

50ml cultures, shaken at 30° K-12 viorc 1030A21.
6-d.

- 2 ~~z~~. glut 5mg, putkastone 1mg, β -alanine 1mg
- 3 ~~z~~. prol 5mg prot. 5mg

- 1. T(0).
- 4. See 202-9.

To test top N22
sample a 10P22.

- 1. Ta 5ml + F(0) 5ml. N. crassa 5531.
- 2. 2a 5ml + F(0) 5ml N. crassa 5531.
- 3. prot series 1r, 5r, 10r " "
- 4. 3a .1ml + 10ml F(0). " "
- 5. " .5ml + 10ml F(0). " "
- 6. 4a 5ml + 5ml F(0). " "

- 11. 1a 5ml + T(B)
- 12 2a " " ²⁵
- 13 Peptide 25r + 1a ~~25~~ .3cc + glut 25r.
- 14 ~~3a 25cc + T(B)~~
3a 0.25cc + T(B) 10ml.

58-3214. inc N23.
±
±
±
±

~~hydrolyze 1-3 (wash 3 cells) in 6N HCl. 14 hours.
total o.d.u.~~

- 1. 10.400
- 2. 12.000
- 3. 4.210

Streak out 207-1 mac coli plate P22. Isolate 10 colonies to H₂O, and test; transfer to ∞ slants. 3P23.

\therefore ~~the~~ threonine not required.

	01	T-B	T-T	T-BT.
1	+	++	-	+
2	+	"	-	"
3	+	"	-	"
4	+	"	-	+
5	+	"	-	+
6	+	"	-	-
7	+	"	-	+
8	+	"	-	-
9	+	"	-	-
10	+	"	-	+

Test 207-2-4.

11P23.

	0	BPMT.	BM	BT	MP	MT.	BP	TP
2	-	++	-	-	-	-	-	-
3	±	++	-	-	-	-	-	±
4	+	++	-	-	-	-	-	+
5	-	++	-	-	-	-	-	±
8	±	++	±	++	±	++	±	++

24h 32h.
++ Maybe a multiple mutant.

Finally: +.

- 2 - BPMT, PT. 679-183. ✓ 212B.
- 3 - BTPT, MT, BPMT. 679-183⁺
- 4 - BPMT, BT, PT. MT(-)! 679-183+ 679-58x? 679-183⁺
- 5 - BPMT, PT. 679-183
- 8

Note: all these strains were isolated from BT plates!

Retest:

	BPMT	BM	PMT	MTB	BPMT	MT.
207-1	/	/	/	/	/	/
207-1A.	/	/	/	/	/	/
207-2	/	/	/	/	/	/
207-4	/	/	/	/	/	/
BM	/	/	/	/	/	/
BP	/	/	/	/	/	/
PT	/	/	/	/	/	/

See 212.

5/25/46.

P25.

From mixed cultures, strains have been obtained which behave peculiarly in their nutritional requirements, behaving for a time like recombinant types. For a demonstration of sex, a stable recombinant type is essential. An analysis must be made of cultures 207-1 and 207-2.

207-1 behaves like a culture of BT with a small % of B cells. Therefore plate heavily + lightly into T(B) + layer \bar{c} "T" after B is detected.

207-2 behaves like a culture of BMPT \bar{c} same PT cell still present.
Plate into 1) PT ~~BT~~ 2) BM. Afterwards layer.

207-4 may have BT ~~BT~~ cells. Plate from 210-4-BT into B. After-layer \bar{c} T.

5/26/46

I Cystine Requirement of Y24.

1A26. A. 10ml T(B, φ) + B: .04r φ 300r

	Cyst.	
1.	10r	89 ¹
2.	30r	84 ¹
3.	100r	83 ²
4.	300r	76 ³
5.	1mg.	68

← 1/2 opt. like 58-309.

11. Methionine: 1mg. ± not parathiotyrosyl.

12. Meth. 1mg + Cyst 1mg. 65.

II Other strains:

	T(0)	HC	VITs.
1	B	Y13.	
2	B	Y14	
3	B	Y18	
4	B	Y20	
5	B	Y21	
6	B	Y22	
7	φB	197-61.	
8	B	197-32	
9	B.	197-33	
10	Bφ	-5	
11	Bφ	-7	
12	Bφ	-21	
13	Bφ	-23	

Analysis of 207-cultures Sex.

5/26/46.

Scrape growth from slant, suspend in H₂O & dilute as indicated. ^{washes well}
 To obtain clean agar, autoclave 3% (agar + T(10) 200% quantity. ^{Thurs. morn. Mutant selection procedure.}

12M-1A26.

~~Use T(10) i.e. 1/2 usual phosphate [T₁]~~

A. 207-1. P27. ^{laymē} Added col. 11 P207. 11A28.
 1:1000 1h:

B ⁻ M ⁺	1.	B	+++
	2.	B	+++
	3.	BT	+++
	4.	T	0
	1:10 ⁶		
	11	B	51
	12	B	39
	13	BT	43
	14	T.	0

Pids to complete 1728.
 For analysis see 219.
 requires B.

207-2.

1:1000 1h:

B.	21	PT	+++
	22	PT	+++
	23	BMBT	+++
	24	BM	—

1:1000000

31.	PT	9	
32.	PT	8	
33.	PT	15	∞ 1 #1
34.	PT	14	∞ 1 #2
35.	BPT	12	
36.	MPT	12	
37.	BMP	0	
38.	BMT	0	
39.	BMP.T.	14	

C. 207-4

1:1000

41.	T	++	
42.	B	0	
	1:10 ⁶		
51.	B	0	
52.	T.	3	'B 1 #3

FPA.

dl-fluorophenylalanine

5/24/46.

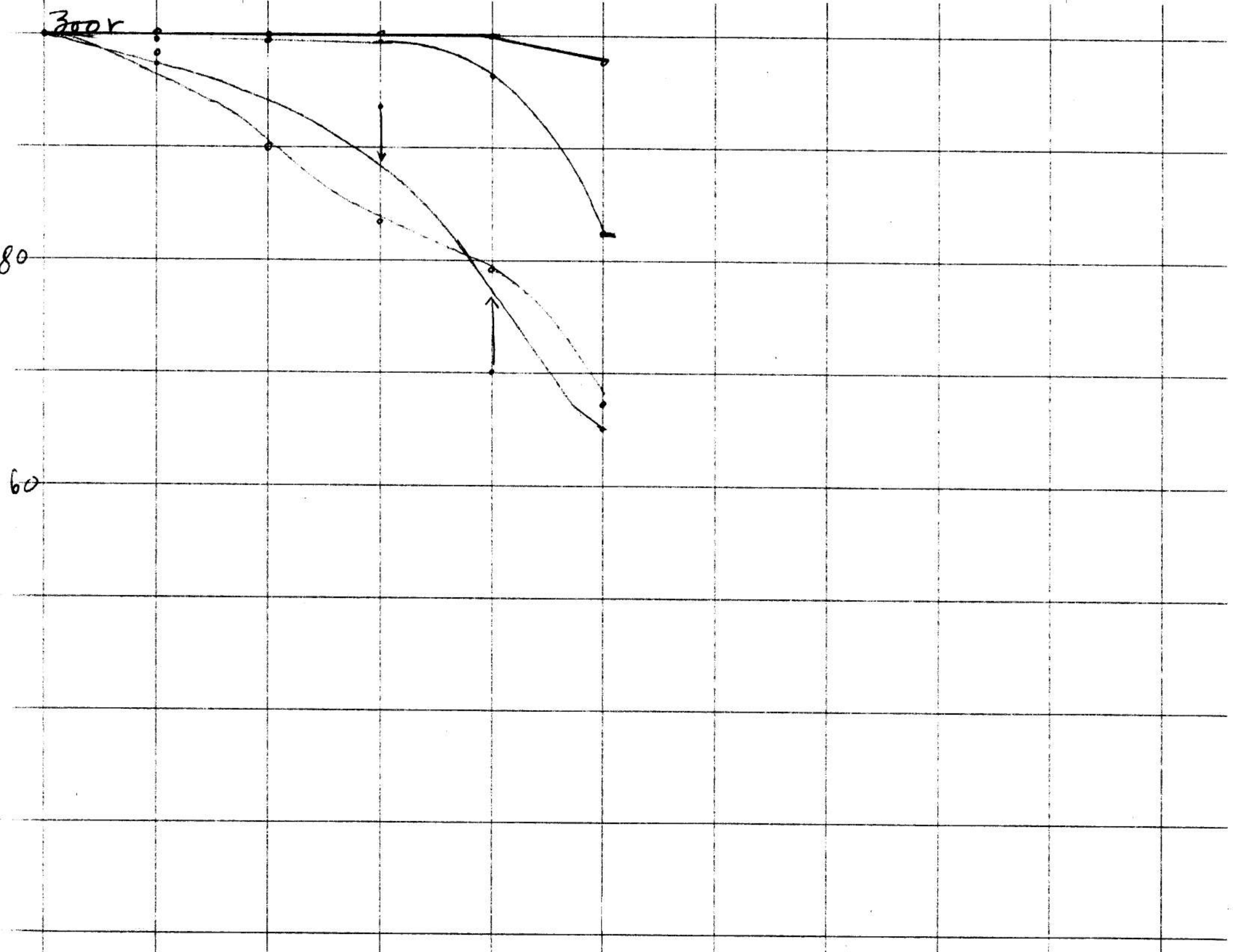
10ml T(B). 30° in columnar tubes.

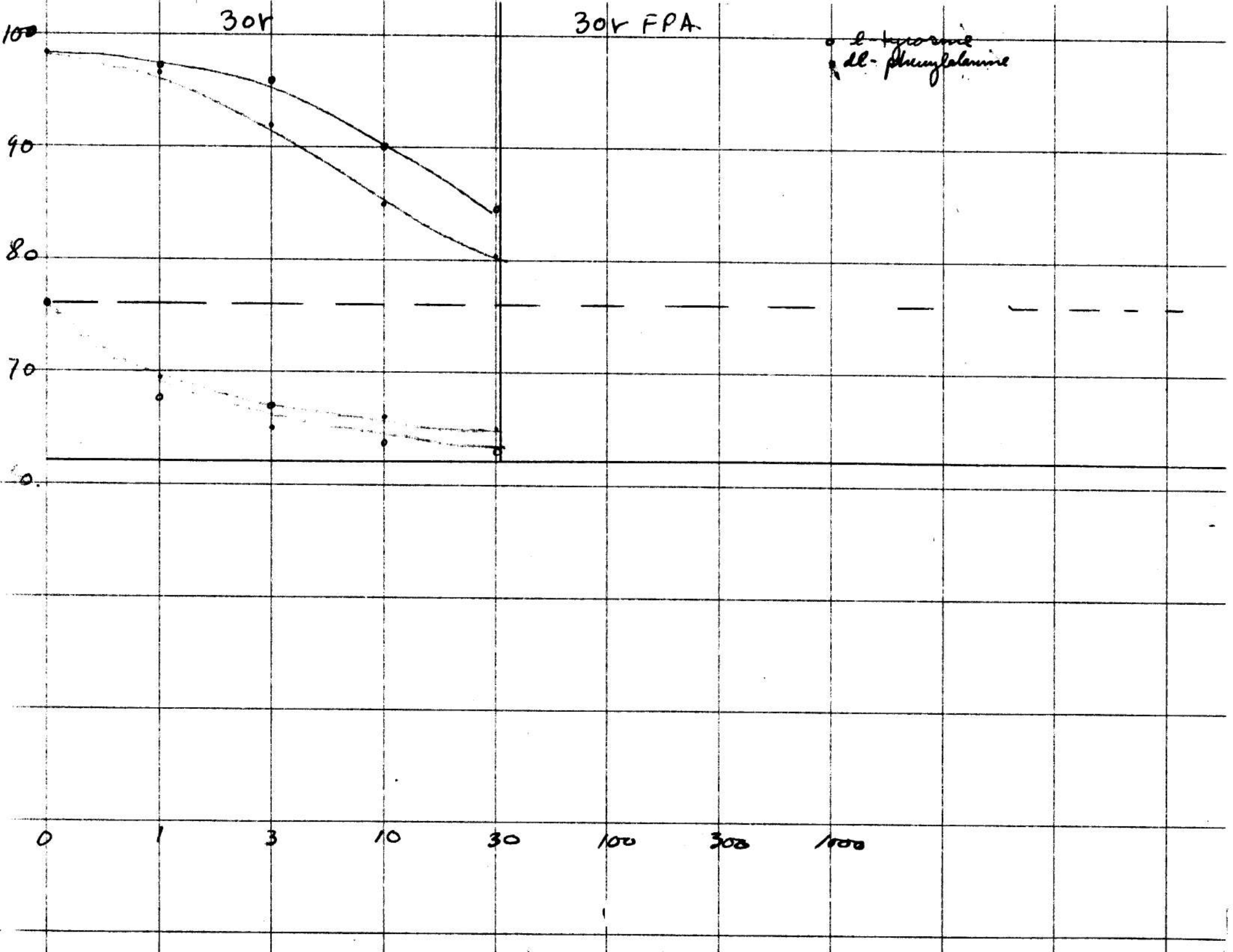
Proc. 1A26.

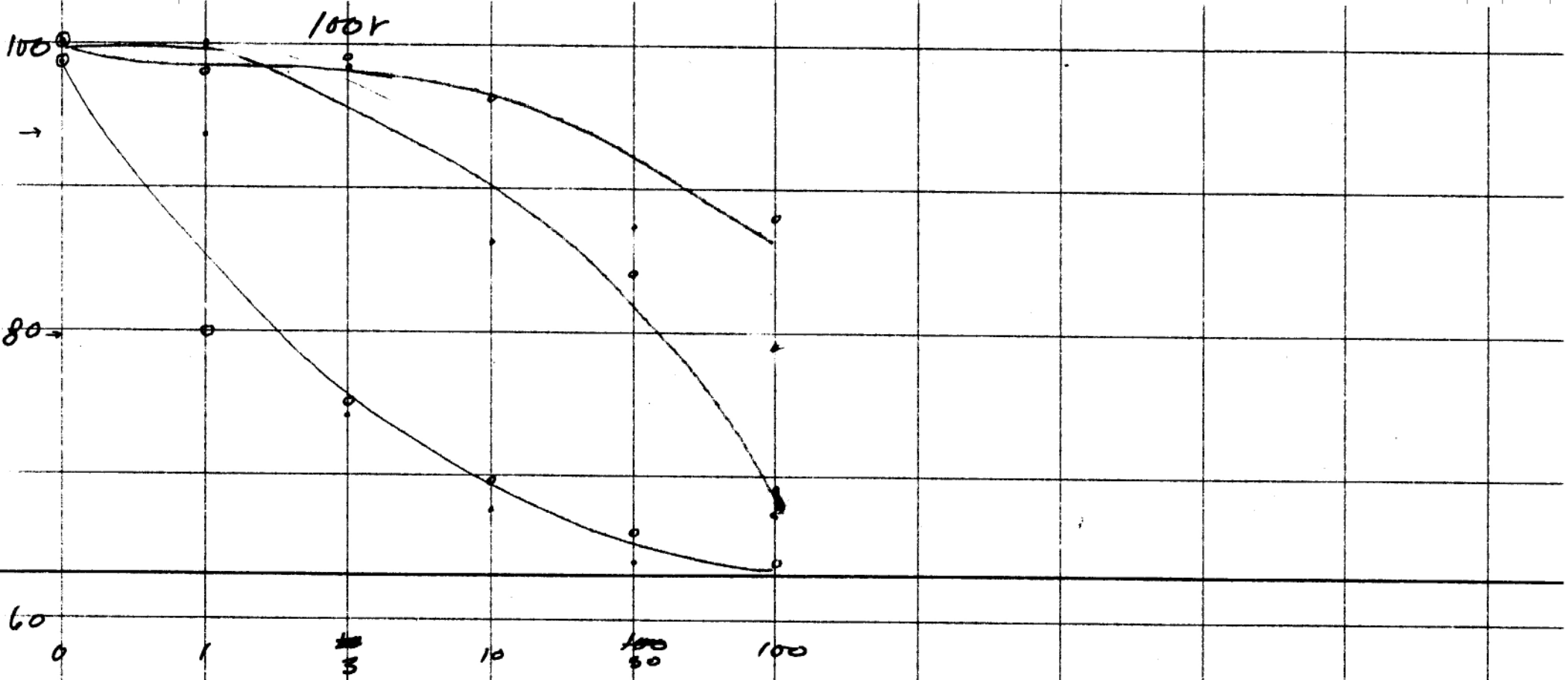
Proc. dl OAL. dl FPA.l-tyr.

				8P27.	1P28
1.	R-12	0	0	67	68'
2.	"	"	10v ✓	80	68'
3	"	"	30v ✓	97	75 ²
4	"	"	100v ✓	100	89'
5	"	"	300v ✓	100	100
6	"	"	1 mg.	100	100
<hr/>					
11.	58-278	100v ✓	0	70	70
12	"	"	10v ✓	66 ³	67'
13	"	"	30v ✓	67	70
14	"	"	100v ✓	69	67
15	"	"	300v ✓	70	69 ³
16	"	"	1 mg.	99	96 ³
<hr/>					
21	58-278	200v ✓	0	72	68 ²
22	"	"	10v ✓	66'	67 ²
23	"	"	30v ✓	67'	68 ²
24	"	"	100v ✓	69	68 ²
25	"	"	300v ✓	69'	70
26	"	"	1 mg.	71 ²	71 ²
<hr/>					
31	58-5030	0	0	0	100 100
32	"	"	0	80v ✓ +++	77' 77 ²
33	"	"	10v ✓	80v ✓	77 78'
34	"	"	100v ✓	80v ✓	78 ³ 79
35	"	"	1 mg.	80v ✓	75' 75 ³
36.		100v ✓	100v ✓	0	- 100 100
37.		100v ✓	0	0	- 100 100
38.		0	100v ✓	0	- 100 100
				76	75 ³

compare E 133.

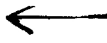






Inhibition of K-12 and Reversal by phenylalanine + tyrosine.

6/2/46.	Inoc K-12	11P2.	25h.	30h.	48h	60h.	
	FPA + dl αα + l-tyr.	4P3	12M3.	1FA4	11P4.	11A5	
1	30	0	0	±	98'	76	63 ²
2		1		±	96 ³	69 ²	65
3		3		±	92	65	68
4	→	10		+	85	66	
5	-	30		+	80	65	
6			1	+	97'	70 / 67 ³	67'
7			3	+	96	69 / 67 ³	
8			10	+	90	70 / 63 ³	
9			30	+	84 ³	63 / 63	68
11	100	0		±	100	98 ³	97'
12		1		±	100	93 ³	67 ²
13		3		±	99'	74	67 ³
14		10		+	86'	67 ³	
15	→	30		+	87'	64	
16		100		++ ^{sic}	69	67'	
17			1		98	80	70
18			3		99	75	66
19			10		96 ²	69 ³	73
20	→	30		+	84	66	68
21		100		+	88	64	69'
31	300	0			100	100	99 ³
32		1			99 ³	97'	78'
33		3			100	100	100
34		10			99 ²	93 ³	72 ²
35		30			96 ²	70	67
36	→	100		+	82'	65	70
37			1		100	98 ²	98 ²
38			3		99 ³	90	70
39			10		100	83 ²	68
40	-	30			100	79'	68'
41	m	100.		±	97 ³	67'	67 ²
51	0	1		±	85	61 ²	65
52		10		±	76	62 ³	66 ²
53		100		±	76	65	69 ²
61			10	+	77	62 ³	67 ²
62			100	+	85	60'	69 ²
63			1000	+	74.	66'	67 ²
				(80)	(80)	(283)	80



Inoc. Medium III FROM 11A6. Proc. 1130P5
 1145 210 450 750 11 220 445 ~~20~~ 550 810
 A6 P6 P6 P6 P6. A7 A7 ¹¹⁵⁰A7 P7 P7

Lag mgt.
1414

Hours + mins		45	190	350	530	820	920	1365	1490	1730	1990	
1	K-12 0	96	92	84 ³	69	58 ³	57 ¹	59				
2	K-12 0	97'	92 ³	84 ²	69	59 ²	58 ¹	61				
3	K-12 v.small 0	98	98 ³	98 ²	98 ²	98	96 ²⁵ → 8.5	73 ³	66	58		144.
4	" / 0	99 ²	100	99 ¹⁰⁰	99 ²	99	95 ³	72	64 ³	57		
5	K-12 FPA 100Y	99	99	99 ²	98 ²	95 ¹	90	67 ³	68 ¹	65 ³	67 ²	220?
6	K-12 FPA 300Y	99	99 ²	99 ²	98 ³	97	96	81 ³	77 ²	68 ¹	67 ¹	OK. 390
7	K-12 1mg. FPA 100	100	100	99 ³	100	100	99 ³	98 ³	100	97 ²	96 ³	
8	K-12 FPA 10	100	98 ³	93 ³	88 ²	76	67	69 ¹	69 ²	69 ³		175
9	K-12 " v.sm.	99 ²	98	93 ³	83 ²	74	67 ³	70 ³	70 ²	70		"
10	K-12 "	100 ²	100	100	100	100	100 ^T	99 ^T	96	87 ²	72 ²	140; 205.

11	K-12 FPA 100	99 ³	99	95 ³	87 ¹	77 ²	68	64 ³				220
12	" " TYA 10	100	99 ¹	96 ²	87 ³	75 ³	68	64 ¹				"
13	v.sm. FPA 300	100	100	100	100	99	99 ^T	96	94 ³	83	68 ³	
14	K-12 FPA 10	97 ¹	93	82 ¹	66 ³	59 ²	62 ¹	66 ³	65 ²	66		
15	K-12 FPA 300	99 ¹	99 ¹	99	98 ²	97 ²	97 ¹	91 ³	89 ²	77	67 ¹	
16	K-12 50% but(T)	100	96 ²	88 ¹	74 ²	69 ¹	71	73	73 ¹	73	73	
		77 ¹	77 ¹	76 ²	77							
				77 ¹	16 ²	76 ²	76 ³	77	76 ²	78 ²		

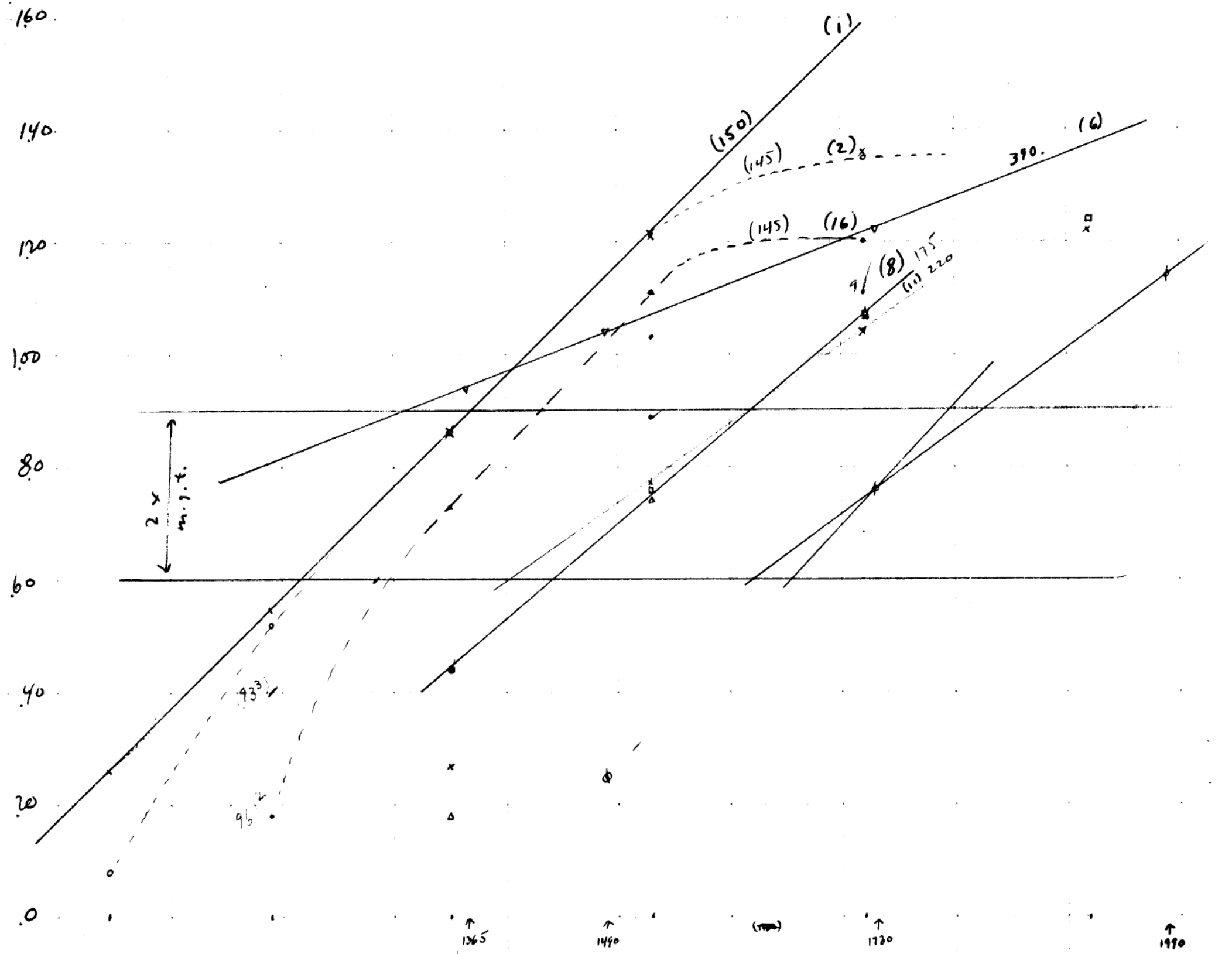
should have separate tubes.

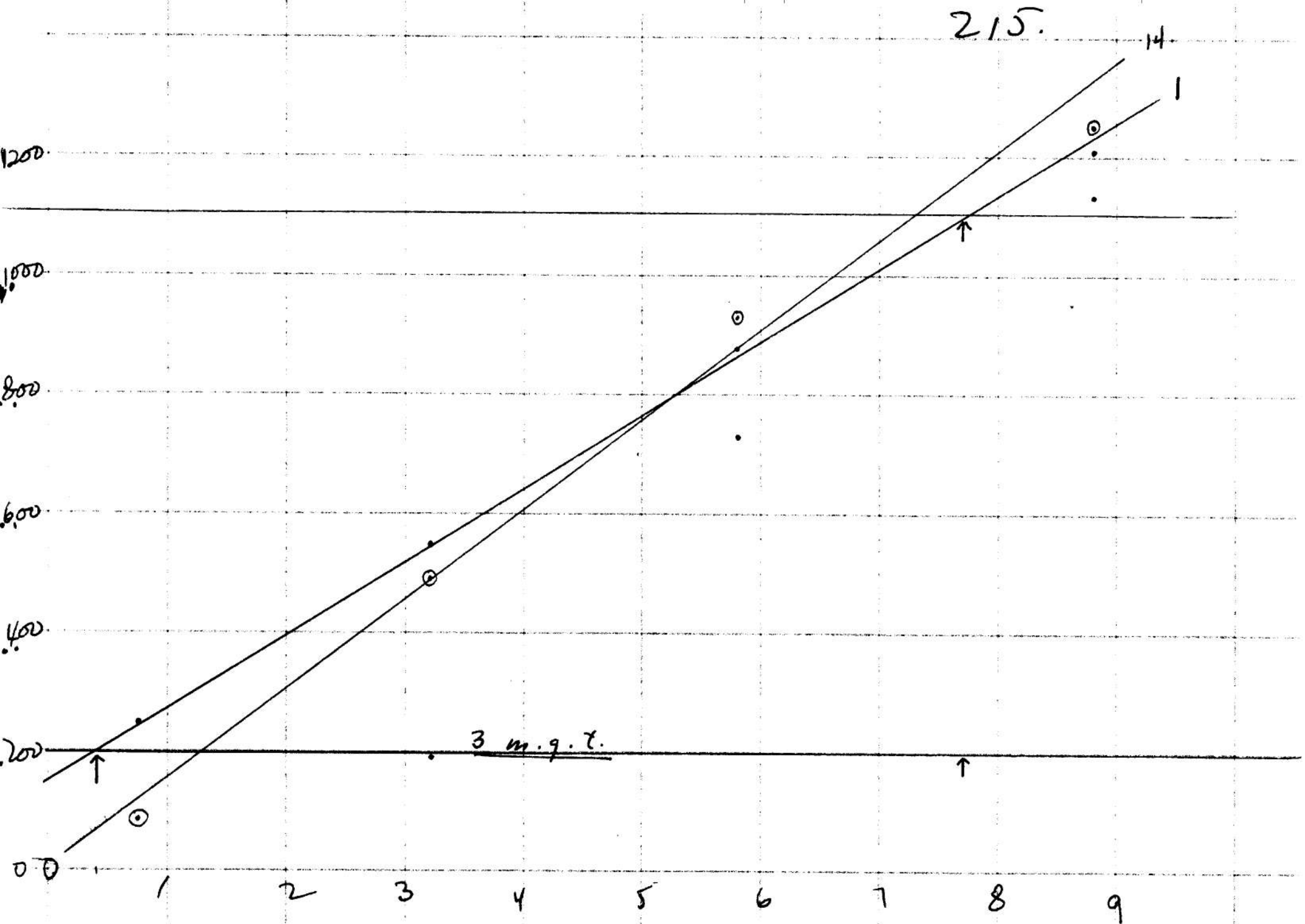
$$d = 2 - \log G$$

$$t = k \log d.$$

$$t = k - \log(\log G/100)$$

$$= k - \log \log G/100$$





B $\frac{1}{2}$ mutants.

Proc. 1245A28.

Y-38	Arginine	Leucine	Citrulline	24h.	48h.
1	0			99 ³	
2	1			94 ¹	
3	3			92 ¹	
4	10			83 ²	
5	30			78 ²	
6	100			61	
7	300			57	
8			1500	100	
Y-39		0		100	100
11		1		"	"
12		3		"	"
13		10		"	"
14		30		"	"
15		100		"	"
16		300		"	"
17				(78)	"

good response curve.
 $\frac{1}{2}$ max = ca. 0.03 mg

all up.

Y-39. 21. L-leucine 100 μ
 dl-leucine 100 μ
 dl-isoleucine 100 μ

Proc 438 in coli ϕ 12428. Grow on shaken @ 30°.

Mix after growth together.

Castids are questionable -

syntrophomonis in plate 13 or canyover?

Six: Analysis of 212 cultures.

219-

Mrs Pichler		BPTM	-B	-P	-T	-M
x P.T.?	1 ✓	+	+	+	-	+
?	2 ✓	-	-	-	-	-
BT?	3					
BT?	4 ✓	+	-	+	+	+

Analyse 219-2

5/31/46.

1. Inoc 50 ml coli α & β (679-183) and β (58-161) and shake at 30° 11P30 - 1130P31.

r/ = Wash and resuspend in 25 ml H₂O.

Turbidity - P4.

Inocul Cell count:

1.	α	10^{-7}	in α	24
2.	β	"	" " "	53

Back-mutation rates: inoc 1 ml. of suspensions.

3	α	in	0.	0	-
4	α	in	Threonine exc.	0	\pm
5	α	in	proline "	2	surface maybe cut but looks like coli
6	β	in	0	0	-
7	β	in	Biotin	0	-
8	β	in	Methionine	0 (T)	+

Syntrophomonas. Inoc in BT.

9.	β	10^{-7}	Control	1	-
10	α	10^{-7}	"	0	-
11	α	10^{-7}	+ 1 ml β control Control	0	-
12	β	10^{-7}	+ 1 ml α "	0	-
13	α	β	1	0 (T)	++
14	"	"	10^{-3}	1	surrounded by local strain. good \pm
15	"	"	10^{-5}	1	" " " " " " " " \pm
16	"	"	10^{-7}	0	\pm <u>no colonies.</u>
17	10^{-3}	1	0	0	-
18		10^{-3}	0	0	-
19		10^{-5}	0	0	-
20		10^{-7}	0	0	-
21	10^{-5}	1	0	0	num. v. sm. col.
22		10^{-3}	0	0	-
23		10^{-5}	0	0	-
24		10^{-7}	0	0	-
25	10^{-7}	1	0	0	num. v. sm. col.
26		10^{-3}	0	0	-
27		10^{-5}	0	0	-
28		10^{-7}	0	0	-

(2)

→ The conclusions that may be drawn from this are limited:
in general diffusion must be limited by the agar to the point
where syntrophism is less effective. It is not due to allosteric
together confirm by plating ^{delete} wild type into concentrated mutant.
Does not adaptation serve the same purpose??

As above but expose by removing transite shutter after 5 min. warm-up period. 5" from tube. Plate initially at $1:10^7$, then take 1 ml samples and determine time for pS of ca. 7.

5:15 P31. 1 min. intervals for 10 min.

t = 0 10^{-7} 65 650,000,000 pS

1	1	} ml	turbid	
2	1		turbid	
3	1			10 ⁵
4	1			ca 10000
5	1			ca 10000 1200
6	.1	} ml	ca 40 40.	
7	.1			7
8	.1			0
9	.1			0
10	.1			0
15	1 ml		0	

→ 6

Because of clumping after ~~exposure~~, the counts at lower mortality (?) maybe disregarded + the initial count should be considered to be

6/1/46.

430731. Broc 50ml coli ∞ and shake at 30° - Y38.
4. 4P1

1. Dil 1:10⁷ into coli ∞ . ca 2000.

Shake
on shaker

irradiate 5 mins as exp. 221. Broc 5ml into 50ml coli ∞ and incubate on shaker. (A). Centrifuge + wash 25 ml + resuspend in saline. Inoculate 0.1 ml into T (Arginine) by detection technique, \bar{c} cover.

Immediately after irradiation:
0.1 ml = ca 2000 colonies.

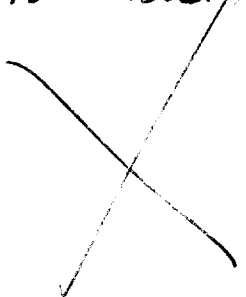
100 mean.

- 1
- 2
- 3
- 4
- 5
- 6
- 7
- 8
- 9
- 10.

$\rho S_{5min.} = \underline{\underline{6.}}$

Dil. A 1:10⁻⁷ and more reabrod.

- 11
- 12
- 13
- 14
- 15
- 16
- 17
- 18
- 19
- 20.



opt. arginine = 3000 dl - .

Phage T-1.

223.

1/2/46.

labelled titre 2×10^{10}

Dilute T-1 as received from Demerec 10^{-9} and plate \approx ca

10^9 cells of:

740P2

1. B/2

Uniform turbidity

2. Y38 (Bh) Agminocless.

Uniform turbidity

3) Plate ca 10^8 Y38 \approx 10^{10} T-1 in colico. for Y38/T1. \rightarrow Large clear center \approx
4) Turbid! ca 400 secondary colonies; puri-
pheny turbid.

5. 10^8 Y38 in colico. When solid, streaks

6. 10^9 B/2 in colico. T-1 over surface.

} no response!

~~12~~ Liquid culture -

4P4 - Inoc colico \approx Y38 shake at 37° .

ca. 10^8 /ml.

1016P - Add 1cc of Phage T-1 " 2×10^{10} ".

12M - no change

9A5 - no change.

broc 50ml coli & flaskes + shake @ 30°. 11P2

6/2/46.

Shaker broken down A3.

1. 58-161

2. 679-183.

3. 58-161 + 679-183.

Leubrid; ca 16 colonies.

11P4 - Centrifuge + wash. broc into T(0) = 2cc. culture.
 pu plate & incubate. also 208-51-2-3. → 0, 0, 0.

As above. No shaking 1130P5. - 1230A7. plate out on T(0).

1158-161

12 679-183

13 } grown
 14 } together

15 } grown
 16 } separately

p7
 0
 0
 4
 5
 0
 0

! Fusion or sexuality? 2

P7. P⁶ colonies from ^⑤ ~~the~~ ~~Flask~~ coli d.; p8 to dants. 1-6.

See: 22?

P10 - numerous additional colonies appear in 13+14
 definite halos around colonies for a diam 2-4mm.

6/5/46.

Inoc. 5 flasks coli $\bar{c} = 438$. Grow shakers 37° 4P41030P 5:Inoculate individually ⁵ min. in same quantity tube & determine killing.

1. $t = 0.10^{-7}$
ca 100.

$t = 5$ Inoc ¹ ~~ml.~~
ca 19,000

2

3

4

~~5~~Killing pS at 6 mins. can be taken as: $\log: 10^4/10^9 = 5$

Y38 - mutants

6/6/46.

1130 P5 from same coli as Y38 Shakes at 37°.

Immediate 6 min. Wash by centr., dilute 1:100 + plate in T (arg.) for mutant detection: ~~200-17~~ 247

- 1
- 2
- 3
- 4
- 5
- 6
- 7
- 8
- 9
- 10

Too heavy !! * * ⊗
 ca 10^{4-5}

Inoc 1 ml into F(10) and cultivate at 37°.

6/9/46. 9 JUN 1946

1A9 - Plate cultures 1-4 of 224 (1-6) at 10^{-6} and 10^{-7} into T(0).

- 1a
- b
- 2a
- b
- 3a
- b.

1A10 - numerous colonies on plates per repetition. Pick sample colonies from surface each plate ~~and~~ to H₂O + to slant colia + test on minimal

Plate:		T(0).	T(10).
1.	1	++	
	2	"	
	3	"	
2	11	"	
	12	"	
	13	"	
3	21	"	
	22	"	
	23	"	
4	31	"	
	32	"	
	33	-	→

#

4
31

see 231.

A10. from oslant to tube of ∞. Wash. Use in PM for plating.

19 JUN 1945

1. Isolated colonies from plate 223-3 to colic. Transfer to slants.

2.
3.
4.
E.

11. Pick agar between colonies from plate 223-3 to the surface of a colic plate streaked with B/r. No lysis. Probably agar too old.

12. hoc colic tube \bar{c} B/r heavily \bar{c} phage \bar{c} !!
10 P. hoc \bar{c} ~~agars~~ ^{T-1 (D)} by 1130 - considerable clearing?

13. " agar plate as above \bar{c} T-1 (D). \rightarrow lytic zone.

2A 13A. Y38 + T-1. \rightarrow lytic zone.

Use Y38 agar for colic in this exp.

9. 14. Streak agar plate of B/r \bar{c} 12. distinct lytic zone.

15. Streak agar plate of Y38 \bar{c} 12 lytic zone (?); later somewhat turbid

all streaks gave clear areas:

\therefore Phage \bar{c} is effective on colic B/r + Y38; also Y38 is sus. to T-1.

21. 11A9. hoc 50ml colic \bar{c} T-1 "10⁰" + 5ml 10hoc \bar{c} colic B/r from colic incubate at 37 in shaker. 1230 P9. - largely cleared.

22. hoc ~~50~~ 50ml \bar{c} 21-1ml + B/r 5ml. 1230 P9. incubate at R.T. mostly. 11P9. - cleared (fairly residual turbidity).
Control completely turbid! Use for stocks T-1. See 230

23. plaque out T-1 (D) on B/r. 10⁻³ to 10⁻⁹
- results obscured by vigorous fermentation + bubble formation.
No plaques found at 10⁻⁴, 10⁻⁶, 10⁻⁷
Numerous at 10⁻⁵ (ca 10³).
Innumerable at 10⁻³. Should use medium with less glucose for phage enumeration?
Gas bubbles in the agar are cones ~~or~~ or lenticulate.

Try Theonine. $\bar{\epsilon}$ recovery added Theonine in view of

very high requirement.

9 JUN 1946,

As 202. K-12 in 50ml T() + suppl. Shake at 37°. Inc 145A9.

	6 (1:10) 84 ¹	d. .744	mg/cc 1.74	Hydrolysate =	Recovery %
1. 0				85 ¹ .693	93%
2. glut. 5mg	81 ³	.875	1.87	foray. (743) 83 ² .783	89%
3. anthr. 5mg	83 ²	.783	1.78		
4. putolact 1mg β-alanine 1mg.	84	.757	1.76		
5. citrulline 5mg.	83 ² (73)	.783		(78 ²)	

See 234.

Harvest 1130P9. Separate 25 ml samples. (a). 16 = 15cc.

Start hydrolysis in 6N HCl 20cc but discontinued due to severe bumping. —
for 1, 2, 5 only. Estimate recovery from volume ←

	Volume after hydr.	%	to	Volume
1.	4.5cc 11.2cc	45%	to	13.5cc
2.	4.5cc	22.5%		5.6cc
5.	6cc.	30%		7.5cc

Neutralize. Dilute to % of 25 cc ind. by recovery
1cc of each solution should give mutant response of magnitude indicated
above, in 10 ml medium. See 234. for assays

Glutamic acid does not increase production of proline in wild type
Citrulline (proportion) of arginine

Note Y38 is a B strain and therefore >100% recovery does not
signify anything.

Requirement is α content. (proline α = c. 3% dry wt.).

6/10/46.

Slide-filter 228-22 for T-1 stocks. Pool \bar{c} 228-21

Suspense aliquots variously; some to K.W. for lyophil preservation.

Plaque out on β 12 at indicated dilutions. 1A18 for filter.

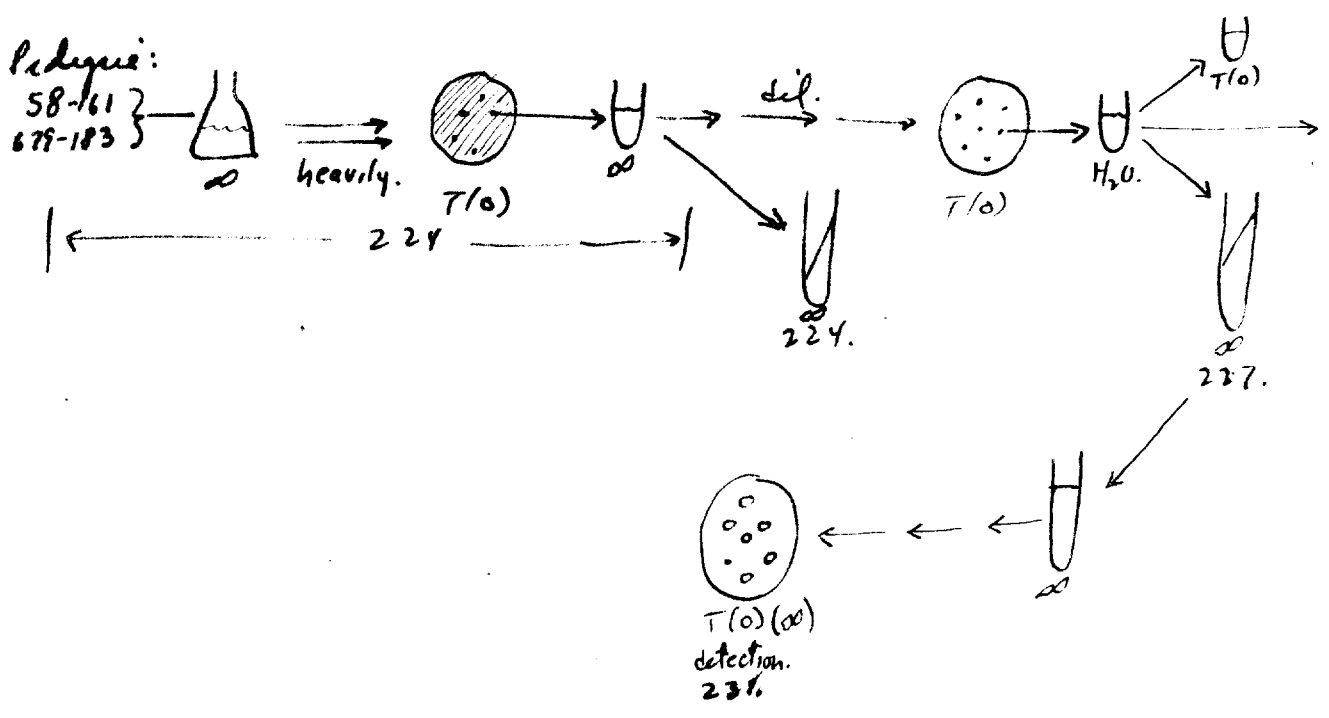
1. 10^6 ca 200
2. 10^8 1
3. 10^{10} 0
4. 10^{12} 0

2/11/46.

Plate out 227-1, etc. detection plates, for mutant segregants.

dil 10^{-6} , 10^{-7} . Unshaken culture in coli from slant.
 1A11. Layer E coli 12N12.

- 227-1
- 2 1
- 3 11
- 4 11
- 5 21
- 6 21
- 7 31.
- 8 31.



1230A13 - Notting signi. seen.
 11A13 - do. (col 0?)

Y38 Mutants.

232

6/12/46.

Knoc coli ∞ 40ml \bar{c} Y38 + shake at 30° 11 P 11

Leibin ride

6/11/46.

Wash and plate heavily into T(10) etc. The cultures:

from P5 coli 02

heavily = 1ml grown culture / plate.

1. 58-161

2. 679-183.

3. 58-161 + 679-183.

SP11 ↓ (6 day cultures).

				1230A13	11A13
1.	1	in	0	0	
2.	1	m	0	0	
3.	2	in	0	0	"
4.	2	in	0	0	
5.	3	m	0	0	
6.	3	m	0	0	
7.	1+2	m	0	0	
8.	1+2	m	0	0	"
9.	1	m	broth	0	
10.	1	m	meth	0	
11.	2	m	threonine	0	
12.	2	m	proline.	0	
13.	3	in	BP	0	"
14.	3	"	BT	0	
15.	3		MP	0	
16.	3		MT.	0	

OK —
throw out.