11/16/50

W2252 = 1895 Hfr lp^s

(-) homodiploid with HFT) ; ~~to~~ obtain Hfr gal-

2341 { 1. Several (-) colonies with N16 HFT gal_2^-
Isolated and purified - HFT λ tests indicate gal_2^-
cross with W902

Results indicate HFT cross - almost solid
smear (with dilute cells also)
no (+) in probably 100,000 colonies

(2.) Streaked against λ - not sens.
518 - not lysogenic = lp_1^r lp_2^r
λ2 - sensitive

(3) "Lysate" prepared - very viscous

0.1 ml / 2281 = no effect, neither plaques nor ~~any~~ opacity
" 518 = " " " (?) 14/9 = ~~no~~ no obvious control area

11/25/50 Repeat with NA-4 lysate 1 & 2

W 2345 = From lysate 1 - one gal_1^- isolated - { tested against N16, NA-1 = gal_1^-

From lysate 2 - 5 gal_1^- isolated - { " " " " = gal_1^-

also nonlysogenic

NA-1
NA-2 } lysates
sterility confirmed

11/30/53

Transductions for segregant analyses

A W1765 gal⁻ (made by ~ transduction) W2373

	λ	control %	plage ½
1.	811	1	30
2.	1210	0	56
3	902 (Amer lysate)	1	11
4	K-12	1	33

W 750

1.	K12	1	46

W1210

1.	K-12	4	21 (23)
2.	750	2	65
3.	811	5	56

12/5/53 See also next page

T18} λs against 2070 - to see if 60% r⁻, 2⁻ is in these apparent doubles.
T19}

(-2) lysates not very active

245B

T18-2 5/11
T19-2 9/2

1. Analysis of transductants of 2070 by T19-2 λ

	segregants /N1	/N16	Focus
1.	+	0	2⁻
2.	+	0	2⁻
3.	0	0	-- ?
4.	+	0	2⁻
5.	+	0	2⁻
6	+	0	2⁻
7	+	0	2⁻
8.	+	0	2⁻
9.	+	0	2⁻
10.	+	0	2⁻
11.	+	0	2⁻
12.	+	0	2⁻
13.	+	0	2⁻
14.	+	0	2⁻
15.	+	0	2⁻
16.	+	0	2⁻

1. Analysis of segregants from 2 transd. of T18/2070

	/N16	/N1	λ Rx	Allele
1.	+	0	R	1⁻
2.	+	0	"	1⁻
3.	0	+	"	2⁻
4.	0	+	"	2⁻
5.	+	0	"	1⁻
6.	+	0	"	1⁻
7.	+	0	"	1⁻
8.	contaminated (+)		"	

12/2/53

A. $\pi\delta^-$ transduced by mutant N16, N1, but not by single λ's

Proposed Constitution: ① 18 pap (gal+) picked and streaked. Many unpure colonies — all appear to be segregating gal+ 2^-1^-, 2^-1^+, 2^+1^- } 3:1 on low should be low slow if gal− present — which it was. Can't test gal segregation

② One of +u on gal streaked out on low, gal — segregating on gal — also on low, (+) appears to be segregating low slow

	on low	Gal+ Segregants	by transd
bio 2^-1^-	s	no	wt.
bio 2^+1^-	s	no	1^-
bio 2^-1^+	+	no	2^-
bio 2^+_2 1^+_1	+	seg low slow	2^-, wt.
bio 2^+_2 1^+_1	s	no	1^-, wt.

See next page 246A for details

B. 2345 / 2251 λ to make HFT, Hfr

0.1 ml 2251 λ = 56, control γ_2 = 0

One second streaking only 4/24 +u — greater stability of transd. Than expected

The 4 restbed and segregants examined

12/9/53 T18 continued.

Transcr. Assay

1. The loc

	Loc	Bond	N1	N116	Loss	
# 1	slow	(−)	o	+	1−	
2	slow	"	o	+	1−	
3	slow	"	o	o	double −	} On standing (12/14) Trace cell pigmentation quite heavily.
4	slow	"	o	o	"	
5	"	"	o	+	1−	
6	"	"	o	o	double −	?
7	"	"	o	o	"	
8	"	"	o	+	1−	
					41−	
					4 (+−)	

Low chloraller not good culture

Ignore

accordance with results [illegible]

Trans. Assay

2. The Cac + ON

	Bloc	Bond	Sqzt	HFT N1	HFT N116	Loss	
# 1	+ (?)	+"	+	o	+	1−	−1 ⊕
2	clay	"	o	o	o (1pp)	−−	−2
3	clay	"	o	o	o streakthrough	−−	−
4	"	"	+	o	+	1−	−1 ⊕
5	"	"	o	o	o	−−	−2
6	"	"	o	+	o	2−	−2 ⊕
7	"	"	+	o	+	1−	−1 ⊕
8	"	"	o	+	o	2−	−2 ⊕
9	"	"	o	o	o	−−	−1 +[1] gave stable + early
10	"	"	o	+	o	2−	−2 ⊕
11	"	"	o	o	o	−−	−11
12	"	"	o	o	o	−−	−2 +[2] gave stable + early
13	"	"	o	o	o	−−	−2
14	"	"	o	+	o	2−	−2 ⊕
15	"	"	+ 1pp	o	+	1−	−1 ⊕ Rev. unstable (det 12 sqzt (−−)
16	"	"	o	o	o	−−	−11
17	"	"	o	o	o	−−	−2
18	"	"	+ 1pp	o	o	? −−	−2
19	"	"	+	o	+	1−	−1 ⊕
20	"	"	o	+	o	2−	−2 ⊕
21	"	"	o	o	o	−−	−2
22	"	"	o	o	o	−−	−2
23	"	"	+	o	+	1−	−2 ⊕
24	"	"	o	o	o	−−	−24

24 loc all picked from here and Sqzt tested

Certain behaviors not likely from loc — need better basis of det. suggest this is [illegible]

Low recession on double ?

[1] gave stable + early

[2] gave stable + early

On 2 examinations the following marked ⊕ were found stable — Why stable?

Omit known — 13 −−
What to — 6 1−
Think about — 5 2−
This.

Correlation of duction above with 1° clap This lysate is not + [illegible] 1pp but I'm not This is a [illegible] 1pp lysate

3/5/54 Lysate made of 246A-15

1 drop		
/200	solid spot	medical HFT gel
/120	0	
/80	solid spot	

6/18/54
246A-15 + streak discarded
246-A-15 on streak

12/6 KS3

W/1210 transduced by K12

Transclone	Transduction 578	Segregant N16	N1	Segregant Rx	Mdel
1.	typ	0	+	r	2-
2.	"	0	+		2-
3.	"	0	+		2-
4.	"	0	+		2-
5.	"	0	+		2-
6.	"	0	+		2-
7.	" pure gal+				
8.	" pure gal+				
9.	"	0	+	r	2-
10.	"	0	+		2-
11.	"		+		2-
12.	"		+		2-
13.	" weak	0	+		2-
14.	"	0	+		2-
15.	" pure gal+				
16.	" pure gal+				
17.	"	0	+	r	2-
18.	" weak pure gal+				
19.	" pure gal+				
20.	" gal+ sc				
21.	" pure gal+				
22.	"	0	+	r	2-
23.	"	0	+		2-
	18/23 +"		15/15 gal-		

See p9 — 21 + 2 = 23 — 6/22 ++

5 R
4- +
2+ -

(247A)

5 R
4* +
2- - → 5 R
4- +
2- +

6/18/54 Stocks discarded
See

Cross x 902		
247A - 4	6/21/54	out
247A - 13	Pure w. streaks	5/13/55
247A - 22	"	"
247A - 33		

Lysates 4/22/54

	Spotted /2.75	/ml	
4	1/<0.1	2/0.5	These two lysates give large numbers of small papillae — possibly not stable? partially HFT?
13	0/<0.1	0.70/0.5	
22	0/<0.1	0.20/0.5	
23	0/<0.1	0.40/0.5	

12/6/53

W1210 & 811

See 24½ for origin

(24½B)

Transd.	Transd. /STR			
1	lys.	NR +	STR 0	
2	"	0	+	
3	"	0	+	
4	"	0	+	
5	"	0	+	
6	"	contaminated with gelt →		
7	"	0	+	
8	"	0	+	
9	"	0	+	
10	"	0	+	
11	"	0	+	
12	"	0	+	
13	"	contaminated with gelt →		
14	" rx	0	+	
15	" bad	0	+	
16	"	0	+	
17	lys	0	+	
18	"	0	+	
19	"	0	+	
20	"	0	+	
21	"	0	+	
22	"	0	+	
23	"	0	+	
24	"	0	+	
	aur +	22 gal$_2^-$		
		1 gal$_4^-$		

Sep Rx r ← 9/10 (54?) NFT segregant found gal$_4^-$ + of NFT segregant found attl.

Cross x 1176, 90v

1
8
14
18

6/18/54 Stocks discarded

2/9/54

Lysalb /SM /2175

-1 0/0 solid smear = NFT gal$_4^-$ (ALSO as the result F$^+$ — x F^x)

See 245

12/6/53
WI-10 t 750

Transd. #	Transd. typ Rx/98	Seg N16 / N18	Seg N1 / N4	Seg typ Rx	Loci
1	typ	0	+	✓	2-
2	"	0	+	✓	2-
3	"	contaminated w/ gal →		✓	2-
4	"	0	+		2-
5	"	0	+		2-
6	"	0	+		2-
7	"	0	+		2-
8	"	0	+		2-
9	"	0	+		2-
10	"	0	+		2-
11	"	0	+		2-
12	"	0	+		2-
13	"	0	+		2-
14	"	contaminated →			(out seg typ disag)
15	"	0	+	✓	2-
16	"	contaminated with gal →			2-
17	"	0	+		2-
18	"	0	+		2-
19	"	0	+		2-
20	"	0	+		2-
21	"	0	+		2-
22	"	+	0		1-
23	"	contaminated with gal ↓			—
24	"	+	0		1-
	acc + a	19 gal 2-			
		2 gal 1-			

6/10/54

		out
Trans on	1	5/13/55
	3	11
other strands	24	11

(24F)

12/5/53

4750 transduced by K12λ

Transd #	Transd 578	/N1	/N16	Alkaline
1	Lyo.	0	+	1-
2	"	pure galt →		
3	"	" " →		
4	"	" " →		
5	"	contaminated with galt →		
6	"	0	+	1-
7	"	0	+	1-
8	"	0	+	1-
9	"	0	+	1-
10	"	0	+	1-
11	"	0	+	1-
12	"	0	+	1-
13	"	0	+	1-
14	"	0	+	1-
15	"	pure galt →		
16	"	0	+	1-
17	"	0	+	1-
18	"	0	+	1-
19	"	0	+	1-
20	"	contaminated with galt →		
21	"	pure galt →		
22	"	0	+	1-
23	"	0	+	1-
24	"	0	+	1-
	19/24 +ᵃ			16 1-

lysate	Spotted 2175	750
1	1/0.5'	0/0
7	2/0.5'	0/0
9	2/0.5'	0/0
11	5/0.5'	0/0

6/20/54

← Only cell-free supernatant from stock 3 was used.

12/8/53 — W2373

(249) A

W1765 gal$_1^-$ (by c-s transd. with NAA lysate) transduced by 841 - See p. 245

Transd.		Stab optos	Sen	
1	— S ✕ arn lys	stable	N1	S18
2	R ✕ lys —	"		
3	R ✕ lys —	"		
4	R ✕ lys —	"		
5	R ✕ lys —	"		
6	R ✕ lys —	unstable	+	+ (?) upper λ^R = gal$_1^-$
7	— S ✕ arn lys	stable	+	
8	R ✕ lys —	"		
9	R ✕ " —	"		
10	— S ✕ arn lys	"		
11	R ✕ lys —	"		
12	R ✕ " —	"		
13	R ✕ " —	"		
14	R ✕ " —	"		
15	R ✕ " —	"		
16	R ✕ " —	"		
17	R ✕ " —	"		
18	R ✕ " —	"		
19 —	skipped	"		
20	R S arn lys —	"		
21	R S " —	"		
22	— S + arn lys	"		
23	— S "	stable		

etha? → stable

Lp^+ = 17 — $\frac{1}{23^+}$

Lp^{++} = 5 — 1 Lp^+

5 s
17 +
0 r

} Ch/ST
streaks discarded

23	22
24	16
12	12
23	13
82	43
23	23
	40
	82

0.5°
59 | 41.0
35 4
560

41
59

12/8/53 W2373

W1765 gal - (by (-) transd ē NA4) transd. by 1210 /

Transd #	Transd Lp R	Transd. stability	Sig-direct N1	Sig-direct N116	Wild
1	Lp S	S			
2	H	U	−	+	λ^R 1⁻
3	H	S			
4	H	S			
5	non lys	S			
6	lys	U			
7	non lys	S	what give ?		
8	lys	U	−	+	λ^R 1⁻
9	H	U	contaminated ē	(+)	
10	non lys	S	+	−	λ^R 2⁻
11	H	S			
12	lys	U			
13	lys	U	−	+	λ^R 1⁻
14	non lys	S			
15	lys	U			
16	H	S			
17	H	S			
18	H	S			
19	non lys	S			
20	lys	U	−	+	λ^R 1⁻
21	H	U	−	+	λ^R 1⁻
22	H	S			
23	H	S			
24	H	U	−	+	λ^R 1⁻
		8/24 Lp^+			6 1⁻
	Lp^+ = 18				
	Lp^{S^r} = 6 = 3 Lp^S	8 Lp^+			1 2⁻

an Lp^R transductin segregating the allelic allele and Lp^R simultaneously.

11/24/54 Probably an error in recording

S may segregate to segregating and wild stable.

see next page as well.

Lysis 249B
10
13
20
21

6/18/54 Strains discard

12/10/53 #12373

1765 gal - transduced by 902

Transd #	Transd. Hp/68	Transd. Strength	Segregant	Segregant	/518
1	- Hp S	S			
2	- "	R			
3	- nm Hp	R			
4	- nm Hp	S			
5	- Hp	R			
6	- Hp S	R			
7	- nm Hp	S			
8	- nm Hp	S			
9	- Hp S	R			
10	- Hp	R			
11	- nm Hp	R			
12	- Hp	R	u	—	+

7 Hp R
2 Hp R S
3 Hp S

6/18/54

(Strain discarded)

Pure in the only segregant obtained

/R 1-

12/10/53 W2373

1765

Wild-75 Sal. transduced by K12

Sig Samples: 23

Transd. #	Lys	λ	Transd. Strs	N16	N1	Allele
1	lys	r	s			
2	lys	r	s			
3	lys	r	s			
4	non lys	s	u	+	-	λ^R I⁻ Lp⁻
5	lys	r	s			
6	non lys	s	u	contam. C (+)		
7	lys	r	s			
8	"	"	u	+	—	λ^R I⁻
9	"	"	s			
10	"	"	s			
11	"	"	s			
12	"	"	u	+	—	λ^R I⁻
13	"	"	s			
14	"	"	u	+	—	λ^R I⁻
15	"	"	u	+	—	λ^R I⁻
16	non lys	s	s			
17	lys	r	u	+	—	λ^R I⁻
18	"	"	s			
19	"	"	s			
20	"	"	u	+	—	λ^R I⁻
21	"	"	u	+	—	λ^R I⁻
22	"	"	u	+	—	λ^R I⁻
23	"	"	s			
Ttl			10/23 +u			

20 Lp⁺ 3 Lp⁻ | 8 Lp⁺ 2 Lp⁻ | 9 Lp⁻

Lysis 4, 8, 17, 14 } were here any others on 1st prep.

{ 6/10/54 Strs clear discarded }

Exp. 642

250
A

F+ F⁻ Lp+ Lp⁻

12/12/53 EMLs cross 1210 × 2234 Gal₂⁻ × Gal₄⁻

1. The small colony "pinkish tinge" (probably requires B)

	M16	S1+	Lp Rx	Locus	
1.	+ (few)	0	s weak	4	
2.	+	0	s	4	
3.	+	0	s	4	
4.	+	0	s weak	4	
5	+	0	s weak ?	4	
6.	0	+	R	2	—
7.	+	0	s	4	
8	+	0	s weak	4	
9.	+	0	s	4	
10	0	+ R	s	2	—
11.	+	0	s weak	4	
12	+	0	s	4	
13	+	0	s	4	
14.	+	0	s	4	
15.	+	0	s	4	
16	+ R	0	s weak	4	
17	0 2	+	s weak	2	—
18	+	0	s weak	4	
19	+	0	s	4	
20	+	0	s	4	
21	+	0	s	4	
22	+	0 (?)	s	4	
23	+	0	s	4	
24	+	0	s	4	
25	+	0	s	4	
26.	0 v	0 (414)	v	?	
27.	0	+	v	2	—
28.	0 v	0 v	s	?	
29.	+	0	s	4	

23 = 4⁻ all Lp⁻
4 = 2⁻ 3 Lp⁺ 1 Lp⁻
2 = 2⁻ 4 Lp⁻

F+ Gal₂⁻ Gal₄⁺ Lp+ × F⁻ Gal₂⁺ Gal₄⁻ Lp⁻

Cont. of 250A.

2. The Large colony puto royale.

	N16	S1t	Lp	Locus
1.	0 ✓	0 ✓	S	— ?
2.	0	+	r	2 —
3.	+	0	S	4
4.	0	+	r	2 —
5.	+	0	S	4
6.	+	0	S	4
7.	+	0	S	4
8.	0	+	r	2 —
9.	0	+	r	2 —
10.	+	0	S	4
11.	+	0	S	4
12.	0	+	r	2 —
13.	+	0	S	4
14.	+	0	S	4
15.	+	0	S	4
16.	0	+	S	4 —
17.	0	+	r	2 —
18.	+	0	S	4
19.	+	0	S	4
20.	+	0	S	4 ✓
21.	0	+	r	2 —
22.	0	+	r	2 —
23.	+	0	S	4
24.	+	0	S	4
25.	0	+	r	2 —
26.	+	0	S	4
27.	+	0	S	4
28.	+	0	S	4
29.	0	+	r	2 —
30.	+	0	S	4
31.	+	0	S	4

20 = 4⁻ all Lp^S

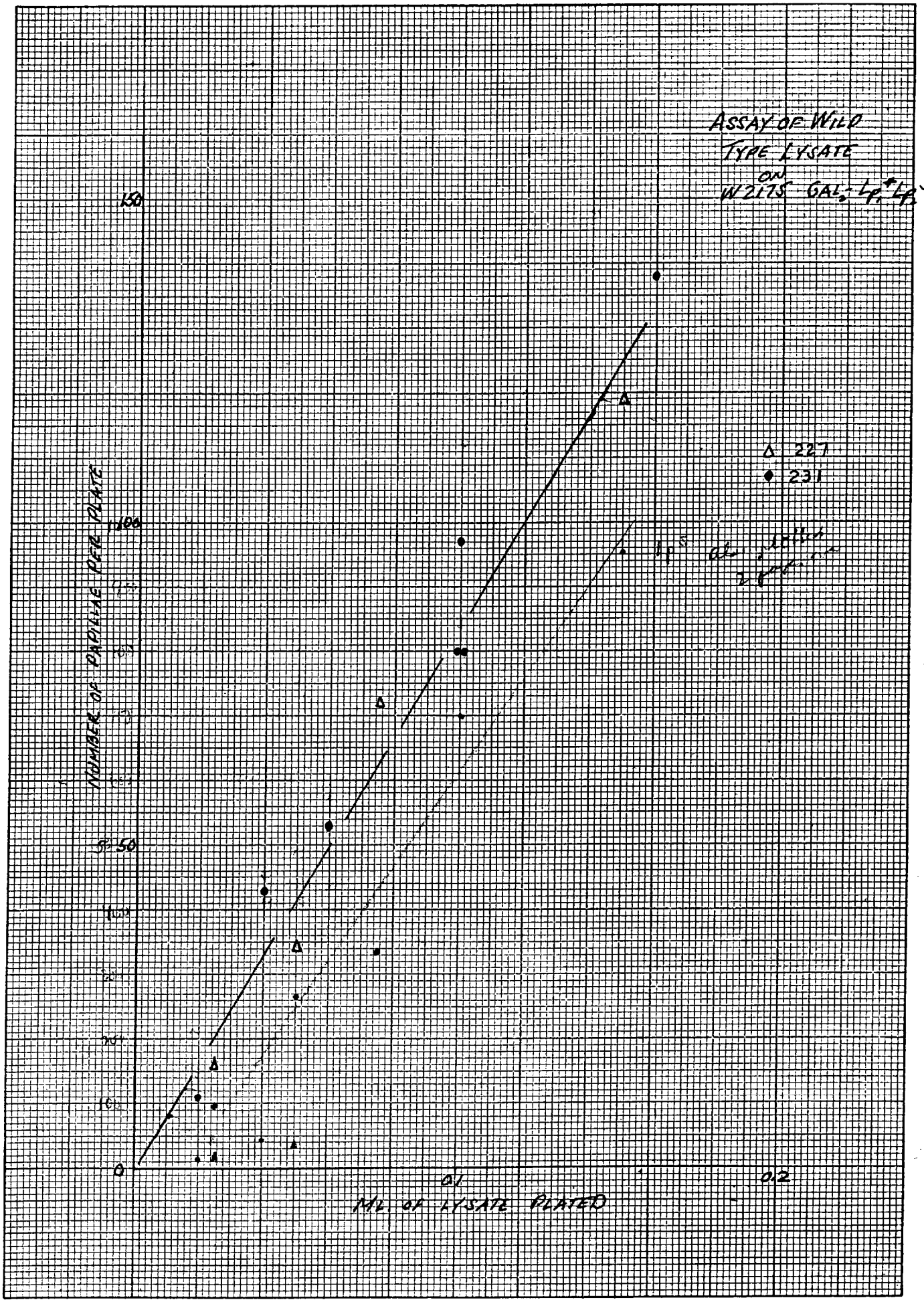
10 = 2⁻ all Lp^r

1 = 2⁻4⁻ Lp^S

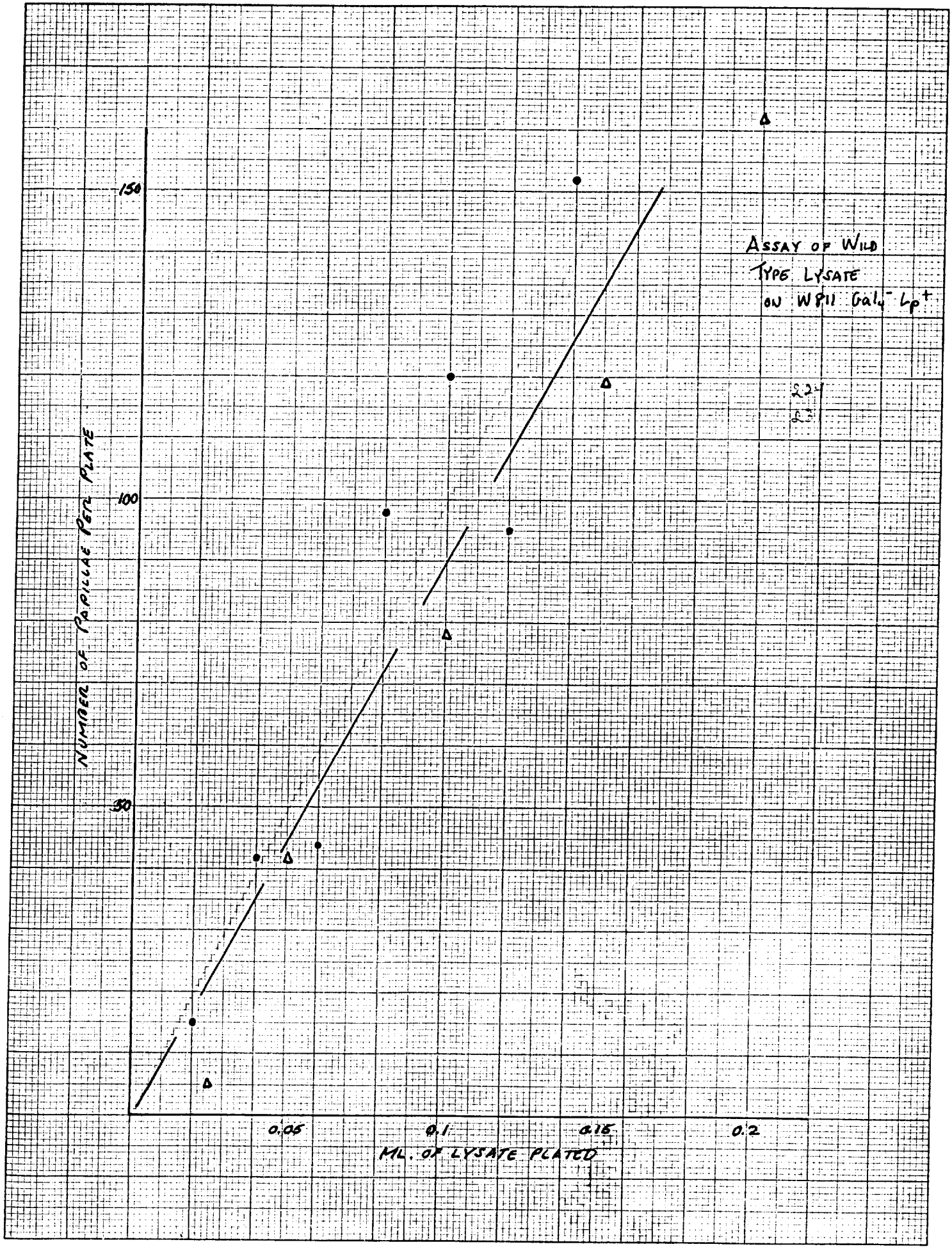
4⁻ Lp^V X 2⁻ Lp^r

Total of 250A + 250B

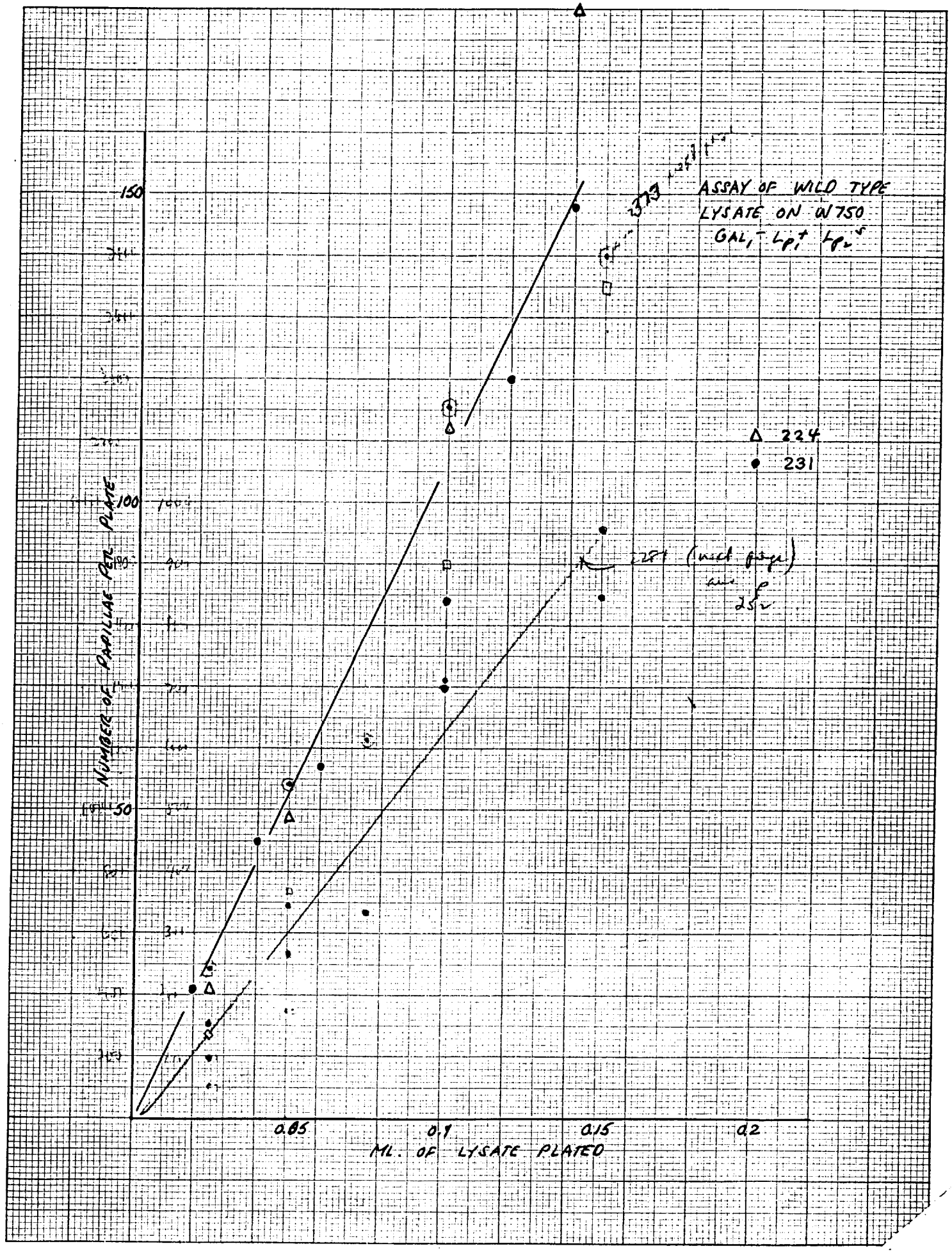
43	4⁻	Lp^V
13	2⁻	Lp^r
1	2⁻	Lp^V
3	2⁻4⁻	Lp^S



Form No. 999 — 20 Squares to Inch
AMERICAN PAD & PAPER CO., HOLYOKE, MASS.



CODEX BOOK COMPANY, INC. NORWOOD, MASSACHUSETTS. PRINTED IN U.S.A.



CODEX BOOK COMPANY, INC. NORWOOD, MASSACHUSETTS. PRINTED IN U.S.A.

Assay - KH2 5/30/52 3.6x10^10 λ/mc

#2175K

ML	227	231
0.025	16	0.02 11
0.05	34	0.04 43
0.075	72	0.06 53
0.100	—	0.1 97
0.15	119	0.16 138
0.20	—	
0.3	255	

Lpk

750

224	231
0.025 - 21	0.02 - 21
0.05 - 59	0.04 - 45
0.1 - 112	0.06 - 57
0.15 - 183	0.08 - —
	0.10 - 84
	0.12 - 120
	0.14 - 148

84

2500

P 253

KH2 12/28

224	231
0.025 5	0.02 - 15
0.05 42	0.04 - 42
0.1 72	0.06 - 44
0.15 119	0.08 - 61
0.2 163	0.1 - 120
	0.12 - 95
	0.14 - 152

0.025 = 5
0.05 = 9
0.1 = 2
0.15 = 2

Exp -

2251

Lp 5

KH2 12/28/53 1.7x10^10 λ/mc

0	0
0.025 - 96	151
0.05 - 266	345
0.075 - 334	—
0.1 - 700	706
0.15 - 955	892

2377

Lp 5

252

KH2 12/28/53 1.7x10^10 λ/mc

0.025 -	483
0.05 -	1079
0.075 -	1225
0.1 -	2314
0.15 -	2799

578

P. 253

KH2 12/28/53 1.7x10^10 λ/mc

0	0
.025	276
.05	738
0.1	1778
0.15	2689

1924

P. 253

KH2 12/28/53 1.7x10^10/mc

0	0
.025	53
.05	72
0.1	143
0.15	249

1.7
1.5
8.5
2.5

12/17/53

2345 - Hfr gal,- Use as a allele tester by

1. growth with prototrophic gal+ for several hours and then plating out on B gal

2. Preliminary encouragement.

Tube	Final titer cells/ml	Plated on B gal after 6 hours (no. gal+/ml)
1. 2175 (= gal+)	0.1 ml	12
2. 2345 (- gal+) Hfr	0.1 ml	1
3. 2175 + 2345	0.1 ml each	166

} suggest method may work

12/20/53 2. Cultures apparently need purification in some places

Exp as above.

pur	Parent	λ	0	Gal+ 2341	Gal+ 2345	Gal+	Gal+	Gal+	Gal+	Gal+	Gal+
✓	2341		47	—	—						
	2345		✓	—	—						
	2343		3	(+) > 500 ✓	(0) (3) ✓						
✓	750		26	(+) > 1000 ✓	(0) 14 ✓						
Lp+	1210	2	9	51 107 ?	(+) > 500						
	2175	2	14	(0) 17	✓ (+) > 1120						
Lp+	2281	2	1	(0) 41	(17) 21						
	578	4	11	(+) > 10⁴ ✓	(+) > 250 ✓						

12/29/53

Are lysates "decaying" at room temperature? Probably to repeat massive dilution 578 and the inability to obtain expected 210.5 of transduction of Lp+ as compared to Lp+ says perhaps. This would also argue that transduction of Lp+ is effected by particles different from those effecting Lp+ transduction

NEW lysates of N16, K12 made - designated 12/28 appropriate titers

1. K12 - $1.67 \times 10 \times 10^7 = 1.67 \times 10^{10}$ against 2201
2. N16 - $>50 \times 10 \times 10^7 = >5 \times 10^9$ " "

N1765 gal,- made by (-) hand. cultured with streak back on

W2373

12/31/53

Assay against λ^s - use of K12 λ 12/28/53

1. 2373 - $gal_2^- - Lp^s$ test Conc. - No. cells in suspension

λ added	pfp/plate	Δ
0.	3	0
0.025	486	483
0.05	1082	1079
0.075	1226	1223
0.1	2314	2311
0.15	2799	2796

$= \frac{137 + 110}{2} = 1243 \times 10^7 = 1.2 \times 10^{10}$.

added / plate = 1.2×10^9

2. 2281 - $gal_2^- - Lp^s$ - No cells in suspension - $2145 \times 10^7 = 2.1 \times 10^{10}$

2.1×10^9 added

λ added	pfp/plate
0	0
0.025	96
0.05	266
~~0.075~~	~~344~~
0.1	700
0.15	955

Suspension appears mixed - large and small clones picked and added - tubs against 5/1 - all req λ^+

1/9/53

Purified 7211 - small colony from cell assay. $\frac{221}{200} \int_{211} \times 10 \times 10^7 = = 2.1 \times 10^{10}$

2.1×10^9 added

0	0
0.025	151
0.05	345
0.1	542
0.1	706

1/4/54

Cells from aerated 10× conc. cultures.

Assays – K12 λ 12/24/53

		Δ	
1. 581			
no add.	40	0	
0.025′	94	54	✓
0.05′	138	98	
0.1	247	207	
0.15′	319	279	
2. 1974		Δ	
no add	27	0	
0.025′	80	53	
0.05′	105′	72	
0.1	170	143	
0.15′	276	249	
3. 2341		Δ	
no add.	34	0	
0.025	217	183	
0.05′	423	389	
0.1	945	911	
0.15′	1398	1364	
4. 518		Δ	
no add.	55′	0	
0.025	331	276 ′	
0.05′	793	738	✓
0.1	1833	1778	
0.15′	2744	2689	

1/11/54

518 Transduction of —, with HFT gal- lysate of 1/9/54

① One ml lysate added to one ml of cell suspension – 1/12/54 incubated at 37C for 10 min centrifuged and resuspended in 1.0 ml broth. diluted and plated.

λ exposed cell titer = 708, 889, 794, 889

3280/4 = 820 × 10 × 100 × 100 × 50 = 410 × 10⁷ = 4.1 × 10⁹/ml

λ exposed cell titer 761, 651, 670, 719

2801/4 = 700 × 10 × 100 × 100 × 50 = 3.5 × 10⁹/ml

② Distribution of cell types - column

	gal (–)	gal (+)	gal – λ culture	Total
1. Control (both exposed)	3280	0	0	3280
2. λ exposed	2801	31	~~58~~ 54	2886

31 (–) picked and streaked /518 - all um lys.
/λ - all sensitive

30(+) picked

#	/518	/λ	Segregating?	/518	/λ	Segregating?	/518	/λ	Segregating?	Summary
Lp+ 1.	lys	R	yes	11. um lys	S	no	21. lys Lp+	R	"	~~R~~ S
Lp+ 2.	"	"	"	12 Lp+ lys	R	yes	22. —	— S	a (–) (no)	23 Lp+
Lp+ 3.	"	"	"	13 Lp+ "	"	"	23. clys Lp+	mix	[illegible] yes	3 Lp^R
Lp+ 4.	"	"	"	14 Lp+ "	"	"	24. lys Lp+	"	"	
Lp+ 5.	"	"	"	15 Lp+ "	"	"	25. " Lp+	"	"	
Lp^R 6.	um. lys.	"	"	Lp^R um lys	R	"	26. " Lp+	"	"	
Lp+ 7.	lys	"	"	17 Lp+ lys	"	"	27 um lys Lp+	small pairs	"	
— 8.	um lys.	S	" ?	18 Lp+ "	"	"	28. lys Lp+	R	"	
— 9.	" "	"	no (wk. R)	19 Lp+ "	"	"	29. lys Lp+	"	"	
— 10.	—	— S	a (–)	20 Lp+ "	"	"	30. lys Lp+	"	"	
							31. lys Lp+	"	"	

2543 Lytic λ – Plaqué (#3) of lytic d lysogen previously reported inadvertently

	um lys	no control spread done
1. 750	3	
2. 2281	9 ← all discarded as slow and slow growing	
3. 2373	4	
4. 518	6 4/4 tested stable	2 appear slow (+)
5. 811	89 ← are stable (+)	

Stocks made

2/24/54 Stocks of these discarded except #s 6, 16, 27

Stocks made of these

254-5 = W2866

254-16 = W2867

1/13/54

Crosses - check on Lp fal segregations

A 7. 1765 X 750 on S fal

39 particles - all (-)

B 2. 2281 X 2035

25 (-)

329 (+)

C 3. 902 X 1655

not counted - many small - appear about 50-50

1/22/53

246A-15 Reversions continued

Reversion	Segregating?	Segregant Character	Proposed Genotype
1	yes	(– –)	1 2 / – – → / + +
2	"	(– –)	
3	"	(– –)	
4	"	(– –)	
5	"	(– –)	
6	"	(– –)	
7	? no	stable +	–
8	? yes	mixed ±	
9	? no	stable +	
10	yes	(– –)	1 2 / + – → / – +
11	"	2–	
12	? no	stable +	
13	yes no	"	
14	"	(– –)	a 1-6, 10
15	"	(– –)	
16	"	(– –)	
17	"	(– –)	

2/4/54

S11 t902 (New location of 902 - 3 bottles)

for the purpose of isolating HFT gal$_4^-$

1. mould ½ - 19
2. 0.1 902 λ - 89

2/8/54

750t902 gal$_2^-$ (segregants coming around about a month) suspected of HFT gal$_3^-$ Origins now cloudy - Originally 3 but one lost Both in need of purification

257-1 → #1 Test for HFT

	Col. 1.
1.	750 NFT
2.	NFT
3.	HFT
4.	"
5.	" (?)
6.	"
7.	"
8.	-
9.	NFT
10.	HFT

257-2 →

257-3 →

257-4 →

#2 Tests for HFT

1.	750 HFT
2.	"
3.	"
4.	NFT
5.	HFT
6.	"
7.	"
8.	NFT
9.	HFT
10.	NFT

9/10/54

NFT cultures examined

257t-1 = gal$_2^-$, +^{17} stbc
257-3 = gal$_3^-$, +^{17} stbc

W2869

S18 transduced with N16 λ 1-9-54 for the purpose of establishing that the transduced colonies contain both 4p$^-$, 4p$^+$ cells and gal$^-$ cells (segregate for both?)

Fresh inoculated with S18, 0.2 ml cells + 0.4 ml λ incubated at room temp. 10'
Then diluted 10^6 and plated B gal

Plate	(+)	(−)	(−) nibbled	total
1.	1*	28	2	31
2.	1**	32	0	33
3.	0	39	2	41
4.	0	43	1	44
	2	142	5	149

37 X 10^1 X 10^6 X 3/1 (10) X (10) ? = 1.0 X 10^{10} ?

1 * a colony such ●
2. ** " " " ◎

both streaked - 10 (+) and 10 (−) colonies picked + λ S18

Opt. 10 16"

	1. (+)	S18 Lysog.	(−)	S18 Lysog.
1.	R	"	R	"
2.	"	"	"	"
3.	"	"	"	"
4.	"	"	"	"
5.	"	"	"	"
6.	"	"	"	"
7.	"	"	"	"
8.	"	"	"	"
9.	"	"	"	"
10.	"	"	"	"

257t-5 +^{17} 4p$^+$

	2. (+)	S18 nonlyg.	(−)	S18 sens. nonlyg.
1.	R	"	"	"
2.	"	"	"	"
3.	"	"	"	"
4.	"	"	"	"
5.	"	"	"	"
6.	"	"	"	"
7.	"	"	"	"
8.	"	"	"	"
9.	"	"	"	"
10.	"	"	"	"

257t-6 +^{17} 4p$^-$

W2868

see p. 262

2/13/54

W2350 - double diechan

parent culture streaked out and 10 colonies examined to eliminate possibility of including reversions as one cocci in the inoculum $- + \rightarrow - +$ and therefore transductible Sg. one of the 2 (-) lysates.

1. All 10 colonies NOT transductible by HFT 2⁻, 4⁻

2. Drop fuls of overnite broth placed on B gal.

2350 = 4⁻ 8⁻

Cells — (c.60 pap) HFT 2 λ | (0 pap.) HFT 4 λ | (c. 20 pap.) HFT₂ HFT 4 λ

4⁻	−	+	
8⁻	+	+	
2⁺	+	−	

Are the pap on HFT 2 due to reversions of the 8⁻ donor being transduced? If so they should segregate 4⁻ segregants predominantly and few (−−). Pick up 12 and see —

258-A. Of the 12 - 6 found unstable Segregants.

	HFT 2⁻	HFT 4⁻			
1.	0	+			
2.	0	+			
3.	contaminated with + (0)		0	+	(2⁻) Now 4⁻ 3 (2⁻)
4.	0	+	−	−	(⁻⁻) explain 2 (−−)
5.	0	+	0	+	(2⁻) 1 (4⁻)
6.	0	0	−	−	(−−)
	+	0	+	0	(4⁻)

3/8/54 Double-decarbt

3. Examine HFT 2 + HFT 4 for double ductn.

258-B pap

19 examined — Segregants streaked

	HFT 2⁻	HFT 4⁻	Loci			
1.	0	0	(−−)			
2.	0	0	(−)			
3.	0	+	2⁻			
4.	0	0	(−−)			
5.	0	+	2⁻			
6.	0	+	2⁻			
7.	0	+	2⁻			
8.	0	0	(−−)			
9.	0	+	2⁻	9	0 (+) pap	
10.	+	0	4⁻	10.	46 (+)	1/6 stable
11.	+	0	4⁻	11	48	1/6 stable
12.	0	0	(−−)			
13.	0	+	2⁻	13.	22	1/6 stable
14.	0	+	2⁻	14.	9 (+) pap	1/6 stable
15.	0	0	(−−)			
16.	+	0	4⁻	16	11 (+) pap	5/6 mixed
17.	0	0	(−)			
18.	0	+	2⁻	18	contam's	
19.	0	+	2⁻	19	event (+)	

Reversions statistically + pap. deleted.

9 2⁻
7 (−−)
3 4⁻

Probe labelled 1L

* reclassify

9 (2⁻ or 4⁻) 2⁻8⁻
7 (2⁻4⁻ or 8⁻4⁻)
3 (4⁻)

3/5/54 Lysate made of #16 Loop spotted

	750	1210	811
a.a.a. =	0	0	0

2/13/54

W2380 lysate on 2070 - After about 4 days c. 20 small plgs on lysate half - 1-5 larger plgs (reversions?) and on control ½ about 5 larger plgs - no smaller -

24 picked to gal for purification - all (-)

2/22/54 Rpt HFT Incubn with N16

N16 lysate 1/9/54 - 0.5 ml

N16 titer - 2×10^9 $\frac{200}{8.9}$ = c. 20x multiplicity 0.5 ml N16 + 0.5 ml cells

Cell titer = $892 \times 10^5 = 8.9 \times 10^7$

EML Analysis EML 867

control plate c. 900/plate - all (-)

expired plate: ½ of (1) = 2/3 = 882 per plate - indicates little killing

½ of (1) 21, 23, 35, 24 = 103/4 = 26 $\frac{0.03}{870 \overline{)24.00}}$ $\frac{2610}{}$ = 3.0% (+)

Set HFT Incubn with N16 1/9/54

Overnight culture 1/10 - 1 ml sedimented - resuspd in 0.5 ml N16 added - about c 3 min centrifuged - Rpt 3 times except as centrifugation after 3rd - dil $10^2, 10^2, 0.5/10$ → plate

See above

Control plates:

37C	371, 341, 319 = 1031/3	$344 \times 10^6 \times 2 = 688 \times 10^6 = 6.8 \times 10^8$		$\frac{36}{344} = 10.4\%$
30C	360, 358, 361 = 1109/3	$369 \times 10^6 \times 2 = 738 \times 10^6 = 7.3 \times 10^8$		$\frac{39}{369} = 10.5\%$

N16 Phage Plates ½ · 36

37°C	$^{63}/_{763}$, $^{22}/_{309}$, $^{23}/_{207}$ = $^{108}/_{1279}$	= 8.4% (+)	/3 = 426	
30°C	$^{31}/_{175}$, $^{45}/_{197}$, $^{41}/_{194}$ = $^{117}/_{566}$		/2 = 189	

Eaton's Page 667

N16 Assay -

Testing 9 (-) on plate - 4 pfx and (+) present on any (-) colony?

30 gal - on vv

λ 10^7 dil = 106

$\frac{130}{236} = 120 \times 10^7 = 1.2 \times 10^9$

7/28 found mixed 4 pfx + 4 pfy on vv on slit

transd.

10^7 dil = plaques = 240

pipettes = 104

↑ why 2-fold diff? 0.2 ml pipette?

No control plate for expts. done on exscript = ~~84~~ ratio $\frac{\text{pap}}{\text{plaques}} = \frac{60}{240} = \frac{1}{4}$

Looking for Recombination between lp^r by selecting for (+) recombinant when two diff. gal- are lp^R

1. This is second attempt - previously no lp^+ found in (+) recombinant between 2341 gal- lp^R x 1924 gal- lp^R

2. In this case 2341 gal_2^- lp^R x W1 (= 1924 + 902 $lp^+ \rightarrow$ xg gal_2^- lp^R) out of 33 gal(+) on a Bgal mixed culture of these (probably includes sp. reversions) none found lp^+.
Not expected? since in 1st test lp^+_x x lp^R_{1924} failed to give lp^+.

What goes here? 2K41 = $\frac{Gal_2 - lp^R}{Gal_2 - lp^+}$

Where did Gal+ come from?

S18 + N16 - again for higher fraction (+)

0.4 ml N16 11/9/54 + 0.1 ml of S18 culture c. 10^8 cells/ml

cell assay of above	$10^2, 10^4 \rightarrow 5'$		3 plate	1 = 600 mw (+)
Failure? why	dilution plate	" "	$\times\frac{1}{5}$ 3 plate	1 = 163 x 5 815 mw (+)

2/19/53

246A-15 gal+$^{R}_{-}$-1
-11

246A-15 - labelled 11 reversing examined

Reversion Group	single seq (+)	HFT (2-)	Structure	Label
1	o	o	++	246A-15+$^{R}_{-}$1A
2	+	o	+-	-a crossover " " -1B
3	o	o	a-1	
4	o	o	a-1	
5	o	o	a-1	
6	o	o	a-1	

Because of above check segregants from culture labelled 246A-15+$^{R}_{-}$1 to see if conforms to original description of 246A-15+$^{R}_{-}$11

		Reversion	Minus characteristics		
261C	3/5/54	1	6/7	(--)	1 (gal$_2$-)
	Lysate of 261C →	2	stable		
		3	7/7	(--)	
750	solid spot } indicate	4	7/7	(--)	
1175	" " } HFT				
511	" " }				

According to this it is 1A

Lysates of 261A + 261B made

Ratio of (+) to minus colonies in	(+)	(-)	Examination of the (-)	
261A	82	44	11 (2-)	4 (--)
261B	103	63	10 (2-)	6 (--)

Acting Lysate	261A	261B
	deep	deep
750	HFT	HFT
2175	HFT	HFT
511	HFT	HFT

(262)

2/21/54

257C-2 S18 + N16 → t^q p^R - Examination of the segregants

Segregant	from this S18 mon lys	N16 HFT 2⁻	S18 HFT 4⁻	t_q p_R			2 + 3
1	"	0	0	S	1	+	- - S (5) (9)
2	"	0	+(?)	S	mt		- + R (6)
3	"	0	+	R	✓	—	+ - S
4	"	+	0	S	?	—	
5	"	0	0	S	?	—	
6	"	0	+ / poor growth	S	mt		
7	"	0	+	S	mt		
8	"	+	0	S	?	—	
9	"	0	+	R	✓	—	
10	"	0	+	R	✓	—	
11	"	0	0	S	3		
12	"	0	0	S	4		
13	"	0	(0)	(S)	5		
14	"	(0)	(+)	R	✓		
15	"	+	0	S	?		
16	"	0	+	R	✓		
17	"	0	+	R	✓		

Not certain that these segregations came from separate (+) -

The vital reactions on P_{12} and in any case from $1^- 4^-$ mutant are not (+).

Since all crosses with 1 - 2 - result in wild type

1 - 2 -
1 + 2 -

On these segregants 1, 2, 6, 7 in the table cured by

1 - 2 -

In which from the [illegible] mutant has been

1 + 2 + 4 -
1 - 2 - 4 +

1 - 2 - 1 +

been observed before

2/20/54 — gal₁ is the original (−) that is depleted for some portion of this gal region

246A-15 — Streaked out and (−) colonies picked and tested against Both HFT gal₁- and gal₂- . (−) about gal₂- found - no gal₁- ← disagrees with earlier classification

The gal- by the present test - were plated and (4) colonies obtained and streaked out on B gal

gal₁ -	Reversion	
1	unst.	+
2	"	+
3	stab.	
4	unstab.	+
5	"	+
6	stab.	
7	unstab.	+
8	"	+
9	stab.	
10	"	
11	unstab.	+
12	stab	

6/18/53

These stocks discarded -

(5 stable 7 unstable)

246A-15 — protrophs stupid! arose with 1503 gal₁⁻ EMS gal

	2	1	
246A-15	−	−	
	+		→
1503 gal⁻	+	−	

246A-15 selected = F⁻

1503 F⁺

control	plate	2	1	(−)
plate	1.	27	3	0
	2.	43		
	3.	25		3

10/26/56

Future check these colonies were from H. or were protrophs.

2/22/54

1503 - (-) direction to obtain gal- in Hfr streaked To streak W2427

1. 2 gal$_2$ - obtained one Lp^s W2427
one Lp^r W2429

2. 1 gal$_1$ - obtained one Lp^s W2430

Also Thes day
2431 = 2478-1
2432 = 1673 gal$_2$-
2433 = T18
2434 = T19

Crosses of gal- . Control streaks on B gal = pure

1. ⊗ 750 (rotated) - c. 30 (small)/plate × 11 = 0+/330

2. ⊗ 1210 (rotated ↑) - 9, 8, 13, 12, 12, 14, 12, 6, 6, 8 = 0+/101

3. ⊗ 811 - 3/17 · 3/14 · 0/15 · 1/28 · 1/17 · 4/15 · 0/16 · 5/23 · 0/29 · 2/16 = 19+/190 - (+) genotype of 18 checked to see if diploids. 7 found (+)

5 of which may be diploid

4 were

Analysis of these diploids by examination of segregants

#2 9 (-) segregants tested against HFT 1-, HFT 4- - all Lp^{tvr} and gal$_4$-

#3 10 (-) segregants " " " " - all Lp^{tvr} and gal$_4$-

#4 8 (-) segregants " " " " - all Lp^{tvr} and gal$_4$-

#6 12 (-) segregants " " " " - all Lp^{tvr} " gal$_4$-

W2360 - a Lp^s from ♂ W2350 gal$_2$- gal$_4$-
6 colonies tested from old broth streaking out from CAC are Lp^s and not transduced by HFT 2 or HFT 4 in two days

Stock made

3/4/54

Examination of Transducher Complexes (+4) for HFT nature if at all
Stocks have been maintained in slants for considerable time (6 mo ?)

	Designation Stock Slant	Loops spotted /750	/R10	/871	characteristic of org. (-)	(+)	mean lysates
	518tK12	solid spot	solid spot	solid spot	19	53	5
	750tK12-1	solid spot	solid spot	solid spot	13	79	8
not stencils	750t902-1	solid spot	solid spot	solid spot	26	110	8
not stencils	750t902-2	0	0	0	14	c.50	4
not stencils?	2175t750-1				4	c.50	4 pure plate 1-2
not stencils?	2175tFW-1				+	2 Kinks ++	?
both Lokelled 2175t871-1	518t750-1	+++	+++	+++	18	83	?
	750tK12-2	+++	+	+++	14	32	5
	750t902-2	+++	+ ck	+++	27	62	6
	2						

HFT Stocks

Designation	Page	Genotype	Comment	Autotrophic Markers
2342 (N16)	192B	gal_2 -		none
2346 (N14)	230	gal_1 -		M -
(D1)	153	gal_2 -	from 592d — x 57F	M -
(S18)	202	gal_4 -	probably NFT now	M -
(#14)	242	gal_2 -		Arg
(#19)	242	gal_2 -		none
246A-15	246A	gal_2 -		none
202-16H	202	gal_2	S16 has an NFT E, it	M -
257-2	257	gal_2 -	has an NFT E it	M -
257-4	257	gal_1 -	" " " "	M -
247B-1	247	gal_4 -		M -
261A	261	+ u		none
261B	261	+ u		none
261C	261	+ u		none
217S2750 + u		+ u	750d — x 217S	none
57BE1412	263	+ u		M -
750t1412-1	263	+ u		M -
750CR02-1	263	+ u		M -

3/21/53

Tests of some lysates for HFT and allele - See also page 265

Lysate	750	1210	811	Comment	Cell Characteristics Making up lys	Stability of +	Yellow crayon #
2342 (3/4/53)	+++	0	+++	Known gal⁻ HFT	—		1
2342 (3/17/53)	+++	0	+++	Known gal⁻ HFT	—	64 12/12 +⁹ ①	2
2175 t811-1	++	0 small	+++		mixed		3
750 t612-2	+++	++	+++	behaves as + HFT	mixed		4
750 t902-2	+++	+	+++	" " " "	mixed		5
2478-1 pop 2	1 pg	2 pg	0	gal⁻ HFT lost?			6
2175 t811-1	0	0	2	NFT?	mixed		7
5Ft 750-1	+++	+++	+++	behaves as + HFT	mixed		8
750 t902-2	1 pg	0	2	NFT?	mixed		9
258-16	5 pg	1 pg small	0	gal⁻ unstable [illegible]			10
2464-15	+++	0	+++	gal⁻ giving +ᵘ revertants			11
2616	+++	++	+++	aggregates (--) principally			12
5Ft K12	+++	c. 100 pg	+++	behaves as + HFT	mixed (+)(-)		13
750 t K12-1	+++	++	+++	" " " "	" " "		14
750 t902-1	18 pg	2 pg	c. 30	NFT?	mixed		15
2346					—	5 4/5 +⁴ ②	

+++ = solid drop
++ = some discret. pop

two dif. cultures mislabelled here

Rev. found?

4/10/54 all typical strain in Pen. 0.5 ml single

The gal⁻ revs is 2175 t811
other 2175 t751

* Gal+ Reversion Stability - no obs., no found +⁴/no. of comments

4/19/54	750 t902-1	0	0	0	16 abs
4/1	750 t912-2	+++	+	—	17 typical strain

3/22/54

Transduction with 2342 (=W16) in NSB

	No (+)	No. (c)	Noc-) contam.	Noc-) Calc gap	dilutions	
(1.) W 518						
Lysate	8	140*	9	1	resusp. in 10 ml, 10^2, 1-50 → 0.1 ml	$8/140 = 143/8.00 \frac{0.057}{\frac{280}{1750}} = 5.7\%$ (+) transd.
Broth	0	679*	0	2	"	on Lp^s

* difference in numbers probably due in part to loss in centrifuge tube decantation

2. W 811

Lysate	1	426*	0	5	above	$426 / \frac{0.0021}{1.000} \frac{852}{1750} = 0.25\%$ (+) transd.
Broth	0	448*	0	1	"	

* two colony type - large and small.

3/25/54 Experiment with TCN - Transmission of "injectate" in cross.

(Hfr $Lp^+ Lp_2^s$ streaked) spread on 2342 λ (X) F⁻ $Lp_1^+ Lp_2^R$ TLB = Gal_2-

Experiment Control run in first place — gal_2

Absorption of 2342 λ on (2341 streptomycin treated) 1 ml phage + 1 ml cell suspension

Before $10^7 = 273, 214 = 2.44 \times 10^9$ $\frac{1}{2} = 1.22 \times 10^9$ (average of four plates on B(c)) ‖ No absorption $\frac{1.2}{1.2} = 0.0$

After $5 \times 10^7 = 70, 26 = 1.15 \times 10^9$

on 2252

Before - above

After $5 \times 10^7 = 15, 12 = 7.0 \times 10^8$] $\frac{7}{12.}$ $12 \left/ \frac{0.58}{\frac{7.0}{100}} \right. =$ 42% absorpt.

3/30/54 Repeat examination of absorption of S^c cells.

W 2341 and W 2342 HFT λ (assays by pour tubes)

1. Pre absorpt. titer = $247, 167 \times 10^7 = 2.07 \times 10^9 = 2.1 \times 10^9$ (compare with above)

2. S^c cells (6.3 ml) + 0.3 ml 2342 λ (NSB prep 1/9/54) - incub. 37 C 10' - 9.4 ml fresh added - centrifuge 5 min.
dil 10^2, 10^4 → 0.1 = 444, 482
titer = $463 \times 10 \times 33 \times 10^4 = 3.3 \times 10^6 = 1.53 \times 10^9$ % Not adsorbed. $\frac{15}{21} = 71\%$ 29% absorpt.