

histidine

→ (Hydrolyzed Casein)

E-8

161-6

ca. 10^{4-5} colonies per plate. They are visible for a radius of 1 cm in both cases around the HC, then thin out somewhat.

161-6 - not scattered large colonies, but quite numerous small. ∴ not "adaptation". ∴ vitamin effect is directly on growth, not adaptation. Sp. of a.a. not clear. Conspicuous faster on e.a.a. than on s.a.a. but this may not be a specific response. Need 15 for control. Inhib. by ketosis??

p 37 ca 10-12 "colonies" full size are seen in the 2-2 plate, presumably adaptants

Try Brewer's inoculum.

See Auxanography - p. 168.

3/28

Strain identifications, preliminary.

163a

Plate histidinless (E-8) and 161-6 heavily into 1% agar.
When solidified, add a loopful of

- (a) HC
- (b) histidine .1% to each.

7P28

Start "washed agar #1" for biotin. Agar washed by 10 transfers through distilled H₂O, + 2x in 95% alcohol, dried in desiccator over ZnCl₂.

Plate "58" (biotinless E coli) in 2% agar unwashed

P31 - no colonies!

" " washed
" " washed + 1% biotin

noc. on surface (streak) 8P31.

A2 - Well developed colonies only where biotin was added. None elsewhere!!!

Auto-antiserum

Test various T-L. standards.

Retest 156-1 142-17.

164

Test 156-2, 3.

4 1130 P28 inoc.
 1 930 A29 .1
 2 12 N
 3 9 P29

	0	TL	TL(HC)	TL+caa	TL+neaa	
✓ 1	142-17	---	---	+++	+++	= 410
✓ 2	-36	---	---	+++	+++	(adapted??)
✓ 3	-54	---	---	++	++	(19)
✓ 4	-57	---	---	++	++	m.g.
5						
✓ 6	0	HC	neaa	leuc ^①	isoleuc ^②	val ^③ + leuc. val ^④ + leuc.
7	156-1	---	---	+++	+++	+++
8						↑ check!
9	0	HC	neaa	caa		
✓ 10	156-3	---	---	+++	+++	
11						
✓ 12	156-4.	---	---	+++	+++	
13						
✓ 14	E-1	0 ^{10A30}	HC	neaa	caa ^{10A30}	
15		---	---	+++	---	
16						
✓ 17	Test 156-9	caa	0	HC	V1, V2, V3, V4	
18	various vitamins	+++	---	+++	---	+++
19						
* 20		neaa	V6	V7	V8	V9
* 21		+++	---	---	---	---
✓ 22	E-6	0 →	---	---	---	---
✓ 23	E-1 + E-6	0 →	---	---	---	---
24	(5x).					

Inoc. 530 P9. Inoc. 1030 P Retest 142-17 - autogram.

9A10 - D+++ A+. Inoc. 2 vitamins.

Many adapted colonies 12M10. No response.

to new plate 4 P10. Incubate 4h. before Inoc. 2 vits. etc.

12M10 - Thiamin.

Antis plate
 incubated
 + turbid

3/29 --

Identify mutants on hand

1. 12N30
2. 4P30

165

12M30 noc.

12M31 noc.

1) 4P31 2) 10A1. 3) 2P1

		val	δ	try	lys	arg	meth	hist
	TL	TL+3	TL-4	TL-5	TL-7	TL-8	TL-9	TL-10
1	142-17					- ±	± ±	+++
2	142-36 N.G.	- +++	+++	- +++	± +++	- +++	- +++	- +++
3	142-54	- - -	- - -	- - -	- - -	- - -	- +++	- - -
4	Postpone							
5	note							
		Checks 4/2/4/6 TL-9 +++ no others.						
		✓ 4/3/6 ∴ probably both methionine.						

0 cuts.

	11	10	1	2	3	4	5	7	8	9
4	156-1			± ± ± ✓	± -					
5	156-3	-	- - ✓	+ +++	± ± +++	- ✓	- ✓	- - ✓	✓	- ✓
6	156-4. ± ±	- -	± +++	± ± ±	-	-	- -	- -	-	- ✓

No further growth by 10A1.

(1) 10A30.

Checks: 4/2 - OK.

* my own paper. 8. 1

other is Tatum's. may not be enough val.

See infra for recheck on 142-17 + 54.

5P4. - Plate 142-17 + 142-54 into T(TL) agar. 1ml vol. incubate to 9P4, then to auxanogram on essential a.g. or 54, only "9" = 17 using double depth agar.

	HZ	B	1	2	3	4	5	6	7	8	9	10	C	D	!	Checks!
17	++	+	-	-	-	-	-	-	-	-	-	-	-	++	!	8PS
54			-	-	-	-	-	-	-	-	+	-	-	-	-	1PS

Check 54 on liquid 10PS. - 12N7(1)

∴ 54 is TL Meth.

	M	MT	ML	MTL	TL
	-	-	-	++	-

Hydrolyzate C.

166a

200 (2+100) 7(10) 72 hour culture. Centrifuge cells down,
put in 10 ml 6N HCl, seal tube + keep in boiling H₂O for 2 1/2 h.

lost during hydrolysis.

Try again.

3/31 - 4/1

167a.

Mutants by ultra-violet irradiation.

10A 31. Inoc 50ml / 125 ml flask coli C-M \bar{e} 58 (Tatum's biotinless coli) and grow on shaker, slowest speed, at room temperature.

① 9A1 - 1ml sample to 50ml coli CM.

②. Irradiate in quartz tube, 11 cm from tube, 15 ~~sec~~ min.

Inoc. 1ml into 50ml coli CM. Grow 1, 2 on shaker.

No appreciable growth in 24 hours. Dosage too high.

Try 5 minis.

1 colony at 1:50 dil.

P2 finally came up.

A2. Use 167A1 + irradiate as above, 5 min. Do in dupl.
11A2.

Estimate (a) before irradiation.

Dil $\frac{1}{500 \cdot 500 \cdot 50}$ + plate into YBG. 1a.

b. after irradiation.

1:1	} 50 ⁿ	710.
1:50		9
1:2500		
1:125 000		
1:6250 000		

c. In (a) prepare last dilution in saline also. Inoculate + compare sal. + H₂O after 48h. Do in dupl. 1c.

~~d. Test colonies from b (1:1) and use for 50/a~~

~~studies:~~

2a + 2c 10P2. O O
3a + c 930P3.

~~15~~ 15 minute irradiation

A1.

Before irradiation, plate counts not made
after 1:50 -- 1.

Proc C-M 2 1 ml. 2 P 3 dilute $1:25 \times 10^6$ + use method II,
plates 1-5 for mutants (T(0) + Dr. Kotei)
Colony test apparent 9 P 4. 10 A 5 Layer YBG. (Agar too soft).

2 P 4. 2 Plates so soft as
to be almost useless.

4
7 (0) 5. Pick colonies before replate. $k_1 > 8$ / radiation -
YBG rest on T(0) - all yeast - resistant.

5 minute irradiation

A2

see 172

Control plate counts: *1a - (last dil in 0)
1: 12,500,000. *1c - last dil in saline.

if said
trivially
steadily

after irradiation, 1b : 1 ml 710 } not in, different incubate &
1:50 9 } shake alternately.

c) Viability of control in sal, H₂O. - Apparently very low in this selection

			i	ii
incubate last dilution	10 P 2	2 a	0	0
flasks of 1a, 1c - 2 sample.	10 P 2	2 c	0	0
Plate in YBG	P 3	3 a	0	0
	P 3	3 c	0	0

I 10A1. halo 2 cm diam. 2 centimegs around H.C.

Nothing over 10^{-3} dilution. By 8P, there was a very faint response to H.
At 8P, 7.5 cm diameter.

II Supplement 10A1. By 2P, a distinct turbidity was visible, ~~was~~ over HC & a faint one over histidine. By 8P this was very distinct & sl. less impressive over histidine. Both ca. 2.5 cm diameter.

3) Supplement other portions of both plates, as above, 8³⁰P. (after prolonged incubation).

10³⁰P - 1B - dist turbidity under HC.
— 1C better.

12 hour incubation probably optimal
inoculum size is also or less optimal + maybe reduced for frequently
occurring types.

3) Add HC 9P4. - No response dead?

Incubation, time at ca 35° unless stated.

1689.

Auto-auranography

8P31 - Plate heavily (ca 10^6) into T(0) 2% agar E-8 (histidine)

I Add, as cooling drops of HC + histidine (10%, .1% resp.)

+ dil. 1:1000 resp.

II Incubate 14h. fast.

III Cover + incubate to pH. then ~~log~~ ^{add ~~XXXXXXXXXX~~ HC.} to determine survival.

P2 - # 48 adapted colonies. ✓

P4 ✓ Try more conc. agar.

IV 4/2-3. Try as above \bar{c} 3, 4, 6% agar.

No difference to spec. of.
 \therefore 2% is opt.

V 4/3 - Use of indicator - plate E-8 as above \bar{c} 10v/cc
Methyl Red (also 20v/ 50v/). Medium is all saline to
the indicator.

4/2/46.

I Plate 161 - Thawed in 20% T(0) agar plate. Add suppl. HC + Biotin to surface 10P

10A - Turbidity increased over HC.

~~Restonably~~ clearer ~~under~~ under Biotin. (logful 1r/ml)

5P. do. The biotin area is definitely less turbid than the rest of the plate. The plate is ~~fairly~~ fairly dense but somewhat darker under HC.

9P4. Differentials essentially disappeared. ∴ 156-9 is not an adaptor, but perhaps a slow grower & perhaps inhibited

by biotin. Check \bar{c} L15! (At least adaptation is not genetic or "mutational")

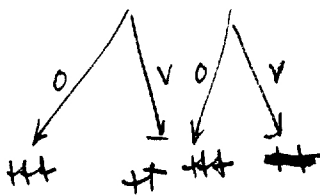
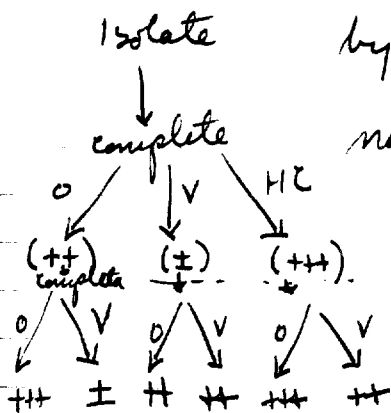
See 173.

Study of 156-9.

1. Isolated by Method C. from X-rayed L15.
2. Responds rapidly to HC, but grows on min.
3. Is slower on vitamins, or on biotin, but eventually grows up.
4. No "adapted" colonies seen on auxanography on (st attempt (163.)
5. Exp. 161. Grown on HC, Vits and O. (Exp. 160). Then transferred to complete slants, growth tested on O + V. To all appearances, that grown on minimal was most sensitive to biotin. Apparently an exposure to HC or to V limits response to V, but not an exp. to complete.

Also, growth on minimal reinforces susceptibility. This is not supported

by the final results on the HC line!!! The fundamental nature of the phenomenon is not clear. It should be more stickering on plates than otherwise. Also, all these results have to be checked.



I

monoclonal 10⁶

[Signature]

Data

	Medium	Date	Require-				
1. 58.	T(0)	4/3	Biotin	1/4. 0	4/5 0		
2. E-6	T(0)	4/3	Methionine	1/4. 0			
3 58-5198	T(0)	4/5	inole	0			
4 58-5417			inole	9			
5 58-5636			inole	>10	(plate microally (inhib) compare 5298)		
6 679-680	T		leucine	7			
7 "	L		leucine	1			
8 "	0		L	0			
9 679	0		threonine	2			
10 58-5631	0		alanine	1			
11 58-161	0		methionine	0			
12 58-7621	0		proline	0			
13 58-5173	0		alanine	0			
14 58-118	0		leucine	0			
15 58-336	0		isoleucine	0			
16 679-682	0		proline	0			
17 679-183	0		proline	1			
18 58-4899	0		leucine	0			
19 58-3214	0		proline	0			
20 58-3336	0		meth	0.			

Mutant Reversions

170a.

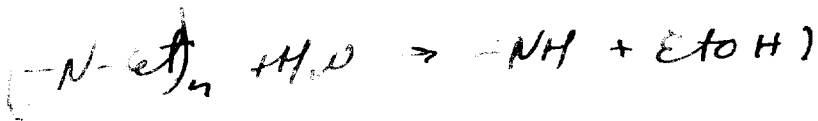
Now ca. 1 ul of inoculum (ca. 10^6) into T/C, as indicated and incubate 48 hours; look for adapted colonies.
Data on 170a.

Mustard

E-3-Histidine sep 168

58 (Pectin).
E-6 Methionine. } Probably excess carryover: plates turbid.

Mustard ident — inconspicuously back mutate also.



Check for sister experiment: 58-565a

Sex.

4/3/46.

5P3 - Cross streaks on minimal plate:

E-6 (methionine), 58 (biotin) + TL.

P5 no growth.

9:30 P5 - In minimal liquid, the following:

1	679-680	T-L
2	"	"
3	"	"
4	58-278	B-Ø
5	"	"
6	Both	"

No growth in any by P9.

N9. Repeat above \bar{c} heavier, fresher mould. in T(0).

1	TL	±	±
2	TL + TL	±	±
11	TL + BØ ₁	±	±
12	TL + BØ ₂	±	±
13	TL + BØ ₁	±	±
14	BØ ₂	±	±
21	BØ ₁	±	±
22	BØ ₁	±	±
31.	In complete TL + BØ ₁	-	+
32	" TL + BØ ₂	-	+
33	In T(0) BØ ₁ + BØ ₂	+	+

~~nothing by P11~~

Note

Try plating out from C-M.

2P9 Cross streaks on T(0) agar

- 41. BØ₁ x TL
- 42. BØ₂ x TL.

nothing by P14.

something happened in the complete cultures between the 1st + 2nd tests in min.

19427 200
 11/10/42
 4/19/43

	Plate #	Total	Mutants	Mutants	# s.y.		Rate
			1230 A7	1130 A.7			
	1	218	3 ¹⁻³	1 ⁴		1	5
	2	213	4 ⁶⁻⁹	5 ¹⁰⁻¹⁴		2	15/11
	3	220	2 ¹¹⁻¹⁸	3 ¹⁹⁻²¹		1	22
	4	194	5 ²³⁻⁷	3 ²⁸⁻³⁰		2	31-2
lost	5	209	2³³⁻⁴	2³⁵⁻⁶		0	
	6	160	1 ³¹	0		0	
37/	7	1214	17	14		6	
		retest					
Y(58--)		T(b)	HC	Vits.	Auxanogram.		
	1	+	✓				
	2	+	✓				
	3	+	✓				
→	4	58-411	-	✓		MC; D?; ?;	
	5	+					
	6	+					
	7	+					
	8	+					
	9	+					
	10	Proline 412	+	±		A. 15 C?	
	11	58-412	-	X+		hybrid plate sl. 14h. MC	
→	12	413	+	-			
	13	58-413	-	X+		hybrid plate	
→	14	58-413	+	-		B?	
	15	414	+				
	16	+	✓				
	17	+	✓				
	18	+	✓				
	20	+	✓				
	21						
	22						
	23						
	24						
	25						
	26						

Ultra-violet radiation: See 167A.

Irradiate 58 coli. 5 mins at 11cm viguacity tube on shaker.

Procellum before irradiation /ml. ~~4×10^9~~ (8×10^{-8}).

$$i. 1a. (455) = 5.7 \times 10^9$$

$$ii. 1a. 97. = 1.23 \times 10^9$$

$$iii. 1a. 300 = 3.75 \times 10^9$$

$$\frac{107}{3}$$

$$= 3.6 \times 10^9$$

$$1c (249) = 3.12 \times 10^9$$

$$1c (129) = 1.64 \times 10^9$$

$$1c. (252) = 3.1 \times 10^9$$

$$\frac{7.8}{3}$$

$$2.6$$

$3 \times 10^9 = \text{mean.}$

$$\text{Survival} = \frac{710}{7111111} = \frac{7 \times 10^2}{3 \times 10^9}$$

$$= 2 \times 10^{-7}$$

.00002 % survival. in 5 min.

[Since there were ca 50/ml after 15 mins, there must be heterogeneity in u-v susceptibility.]

Shake 48 hours + diff: 12.5×10^6 . (9p4). Plate 4 in T(13).

2:30 P6. Layer.

12:30 A7. Pick mutants (ca. 10%) see data.

[See 183]

All mutants this time were picked 21 hours after lagging.

score in this group is 5/12.

3 plaque.

5 mutants in 1200 cells.

comp. \bar{x} part.
5/6000.

	found	Exp.
sp	5.	1.7.
uv.	5.	8

$\frac{10}{7200}$ Exp. are too small.

$$\frac{(8-5)^2}{5} = 1.8 = \chi^2 \text{ too small for sign.}$$

also test:

			I (Brd).	HC	V. ts.	Auxiliary.	✓
Proline	27 28	y15 58-444	-	✓			
Proline	29	y16	+	✓		HC 12; ?;	
	30	58-445	-	✓		HC large zone; D; A	
	31		+	✓			
	32		+	✓			
	33		+				
	34		+				
	35		+				
	36		+				
	37		+	✓			
Tatum's sp. control.			+	✓			
	58-278	"	-	✓			
Total: 5 stable							
1214 colonies.							
5580.						HC; AAA???	

4/9/46

156-9 vs. L15.

Biotin

noz. 3P 9 asund.

(1) 9A10.
(2) 9P10

	0	0	V.+	Brot.	value
L15	+++ ✓	+++ ✓	- ++	+ +++	+++ ✓
156-9.	+++ ✓	+++ ✓	- +++ ±	++ ++	
K12	+++ ✓	+ ++	- +++	- ++	
Biotin is then not the only factor + there is some inhibition by a vitamin of 156-9.					

Add 1 drop 1N HCl / 10cc
for preservation?

Also consider:

Eff. pH in drops (for presp.)
benzene

Concentration & amt. of substrate.
(Res might be sharper after pre-incubation).

4/9/46.

Auxanography

Optimal inoculum size. Plate into T(B) varying samples of an inoculum contg. ca. 10^8 /cc of 58-278 (B- ϕ)

Incubate 12 hours & supplement \bar{E} HC + \bar{E} ϕ . T(B) = Phenylalanine

noz: 530P9 Suppl. 1030 9A10.

	Diam. ϕ	Heain H.C.
1 ml	1.8 ++	2.3 +++
.1 ml	22.5 ++	3cm ++
.02 ml	32cm ++	5cm +
.002 ml	4 ? \pm	2cm +
.00004 ml	5 ϕ 2	+ 4cm +

The method can be used at any inoculum size, but is most sensitive & heavy inocula. For very adaptable strains, it may be important.

↑
↓
distinctness

OPTIMAL AGAR DEPTH. 4/10. noz. 1ml undil culture into varying agar depths. noz ~~1030~~ ~~incubate~~ to 1130P9. Supplement A - ϕ - at t=0. B: ϕ at 1230P11.

	A (11A) 730P	B. 230P
1 5ml	nothing!	fair response.
2 10ml		
3 15ml		
4 30ml		

It makes very little difference what depth agar is used 10-15 ml is quite OK.

TIME OF INCUBATION. noz. 58-278 into T(B) agar 9P11

Suppl.	t	9A12	7P12
	0 10P	0	\pm
	0 12M	+	+
	0 10A12		
	0 10A12	##	-
	0		
	0		
	0		

quality of zone.

optimal < 12h.

1:25,0000

sample is 6000.
ca. 5 mutants.

compare 175.

~~hydrolyse A 1:10.~~

* Contam. = Neurospora + thermophil!

4/9 - 10... 1/56.

Spontaneous mutants in 58.

530P9 broz t.t. E coli CM E 58. Shake at RT.

9P10 - dil 1:12,500,000 + plate out by method II in T (100% in).
 Incubate at 35° to 11P11. Plates covered!
 Count 1130A12 6P13.

*1

*2

*3

4 1000 1

5 " 2-5

6 " 6

7 " 7 8-9

8 " 10

9 " 11,12

*10

12h. 30h.

Add. 1ml of "A/10" in lieu of bacteria :-

11.

Picks to CM 830P14 n.v. satisfactory

None of the 1st series would have qualified except by sterility
 criterion. (B). v. in. inoculum from colligrid.

1 +

2 +

3 -

4 +

5 +

6 +

7 -

8 -

9 -

10 no growth in CM

11 -

12 n.v. in CM.

Done 1-15

July.

921

4/12/46

Proc. C-M E:		4 shalae			A12 - P13	
	1.	FL				
	2.	B φ ₁				
	3.	TL+B φ ₁				
					A18	P19.
12 M13.		T(10)	P16.	P17	# P17: add	
Test 1	4	-	-	-	1 loopful of	-
2	5	-	-	-	complete	-
3	6	±	+++		coli medium	
1+2	7	±	+++		to tubes	
					4, 5 and 9	
3	8	-	-	-	13, 15+17	-
1+2	9	-	-	# -		- killed?

See 171. Plate 31. into T(10) + cover. M13. (TL+B φ₁ in C-M).

1: 25000
~~1: 500~~

10 } No colonies // Cover φ A1+B.

1: 12,500,000

11 } P16 // Nothing came up

P17 A18 P18

12 3 into CM, 5 shalae at 30° 1030 A13

small mic. 13 Test: 0. - - # +

large mic. 14 ++ +++

P16. Test 6 mm 15 - ++ # ∴ the transfer is as infert.
6 large 16 - ++ as in culture count of cells!
1 mm 17 - - # These have no recombination.
1 large 18 - - -

Preservatives.

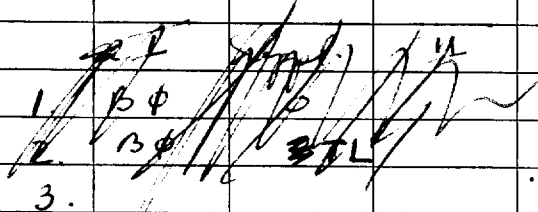
I Plate K-12 in T(10). Add drops of HCl - detected.

Center #	Conc HCl (10N)	Yeast inhibition	
2	1:10	1 cm width.	
4	1:100	No inhibition	try 1 drop HCl / 10 cc medium, etc.
6	1:1000	No inch	
8	1:10 ⁴	No inch.	

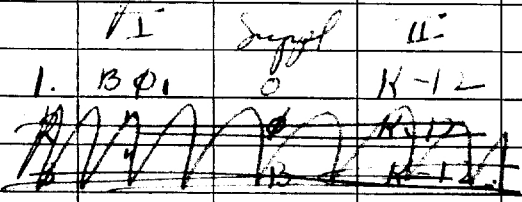
II. Benzene.

1.	Glycer - pure Benzene	Clear zone
2.	10 - Benzene-water.	OK
3.	5 - Alcohol 95%.	OK
4.	10 - Benzene Chloroform	OK.

Pour plates \approx ca 10^6 organisms mixed + suggest. where hard, streaks 11" on surface.



4/17



See N-363 for exp. on symbiosis with N: 33757.
~~E~~ E 181.

A19. About each streak is a clear zone, ringed \approx a range of greater density, fading off to a small turbidity.

As complete P19 - No differential zone found. K-12 somewhat inhibited.

Salt resistance mutants.

180

after devereux & Turner.

4/17/46. 581

1. Plate ~~H₂O~~, ca 10^3 , into complete plates +

1. CuSO_4 ~~5mg~~ 10mg

2 HgCl_2 .1mg

A19. Nothing came up. use less.

A *Penicillium* sp. contaminant
did grow on D!

4/19/46

held!

As before:

		Supplements.		FPA.
		Ø	Tyr.	
5	broc	Ø		
51	K-12	0	0	10r
52		0	0	100r
53		10r	0	100r
54		100r	0	100r
55		0	10r	100r
56		0	100r	100r
57		10r	10r	100r
58	58-278	1r	0	0
59		10r	0	0
60		100r	0	0
61		10r	100r	0
62		10r	0	10r
63		10r	0	50r
64		10r	0	100r
65		0	10	100r
66		100r	0	10r
67		100r	0	50r
68		100r	0	100r
69		10r	10r	100r
70		0	0	50r 100r
71	58-4899	0	0	100r
72	58-5030	0	0	100r
73		0	10	0
74		100	10	0

Many aspects of this experiment are consistent with the investigation of the aromatic a.a. mutants + are to be postponed until this is carried through by ELT, et al.

4/19. Utilization of FPA:

		430P20	³⁰ 3P22	9P22	P23
81	58-278	10	-	96	-
82		100	-	97	-
83	58-4899	10	-	96 ²	-
84		100	-	97	-
85	58-5030	10	±?	95	75 ³ †
86		100	-	67	65 ¹ †

† test on minimal: they grow.

Mutant detection:

Viability

(with shaker)

* Pour 58-278, dil. to 1:12,500,000 into T(6) plates + cover as in mutant detection. 830 P19. Cover complete at time t. Colony diameter recorded at t. Incubate at 38°.

St.	hour	h ⁺ 430P20	h ⁺ 830P	10A21	4P21	8P21	12M21	Count.
1	9P19	1	19	5	+++	irriduable variation.		
2	"	1		4		← do		362
3	1130	3	17		+++			352
4	"	3		6		← do		
5	930A20	12	7	< #1.	+++			379
6	"	12		3		← do.		407 (41?)
7	830P20	23			++	+++		363
8	"	23						5
9	10A21	36			-	±±±	+	380. ← too many bottom colonies
10	8P21	48.						349
11 1130 10								353
12	+ 1mg β		6					

Bottom colonies troublesome. 7 hours is barely too long for period II. Past runs u.g.

Repeat 9P20: clearer agar. as above. Pour bottom layer. Shaker culture is complete. Incubate 30°.

St.	hour	7:15P.	130P.	10A21	4P21	8P21	12M21	10A22	P23	Count.
21	9P20	+	++	++	+++					131
22	9P20.	+	++	++	+++					139
23	1P21		±	+	++					138
24	1P21.		±	+	++					151
25	8P20			-	+	+++				23
26	8P20			-	+	+++				23
27	930A22					++				36
28	"					++				36.
29	3P23					++				64
30	1130A24.	25°	130P.							54
31	"	30°	colony							156.
32	" (2x)	38°	mini distinct !!!							145
33	CM.-P.C.									135
34	Plate Count									340. (2x)
35	"									146

Uneven pour: lumpy. Colonies quite distinct mini. indicated (3-5h) 1mm. colonies 2mm colonies 3 1/2 da. 86.

Count. ΣΔ² = m = 140. 1097. σ = 8.85 σ_{calc.} = 14.

Viability of 58-278 at 38° is excellent for 48 hours.

* 7 small colonies noted 10P27. puncture.

Rick colonies 11A30. see 194.

* 15 units = 1mm.