

3/28/54

SIFE 750-1 (Look up origin) Nature of segregant.

Seq.	HFT-1	HFT-4	Locus
1	o	+	1-
2	o	+	"
3	o	+	"
4	o	+	"
5	o	+	"
6	o	+	"
7	o	+	"
8	o	+	"
9	o	+	"
10	o	+	"
11	o	+	"
12	o	+	"
13	o	+	"
14	o	+	"

what happened to 4-?

all d^R

By site made of here to and of HFT-Hft & has any additional properties

2341 *trichoductin* (for T₁) using his culture which was found 4⁵)

7. K12 lambda

1. mode = 4
2. 0.1 K12 (4%) = $32 \times 58 = 116 = 1856$

Use in cross of added segment as a test of elim. and passage of segment in 5th cross.

not done beyond

2341 (abm) 2. 902 lambda - (indate HFT Hft)

7. in add 2
2. 0.1 902 42

4/7/54

In relation to SIFE 750-1 Above. Transduction of 811 with HFT 1- 2346
 Examine: stability - Do the 1- 1- combined +? Does 1- and 4- segregate
 from such a complex? Are the observed (+) in 4, 1- transductions
 (+) after elim. of (-) allele?

1. 8 peculiar examined 4/8 stable.

	HFT 1-	HFT 4-	HFT/1-	HFT/4-
A. 1.	-	+	+	-
2.	-	+	+	-
3.	-	+	+	-

269-3

3/28/54

Examination of HFT stocks

1. Previously noted (in last week) that cultures of N16 (= 2342) and NA-4 (= 2346) were segregating NFT lines (opt plate test) (unad. on plate possible - loopful added to indicator, plate unadicated). On each case of 10 colonies 8 were HFT - From each a NFT and HFT selected and plated on slants and stored in refrigerator to lessen segregation. A portion of each plated on Royal and reversion examined - (See pg 267)

2. 241-14 (a gas₂ - HFT) tested against 1210, 750 - 10 colonies
 against 750

⊙	⊙	⊙	⊙
⊙	⊙	⊙	⊙

 - 8/10 HFT. Stocks made of an HFT, NFT
 against 1210

⊙	⊙	⊙	⊙
⊙	⊙	⊙	⊙

3. 241-19 (a gas₂ - HFT) as 2 above
 against 750

⊙	⊙	⊙	⊙
⊙	⊙	⊙	⊙

 against 1210

⊙	⊙	⊙	⊙
⊙	⊙	⊙	⊙

Examination of NFT, HFT stocks above - also 2342 2346 - See page 267 also

Cultures	No. Reversions	Stability	Transmission units?				K/Slide
			1/4 HFT	1/1 HFT	1/2 HFT	4 HFT	
241-14 HFT	19 (Square)	12/12 + ⁿ	(+)	+	0	(+)	2-
" NFT	12	12/12 stable	+	+	0	+	2-
241-19 HFT	12	12/12 + ⁿ	(+)	+	0	(-)	2-
" NFT	0	-	+	0	0	12/12 +	1-2
2342 HFT	62	12/12 + ⁿ	+	0	0	+	1-2
2342 NFT	0	-	+	0	0	+	1-2
2346 HFT	15	4/5 + ⁿ	+	0	0	+	1-2
2346 NFT	8	8/8 stable	+	0	+	+	1-1

4/6/54

Examination of HFT by plate

1. 7506K12-2 - Assay on different indicators - 0.1 ml of overnight unassociated cult. unal.

dil	no. pop.	Assay cult	dil	no. pop.	Assay cult.	dil.	no. pop.	Assay cult
na	0	750	na	0	2438 (r2)	na	3	12-10
10 ²	+++	"	10 ²	++	"	10 ²	2	"
10 ⁴	565	"	10 ⁴	214	"	10 ⁴	23	"
10 ⁶	12	"	10 ⁶	4	"	10 ⁶	1	"
<u>titers</u>	1.2 x 10 ⁸		4 x 10 ⁷	2.1		2 x 10 ⁶		

2. 7506902-2 4/8/54 (lysate that worked)
 Filtered on U2 (sp. 1st) 10⁷ = 717 = 7.2 x 10⁹ - lysate tested stock prob

Assay	750	1210	2433	Cristiditria presumed
no. cells	0	3	0	1-2+
0.1 ml of dil 10 ²	+++	1200-1600	183	1+
" 10 ⁴	++	13	1	
" 10 ⁶	12	?	0	
<u>titers</u>	1.2 x 10 ⁸	1.0 x 10 ⁶	1.83 x 10 ⁵	

suggests

4/10/54

U2 - spmt 10⁵? from 2175T511 → (-) seg 10⁵ seg 4 - Only
 useful marker for elim. possibility of contamination to
 path. proph. This culture grows on EMJ. gel. Presumably
 this is a valid occurrence of 10⁵ - No other seg 4 - 10⁵
 proph. about at time of isolation.

4/5/54

In search of triple (-) 1-2-4-

7. W2350 ^{related to} ~~isolated by~~ HFT 1- (2346) - Segregants skinned

Seg #.	HFT 1	HFT 2	HFT 4
1	+	0	+
2	+	0	+
3	0	+	+(weak)
4	0	+	+(weak)
5	+	0	+
6	+	0	+
7	+	0	+
8	+	0	+
9	+	0	+
10	+	0	+

2-
1+
4-
X-
-
+

In these cases
a few small papillae noted
similar to the 1% of 2350 against
HFT 2 - Significance?

8 2-
2 1-

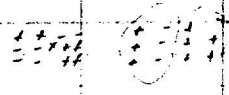
8. T18 = W2437 ^{related to} ~~isolated by~~ HFT 4- (2478-1) - Segregants skinned

Seg #.	HFT 1	HFT 2	HFT 4
1	0	0	+
2	0	0	+
3	0	0	+
4	0	0	+
5	0	0	+
6	0	0	+
7	0	0	+
8	0	0	+
9	0	0	+
10	0	0	+

272-2

all 1-2-

2-
1-
4+



4/7/54 - and several days earlier

Recovery of l_p from l_p cultures

W902

1. dominant = 2346 HFT 1-1 - Reaction suggest family large fraction of l_p present.
2. checked but B got - seg + observed.
3. just for seg anal (c.b.) all l_p
4. One titer for nutrition -

D(0)	no growth
+ T2	growth 2 days
+ LB	no growth
+ TB ₁	no growth
+ T2 ₂	growth 24 hrs.
5. Examination of imm segs and

nutrition sh. tested on
make sh. 13. Mal = met+
 W2584

W945

1. dominant = 2346 HFT 1-1 - React. also suggest family large no l_p present.
2. checked B got - +² observed.
3. 6 seg titer / 2346 / 2342 - all l_p
4. Examination of the culture for sugar reactions.
 Mal + Xyl - Mtl -
 Bacteriological culture also found Mal +

W2580

4/9/54

In search of phage mutants.

Platings on E. coli - all platings gave no plaque.
 With the following cultures.

Assay on NSA (pumps)	10^7 dil	10^8 dil
2281	274	10
1485	27	10
578	281	6

4/10/54

Repeated at lower dilutions

Sp. l_p →	U2	10^3	10^5
578	0	0	0

4/14/54

Is the complex $\frac{1+4^+}{1-4^+}$ phen. (+)? All (+) appear to be secondary - segregating one (-)

SIF # 2746 λ - Single exposure to phage - unacrated fish, sea water. per cent.

①

Treat	SIF cell	No. phage (+)	No (-)	No. pure (-)	Total
1. broth	10 ⁶	0	0	67	67
2. 2746/HFT 1	"	9	2 (populab)	41	52

② Examine (-) on treated plate to see what they are segregating

- Single (-) appearing colony (up to 2 days - then giving a populating appearance thus \odot , and turning darker in the following day)
- Streaked out and giving

- (+) some of which appear to be segregating (-)
- (-) non populating (?)
- parental type (-) which papillate

3. 10 (-) colonies picked and streaked against HFT 1 + HFT 4

#	HFT 1	4-	lp Rk	Comment	Allele
1.	+	0	5	} lp to segregating also	4-
2.	+	0	5		4- (?)
3.	mixed (+) and (-)	1 ⁺	5		4- (?)
4.	+	0	5		-
5.	mixed (+) and (-)	1 ⁺	5		-
6.	"	"	5		-
7.	"	"	5		-
8.	0	+	5		-
9.	intermed (+) turning dark after 3 days	"	5		-
10.	"	"	5		-

Only Segregants are 4- lp
lp to Rk remaining segregating 7 1-

parental
papillate
from
parental
segregants
parental
parental

4/28/54

902 Mal- → Mal+ = Ap_2^R → lp_2^S ?

10 well isolated colonies (-) on Mal, tested and found lp_2^R checked on Mal for reversion exam. (1 / streak) and in gal coated with HFT 1-

1. Exam parital on gal for instability, indicating lp_2^S + transduction.
 2. Exam Mal + for lp_2 reaction.

Original Colony	No. reversions B mal	lp_2 Reaction of Mal+	No. pop. B gal + HFT 1	Stability of Gal pop. (checked)	Mal + lp_2^S sample characteristics	Transd. of seg. out from HFT 1/HFT 2
1	+	lp_2^S	31	1/6 stabl	mal- lp_2^R	0 0
2	0	—	16	4/6 stabl	" "	0 0
3	+	lp_2^S	18	0/6 stabl	" "	0 0
4	+ intermed (+)	faint reaction	12	5/6 stabl	" "	0 0
5	+	lp_2^S	14	2/6 stabl	" "	0 0
6	0	—	10	3/6 stabl	" "	0 0
7	+ intermed (+)	faint reaction	16	3/6 stabl	" "	0 0
8	+	lp_2^S	26	1/6 stabl	" "	0 0
9	+	lp_2^S	12	1/6 stabl	" "	0 0
10	+	lp_2^S	21	4/6 stabl.	" "	0 0

6 lp_2^S
 2 intermed seg.

All run transducible

Σ. mal+ reversions are lp_2^S or intermediate sensitive

Selection here may not be for the occurrence of lp_2^S under conditions of extreme purity of cell population but may be for host range mutants in the HFT phage population.

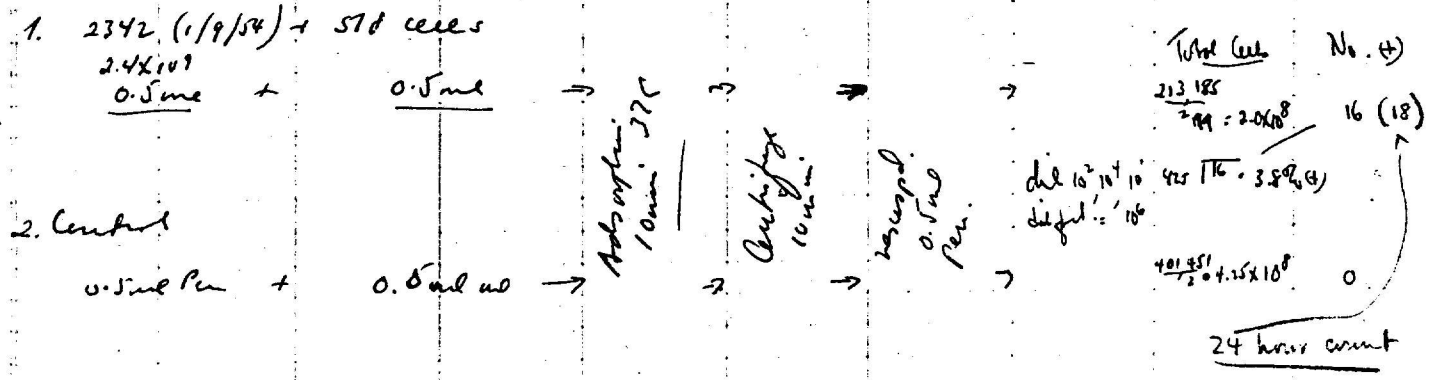
③ Lysozyme made of #1 (unstable, and seq(-))

atml/	results
1. 750	solid, amor
2. 578	" "
3.	

5/6/54 W2331 mal- tested for band. with N16 4/9/54 (HFT) not found band.

5/4/54

J.L. Single cell cft.



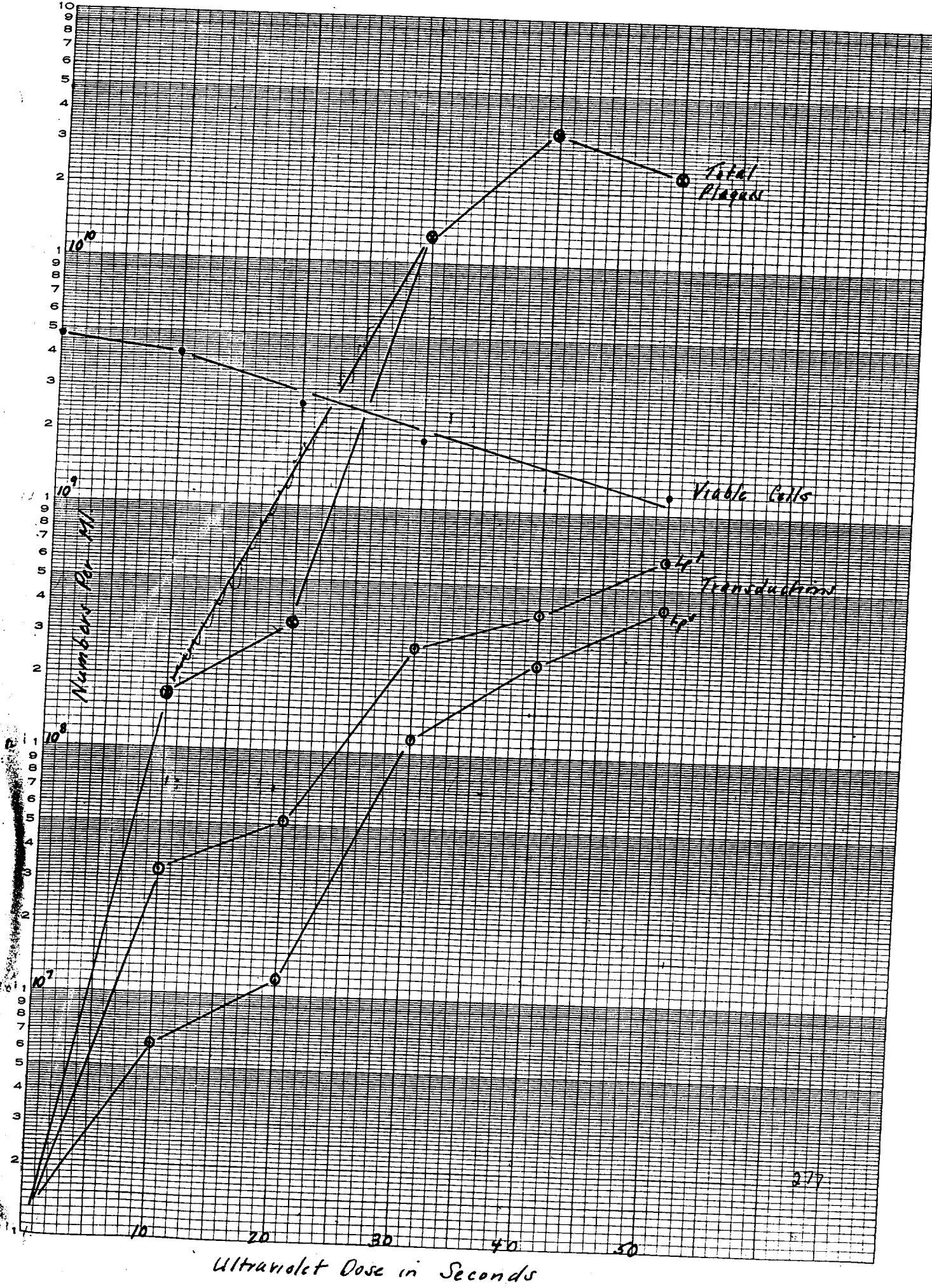
It reports no. found present in his sample of cells studied in the micrograph.

5/6/54

This day 2252x Hft gal⁻ (247134) entered in stock book as W2487 presumably made and tested c. Mar. 24, 1954 - Culture to TCN at this time also.

5/6/54 For the purpose of checking the transmission of the plaque in crosses. No sugg. by Jk. Parent 2252 (=Hft gal⁻ gal⁺) to $\frac{++}{-+}$ and then cross to $\frac{+-}{-+}$ ⊕ $\frac{+-}{-+}$

1. 2252 treated Hft gal⁻ (with $\frac{Hft}{100}$) colonies examined for "phage center" appearance - some found treated out.
2. on second streaking, all + still



5/26/54

2342. Test on survival to see if inability to get lysates & high titers is caused by application of wrong a.v. dose.

1. Aged culture, nearly saturated - centrifuged and respl. in saline

2. Inoc

0
10
20
30
40
50

0.1 ml
addback
to
10 ml
Pen
Pen
Inoculated

Concn	Single plate	N/N ₀	Plaque (later) 1-100 dil	Transd. plate x 10 ⁸	Plaque / 1-100 dil
0	477	1.0	0	50	0
10	417	0.88	103 x 10 ⁶ · 1648 x 10 ⁵	112	62
20	263	0.55	c. 2 x 10 ⁶	164	114
30	194	0.41	1284 x 10 ⁶ · 1.5 x 10 ⁸	1200	115
40	(256) -	-	5456 x 10 ⁵ x 10 ²	2368	238
50	117	0.25	-	4160	418

1-100

10⁶

Trans	Sample Cell Count	Inoculum Cells	Trans. produced	Plaque produced
0	4.77 x 10 ⁹	0	0	0
10	4.17 x 10 ⁹	0.6 x 10 ⁹	6.2 x 10 ⁶	1.65 x 10 ⁸
20	2.63 x 10 ⁹	2.14 x 10 ⁹	1.14 x 10 ⁷	2.3 x 10 ⁸
30	1.94 x 10 ⁹	2.83 x 10 ⁹	1.15 x 10 ⁸	1.24 x 10 ¹⁰
40	1.59 x 10 ⁹	3.12 x 10 ⁹	2.32 x 10 ⁸	2.46 x 10 ¹⁰
50	1.17 x 10 ⁹	3.60 x 10 ⁹	4.1 x 10 ⁸	2.4 x 10 ¹⁰

Transd. assay on PH

no a.v. Inoc. Proc

17

10'
20'
30'
40'
50'

16 x 200 x 10⁴ = 3.2 x 10⁷
32 x 163 x 10⁴ = 5.2 x 10⁷
169 x 16 x 10⁵ = 2.7 x 10⁸
379 x 10⁶ = 3.8 x 10⁸
654 x 10⁶ = 6.5 x 10⁸

Plaque Transd.

16.5/3.2
33/5.2
103/2.7
340/3.8
240/6.5

Ratio

5.2
6.4
38
90
37.

6/1/54 Transduction of l_p^R with HFT lysates - An indication of host range change in λ ?

1. 1912 (l_p^S l_p^R Gal⁻) transid = 2342 λ
 no add 40
 23% λ 456

- #1 #2 2. 30 papillae picked - after 2 streakings, all stable in Gal
 #3 3. 24 Gal⁻ papillae - after #2 2 unstable - ~~65%~~ non-lys in STP, Gal⁻
 #4 #5 4. 24 " " " " 2 streakings 1 " - ~~65%~~ Gal⁻, non-lys in STP
 5. 24 " " " " 2 " - ~~65%~~ Gal⁻, non-lys in STP
 also 23 stable Gal⁺, non-lys in STP

6. Examine Mal + Reversion of above and see if sens.

Mal Reversion	Gal	lys	Mal ⁺ Revert	Gal ⁺ of Mal ⁺	lys	Revert ⁺ of Gal ⁺	Gal ⁺ of Revert ⁺
intermed #1	+	L	full	full sens.	2	non-lys	+
full #2	+	L	"	"	3	non-lys	-
full #3	+	R	"	"	4	"	+
intermed #4	+	R	intermed	intermed sens.	2	"	+
full #5	+	R	"	"	?	?	?

Suggests that transid accomplished by l_p^R forming phage

278

Eliminate to see if reversion of l_p occurring - Tested against λ , STP

Seq. Tested	STP	STP	STP	STP
1. 5 non-lys	+	+	+	+
2. 5 "	+	+	+	+
3. 5 "	+	+	+	+
4. 5 "	+	+	+	+
5. 5 "	+	+	+	+

Indicate some cultures look for further phage

Strains made as 278-1, 2, 3, 4

7/7/54 Lysate made for full⁺ Mal⁺ l_p^R - no plaques found before irradiation
 #1 per med. streak indicates mixed for gal - lysate showed no evidence of plaque or transid in STP

10/26/54 - In looking for whether selection for (Mal⁺ l_p^R) intermediate from admixing λ selected by HFT except above. Strains found to be reversion of Mal and l_p^R . Two fully λ^+ l_p^R found from #3 #4, (30% reversion of l_p^R)
 both found stable such that it is likely more l_p^R are not digested

STP λ for hybrid lambda prep. Labeled 6/1/54
 1. Titer = $209 \times 10^8 = 2.1 \times 10^{10}$

STP λ 2342 - Attempts to get higher fraction of transducing

1.0ml cells	(-)	(+)	total	cell count	
add to	① No add - 216	0	216	$\frac{215}{3} \times 10^5 = 7.2 \times 10^6$	
lysis	② 2342 lysate (65%) 206	10	215	$\frac{215}{3} \times 10^5 = 7.1 \times 10^6$	c. 5% (+)

Dec. 5 min
 centrifuge precip
 resuspend in 1 ml
 saline

W2877
 2878
 2879
 2880

6/17/54

279-1 14/12/2342 plate - the sporoblasts (+) - Revert ^{if picked} and examine by ^{for Mal} Koch.

279-2 14/12/2342 plate - 12 picked from by site pulsed of plate - to see return of by of the stable gal (+) - by site pulsed of plate:

1.	2.	3.	4.	5.	6.	7.	8.	9.	10.	11.	12.	13.	14.
1. All 12 mal-gal + were found to be L ² R	1. 100% by	R ³	5 wk	+	+	+	+	+	+	+	+	+	+
2. DeMalt ^P (of varying degrees + rev)	2. "	R	5	sq	+	+	+	+	+	+	+	+	+
3. "	3. "	R	5	sq	+	+	+	+	+	+	+	+	+
4. "	4. "	5 wt confined	5 wk	sq: 1/4	rev	sq: 1/4	+	R-	+	+	+	+	+
5. "	5. "	5	5	+	+	+	+	+	+	+	+	+	+
6. "	6. "	5	5	+	+	+	+	+	+	+	+	+	+
7. "	7. "	R	5	sq	+	+	+	+	+	+	+	+	+
8. "	8. "	R	5 wk	+	rev	stbl, + R	+	+	+	+	+	+	+
9. "	9. "	5	5	(-)	+	+	+	+	+	+	+	+	+
10. "	10. "	5 wt confined	5 wk	+	rev	stbl, S-	+	+	+	+	+	+	+
11. "	11. "	5	5	+	+	+	+	+	+	+	+	+	+
12. "	12. "	5	5 wk	+	rev	stbl, R-	+	+	+	+	+	+	+
Σ	Σ	4 spore + 1 ^S	3 trans. + 1 ^R	4 " + 1 ^R	Σ	14 spore (+) 1 ^S							

General Summary
 trans. appear to be gal + 1^R
 gal + 1^R → sq gal- 1^R
 gal + 1^R → sq gal- 1^S

279-3 2580 t K12 F⁻ X 1321 3 plates
 Control plate ~ 2580 t K12 F⁻ for distribution of (+), (-) gal (100% 5X-)

1. Three plates from "pneumophs" early (1st day) - all (+)
 After 2 days - c. 2000 / plate - 1 apparent (+) instead of (+) on streaking on B gal - Squiggle gal - mixed type (2)
 6 sequences of this (+) tested against HFT 2, 4 - all transduced by 4, ∴ Gal⁻ from 2580 t K12 F⁻

279-4 2580 t K12 F⁻ X 518 3 plates (discarded one, similar to others because of mixture.)

1.	2.	3.	4.	5.
1. Control on (+), (-) in 2580 t K12	15 (+)	13 (-)	30 (+)	30 (+)
2. Plate 1.	300	53	353	30, 4 were apparently stable (+), of the remaining 26, 25 tested microscopically against HFT 2, HFT 4 - all transd. by HFT 4, ∴ Gal ⁻ from 2580 t K12 F ⁻
3. Plate 2.	241	46	287	
4. Plate 3.	541	99		

6/21/54 2342 purified of 10 colonies examined 7 were HFT on 84 - new stock prepared

For reporting above transmission experiments
 1321 transd. by K12 →

1321 t K12 F⁻
 Stock - 279-5

6/24/54

280-1 Are the bands of l_p^+ really of l_p^R/l_p^+ nature? Examine on l_p^+ bands. W 2570542

- ① The Gal (-) say (20)
- ② The Gal +ⁿ (20)

-(280-1) the Gal- (In General these were from tests) all found to be l_p^+ and l^R

(280-2) the Galⁿ is found to be l_p^+ and l^R . 2 tests were too poor to judge

980-3 Lytic λ - 1485 - 0.5 ml of this lysate into Per - sterile after 2 days.

① 1485 grown and NSB overnight - then aerated culture in NSB started until nearly full density, - centrifuged rapid in 3.0 ml. Sd λ (ambly) (2.78) tube - 2.1×10^{10} - Centrifuge rapid in c. 20 ml NSB aerated. After 1 hour partially cleared. Centrifuged and dec'd

② Assay

$10^7 = 757$ - tube = 7.6×10^7

③ Transduction with

in D(0) 0.1 ml this lys + 0.1 ml 81 overnight broth cells. - no clones after 48 hours - No M ductin

④ Coelution

	Expt/Control	Stability	
1. W 750	2/0		After 2 shakings - 2 was neg stbl (+) directly 9 stbl (+) after 2 shakings
2. W 1110	2/1	2 very faint (+)	
3. W 81	10/14	1, a very faint (+)	

Further evidence that lytic λ does not transduce

Account 2342 deleted to finance restricted by in volume
 had. all 0:10 + 10 min

287-1

Time	Cell Assay	HFT Infective Centers #	- Array Tinned Centers (n=49)	plates Tinned 1-100	Plaque 1-100	A	HFT/NFT	Array of Colony of Each time
0	622	3 0	45 0	97 72	6	0 0	0	3/25 = 0.12
10	381	193 140	51 2	47 22	11	5 500	2.6	10/25 = 0.4
20	170	372 364	87 38	38 13	28	41 4100	11	9/25 = 0.36
30	63	517 516	136 87	28 3	68	62 6200	12	9/25 = 0.36
40	20	449 446	128 79	25 0	59	53 5300	12	4/20 = 0.20
60	1	259 256	98 49	26 1	35	29 2900	11	
70	1	230 227	80 31	16 0	24	18 1800	9	

* Some of these infective centers appear to have papillae in them. These also appear to be papillae cont. infective centers about them.

↑ these values cannot be trusted since they represent some growth effect after mechanism of genome

Time	Total N/N ₀	P/N ₀	T/N ₀
0	1.0	-	- x3.5
10	0.61	0.31	0.0032 0.11
20	0.27	0.59	0.061 0.21
30	0.10	0.83	0.14 0.48
40	0.032	0.72	0.13 0.45
60	0.0016	0.41	0.027 0.27
70	-	0.36	0.050 0.17

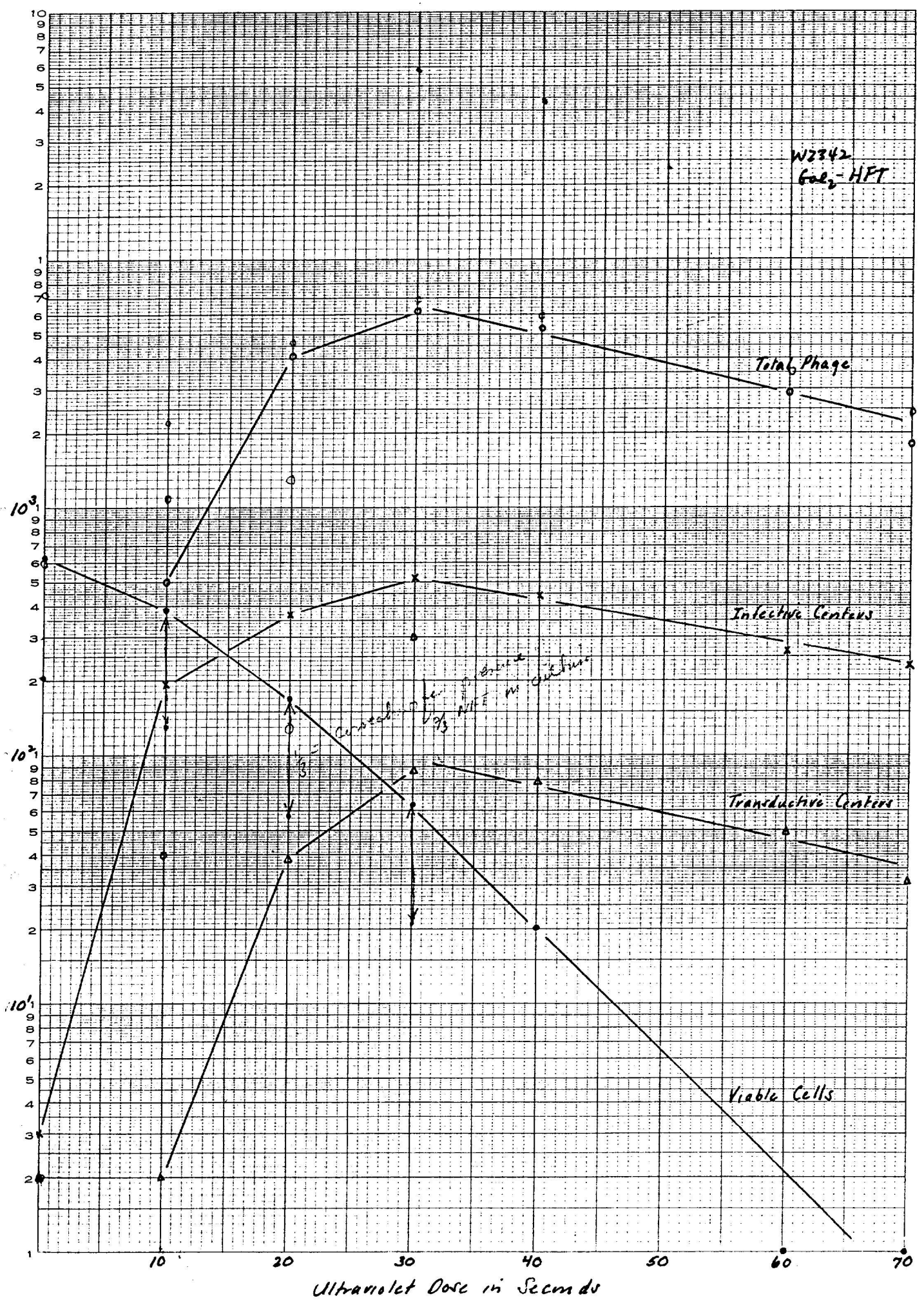
Estimate of frozen HFT

$$\frac{3+10+9+9+5}{25} = \frac{36}{25} = \frac{7.2}{25} = 0.29 \text{ HFT}$$

Corrected for LFT

Time	N/N ₀	Infective Centers	P/N ₀
0	1.0	0	
10	0.61	55	
20	0.27	103	
30	0.10	150	
40	0.032	129	
60	0.0016	74	
70	-	66	

N₀ = 622 x 0.29 = 180



282-1 In the 1st data so far suggests lysates of 1st carry more often than not the (+) of the segment. Really so?

1. 2175E750-1 Sheathed out and 5 colonies picked - Cultures made and lysates of these prepared.

Culture	W 282-1 Assay/750	no. cells - 4500 Assay/1210	Plaque/SIT	Comment	Ratio #1	Comment 2
1	1124/448	49 (small)	533x10 ⁶ x 2.1x10 ⁸		2132/497 = 4.3	Higher titer in Gal ₁ than 2 suggest that fragment is 2- or Gal ₂ ⁺
Lysate Culture 2	523/460	29 (small)	307x4x10 ⁶ x 1.2x10 ⁸	(omit 2x as many minutes plating)	1225/466 = 2.6	
3	336	36 (")	134x4x10 ⁶ x 5.4x10 ⁸		544/372 = 1.5	
4	281	57 (")	217x4x10 ⁶ x 2.7x10 ⁸		868/337 = 2.6	
Lysate Culture 5	382	34 (")	191x4x10 ⁶ x 7.2x10 ⁸		764/416 = 1.8	

due to lower titer with 1210? possibly a 12175

2. Single colony isolated from each and tested for allele - all were gal⁻

282-2 Transd. of W282 with HFT 1st → Lysate = 282-2
408 (-), 2 (+)

also gal⁺ derivative cultured

W282 | W2851

① W282 cells exposed to HFT 1st λ (NA-4 prep?) plated out. Many partially lysed colonies after 3 days & colonies showing some evidence of ⊕ possible transd., but predominantly ⊕ in streaks, also appear to be segregating. Original colonies were not ⊕, but pink in color, also of large size as if had selection advantage in Gal.

② Search for ¹⁻⁴⁺/₁₋₄₋
 (a) 1 discarded cultures #1. Lysate prepared - streak of culture before done c. 50-50 ⊕/(-)
 started from (-) colony - assay of lysate (spores) in 575, 1210, 750 - all HFT working
 (b) Single (-) pop. colony (the original) → streaked out → single ⊕ colonies streaked out: 24 in number
 (c) 4/14 stable ⊕

Remarks	HFT 1 st	HFT 4 th	Result	Comment	HFT 1 st	HFT 4 th	Result	Comment
Remnants from (-) came from pure ⊕	+	⊕	Result	Gal ⁺ - ④	0	+	Result	Gal ⁻ ①
1	+	⊕	Result	Gal ⁺ - ④	+	0	"	Gal ⁻ ②
2	not gal ⁻ (-), but revertible	"	"	parental	+	0	"	Gal ⁻ ③
3	"	"	"	"	+	0	"	Gal ⁻ ④
4	⊕	+	Result	Gal ⁻ ①	not gal ⁻ (-) but revertible	-	-	parental 1-4+ / 2-4+
W2851	+	0	"	Gal ⁺ ④	Σ	parental	4	} all tested against
283-1 →	0	0	"	Gal ⁻ Gal ⁻ ②	Gal ⁻ -	6	Gal ⁻ -	
7	⊕	+	"	Gal ⁻ ①	Gal ⁻ -	5		
8	⊕	⊕	Result	?	Gal ⁻ -	2	} all tested against	
Discarded	+	0	"	Gal ⁻ ④	Gal ⁻ Gal ⁻ -	1		
283-2 →	0	0	Result	Gal ⁻ Gal ⁻ ②	Unobtainable	1	} all tested against	
5/13/55	0	+	"	Gal ⁻ ①		1/8		
11	+	0	"	Gal ⁻ ④				
12	+	0	"	Gal ⁻ ①			} all tested against	
13	0	+	"	parental				
14	not gal ⁻ (-), but revertible	"	"	parental				

These are not derived from 282-1 or 283-1
 HFT

7/11/54

283

283-1 } from previous page - are presumably Gut. - Gooey
283-2 }

283-1 Transd. \bar{c} HFT \bar{c} to get type \bar{c} .

- (1) Reaction weak - (+) not strong in cross break - segregation $\frac{1-4-2+}{10402-}$ is not pure + ?
- (2) Stealing out - suggestion that (+) may be of two classes. - one pure (+), other unlabeled.

A 283-1 cross A = \odot 2234
B cross B = \odot 902 gal,

(A) 1st cross left prod - many 616 prod. - no (+) observed

(B) $\frac{1}{8} = 83$, = 667×10^6 plates = 6670 (-), no (+) observed.

c 283-1 Further test - transd. 283-1 \bar{c} HFT pre + λ and obtain (+) - this should then segregate from σ type (-) which revert rapidly - whose genetic constitution is $\frac{1+4-}{10402-}$

(1) Transd. \bar{c} σ + K12 (#13) p. 267 - dil 10^6 = solid mass, dil 10^8 = cross, dil 10^6 = 20 (loop free)

(2) 24 (-) seg checked out - all appear to be stable (-) - allele = all found 1-4-

(3) rechecked 12 (-) seg all stable (-)

283-3 Continuation of 283-1 - #2 lysate. The small plaque? a σ mutant. Attempt to obtain lysogenic culture of - Plaque picked and streaked out, colonies tested

1. 1st 24 colonies found non lys. / 518
2. 2nd / 518

7/16/54

- A Repeat examination of +4 - after picking single colonies -
 check on lysate cell population

	14	14	20	15	32
1. 578E902-1 6(C) cl.	sterile		17	12	23
2. 578E902-1 neg, mostly +		0	0	9	26
3. 578EK12	not sterile		9	4	26
4. 750E902-1	sterile	2	2	11	6
5. 750EK12-2		5	5	11	6
6. 217E902-1		3	3	11	6
7. 217E902-1		9	9	15	10
8. 217E902-1					27

delivered to journal

Repeat array 10⁶ dilution

Hydrol	13	13	13	13	13	13	13	13
1. 578E902	0	0	0	0	0	0	0	0
2. 578E902-1	0	0	0	0	0	0	0	0
3. 578EK12	0	0	0	0	0	0	0	0
4. 750E902-1	1	1	1	1	1	1	1	1
5. 750EK12-2	0	0	0	0	0	0	0	0
6. 217E902-1	0	0	0	0	0	0	0	0
7. 217E902-1	24 (C)	24 (C)	24 (C)	24 (C)	24 (C)	24 (C)	24 (C)	24 (C)
8. 217E902-1	0	0	0	0	0	0	0	0

no. cells	wt	total	total	total
148 = 1.5 x 10 ⁸	88	75	111	74 - active in 1, 2, 4
62 = 6.2 x 10 ⁷	18	5	45	8 -
113 = 1.1 x 10 ⁸	10	0	39	2 -
99 = 9.9 x 10 ⁷	24	0	25	0
577 = 5.7 x 10 ⁸	32	18	50	13 - active in 2, 4
344 = 3.4 x 10 ⁸	10	0	35	0
615 = 6.2 x 10 ⁸	160	147	170	133 - active in 1, 2, 4
116 = 1.1 x 10 ⁸	12	0	30	0

Assay 250 (11)⁸ = 501, 5 x 10⁷
 Control = 0
 (5) 10⁴ = 24 = 2.4 x 10⁶
 (7) 10⁶ = 433 = 4.3 x 10⁷

Presumably activity within cell
 such that assay expect - other
 experiment about this time
 suggest this is true activity

284-B

1. 2487 x 2593 - Gray-146 x Gray-146 - to occur directly? not given (4)?
 2. Culture given by gelatin c. 3 hours. plated out ETS low
 10² dil - c. 10³ prob. high
 10⁴ dil - c. 1.50
 10⁶ dil - c. 20-70
 3. Streaking with of resp. c. 5 min - slightly purple in appearance, large 14 number
- | | | |
|-----|--------------|--------------|
| 1. | (-) | (-) |
| 2. | (-) | no growth |
| 3. | (-) one | no growth |
| 4. | (-) pap. cd. | no growth |
| 5. | + ches. | + (free) |
| 6. | + " " | + (unknown) |
| 7. | + " " | + (unknown) |
| 8. | - | no growth |
| 9. | + also (-) | + also (-) |
| 10. | + " " | one (-) col. |
| 11. | (-) | micro. cd. |
| 12. | (-) | (-) |
| 13. | (-) | (-) |
| 14. | (-) | (-) |

7/18/54

22799 = Gal, - tp^s - Repeat of 578t HFT, using Gal, cells instead of Galy

1. Actively growing culture of 22799 in Pan. c. 10^8 cells per 0.5 ml cells + 0.5 ml NA-4 prep. 2. Adverts 10', dia 10', 10', 10'. Control cells both treated

(a) Control cell. - one of three plates - 155 = $1.55 \times 10^2 \times 10 \times 10 \times 10^4 \times 1.6 \times 10^4$ no (P), lysed, or phase spread colonies observed, nor any weak (P) colonies

(b) exptl. cell. - one of three plates - 118 on 38 partially lysed colonies observed no through 2. Colonies papulating in appearance, slightly pink toward similar to appearance of Galy band. E HFT 1 = the 2 colonies in streaking out gave many (+), and parental (-), and possibly non-parental (-). Some of the (+) appear to be segregating

285-1
285-2
Streaks

Picking (P) to examine w/ seq.	(+) colony	stability	Seq. test	Allele	285-2 HFT	Seg. test	Notes	2nd batch
285-1	1.	S. stabl	0 - +	-	u	parental		5-1229
	2.	u - unstbl	0 + +	1-	S	-		subset from Galt
	3.	S	-	-	S	-		resequenced
	4.	u	+ var 0	4-	u	parental		9 in all
	5.	u	0 +	1-	u	parental		6 were parental
	6.	u	+ 0	4-	u	parental		3 were Galt tp^s
	7.	u	0 +	1-	u	0 + tp^s		
	8.	u	0 +	1-	S	-		
	9.	S	-	-	u	parental		
	10.	S	-	-	u	"		
	11.	u	reverting to + also trans. by 4?	parental?	u	"	Many of parental show evidence of lysin- Segregating is that this is a transducing segregating	As in the case of the 1st b
	12.	u	0 +	1-	u	-		Galy \rightarrow x Galy
	13.	S	-	-	S	-		where the transduction looks also tp^s and segregating tp^s homotype
	14.	S	-	-	u	parental		no heterotype
	15.	S	-	-	S	-		recovered
	16.	u	0 +	1-	S	-		Crossing over apparently
	17.	S	-	-	u	parental		little formed
	18.	S	-	-	u	-		with
	19.	S	-	-	S	-		
	20.	u	0 +	1-	u	parental		
	21.	S	-	-	u	-		
	22.	u	0 +	1-	S	-		
	23.	u	0 +	1-	u	parental		
	24.	u	0 +	1-	u	-		
	11 stabl			10 1-	8 stabl	1 1+?		
	13 unstbl			2 4-	16 unstbl	15 parental		
				1 parental?				

From the above information it appears that these (-) seg. are not derived from separate crosses

7/22/54

Linearity of G_{ad}^- in G_{ad}^- and G_{ad}^- ?

3/17/54
902A

Δ	Δ	Δ	Δ
0.025	62	60	357
0.05	81	77	415
0.075	206	204	725
0.100	243	241	1392
0.125	157	155	1734
0.150	130	128	1566

From the above 12 plaques picked - to observe if trans. of tp^+ are by tp^+ particles
 examine hard for lysogenicity and also try to confirm the mechanism
 of reduced lysogenicity with in G_{ad}^- trans. Examine say the use of tp^+ genes
 are tp^+ or tp^+ . Also try one of each to see behavior of + with. Make
 check.

286-1 = 750E902 | of the 12 picked - after 2 shh. $1/12$ with (+)
 286-2 = FU E902 $6/12$.. (+)

1. In both cases - no clear distinction as regards lysogenicity response
 between transduction and segregants. In cases of tp^+ segregants observed.
 On the assumption that tp^+ particles are transducing tp^+ cells, perhaps the
 heterotypic segregants should be tp^+ . However previous evidence
 indicates that such segregants need not be tp^+ , since lysates were
 made of them - see FU E 902

286-1 titer against S18 10^7 ph = 1 plaque = c. 1×10^7

Assay

	10^2	10^4	10^6	ph
750 (cont. 10)	44	18	0	2.8×10^6
2175 (cont. 11)	63	11	13	6.3×10^4

750E902

286-2 titer against S18 10^7 ph = 73 = 7.3×10^8

Assay

	10^2	10^4	10^6	ph
750 (cont. 10)	44	251	2	2.5×10^7
871 (cont. 11)	276	13	14	2.8×10^5

871E902

Assay with

Popular plate

Assay of G_{21} -
in G_{21} and
 G_{21}

G_{21} WBU

G_{21}
new
10X
new

G_{21} WBU

ml of lysate plated

13500

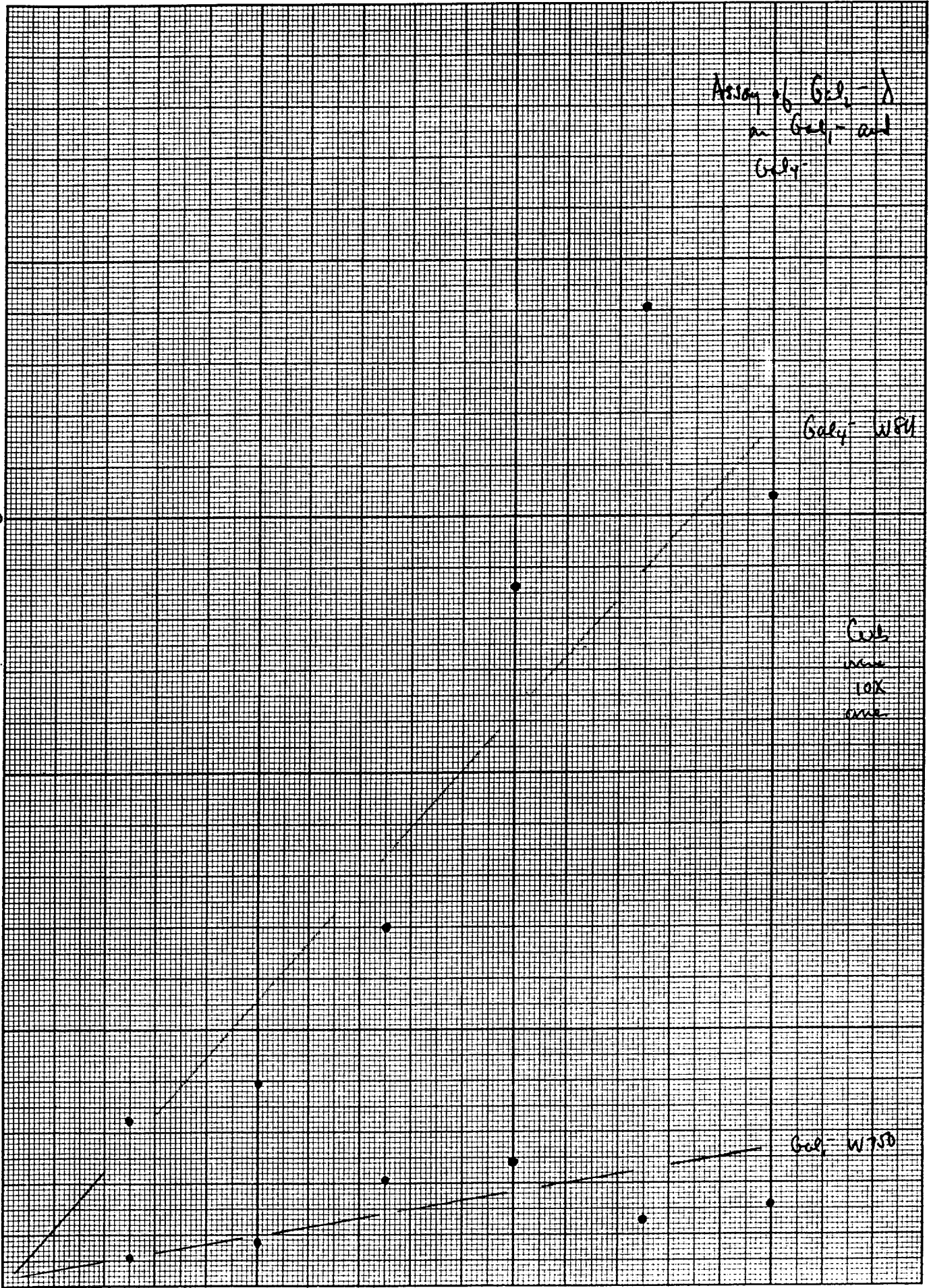
1000

500

0.05

0.1

0.15



9/9/54

2341 Its constitution. Isolated on ~~1924~~ ^{Hfr} gel - $4p^R$ - T₁N isolated by $4p^R$ against $4p^S$. Is. run on indicator of $4p^R/4p^S$?

- ① Shook out and single wt tested.
 - $9p^R$
 - $1p^S$
- ② One $4p^R$ and ten $4p^S$ grown up and plates B gel to obtain reassess.
- ③ Reassess (obtained from ~~same~~ ^{same} ~~source~~ ^{source})
 - ④ $4p^R$ - 8/8 appear to be neg ± on streaking
 - ⑤ $4p^S$ - 6/6 " " " stable on streaking

∴ 2341 is $\frac{2-4p^S}{2-4p^R}$ coming from the following Rx

$$\frac{2-4p^R}{2-4p^R} \times \frac{+p^S}{+p^S} \rightarrow \frac{+S}{-R} \rightarrow \frac{-S}{-R}$$

9/10/54 7 ml of 2341 stock in stock
Labelled as 2341R

9/12/54 lysate of 2341 $4p^R$ (made in with an $4p^+$)
analy against SR
no plaques
no transductions

9/13/54

1924EK₁₂ - Recheck and cross ϵ 1436 to see if fragment is $\pm R$ by reversion? $\frac{-S}{+R}$

① possible trans. obtained - all non lyogenic / SR
② crossed ϵ 1436 - both cultures not aerated. B gel.
3 plates -

$\frac{1}{8}$ of plate = $\frac{106(+) / 39(-)}$

The (-) 23 picked ^{shook out more} and tested (2 may be ~~SR~~ ^{SR}) - 21 $4p^S$ 2 $4p^R$

The (+) 24 picked streaked out 2^o, (-) neg (tested) = 24 (3 appear extra mixed ^{after} rechecked _{mixed})
21 were SR

stock plate of this
saved for further

9/20/54

2341 - 2344

2344 is a V_1^R derivative of 2341. Since 2341 is $\frac{2-L_1^J}{2-L_1^C}$ what is 2344 with regard to V_1 ?
① taking colonies of 2344 streaked out

17 colonies tested / d - all L_1^J . Suggestion that only L_1^J seg. of 2341 can be made V_1^R since according to Caselli suggestion crossing over between fragment and chromosomal V_1 results in duplicate for V_1 and L_1^J ?
② Plate 2341 / T7 see if any $L_1^J V_1^R$ start.

③ ~~24~~ V_1^R colonies picked, purified once and tested against 1) 10 were mixed and phage reaction suspect
7 lambda Sars
20 L_1^J

Better repeat with reported Gel no as to judge segregation

581 - Metamix HFT by adding fragment?
578 (9/16/54) -

the run with 578 opposed with HFT 4 - 48 tested found NFT
against 2175 -

84 (9/20/54) - 24 colonies tested against 2175 - all NFT
sample of 18 NFT
run 2
9/30/54 20
1210 - all NFT

9/18/54

518

To make HFT blocks of other gal

289-1 ← U2870

289-1 →

① 90 against EM's 2238 d - of 20 py picked (very ^{small} py in exposed portion of plate) found segregating

10 segregants tested all transduced by HFT 4, must be gal⁻?
lycote made - tested for transd. / 2297, 811, 0.12 of gal, 106

② 2070 d - m + a found } 24 tested

84	25	73	33	48 x 106
2297	223	313	251	9.0 x 106

9/23/54

2350 - stock to see contaminated to reverse in

10 colonies tested against 2 & 4 HFT

} H make gal⁻ - HFT by segregation from double ducker

① after 16 hours in transduction:

② " 72 " transductions by HFT 2 - of 4+ reversions?

This behavior is consistent with previous.

9/23/54.

Recheck on probability that virus in goats give more stable reproduction than wild type as typified by K-12

1. The hands down:

		lysate	Control	0.1 lysate	Stability
290-2	750	750g ⁺ -1	0	384	21/24 +u
291-1		2175g ⁺ -1	0	80	22/27 +u
292-3	2175	750g ⁺ -1	3	540	21/24 +u
293-4		2175g ⁺ -1	3	93	23/27 +u

W's
out
stock
out 5/12/55

Storks 290-178

1210 - hand. E 2070.1 to obtain $\frac{2-6^+}{2-6-}$ 0
 290-5 1. 1210 lysate Cont 1 0.1 ml 17 9/12 +u

Single day clone of chet- lysate prepared

7 seq from seq hand. tested against HFT 2 - see wt hand. F 2)

1210 E 84 for Carvill Sept
 292-A Culture 81K 3
 62

Saved 5/13/55

[2, 3 cultures made (3/11/55)] 291A1, 291A2, 291A3

1. 291A-10 8 day tested by uv. irradiation on 811 - all NFT
 8th spotted on 8th for all of reversions stable
 ① 12 cultures taken on 8th HFT - none HFT
 ② c. 8 found stable
 ③ c. 5 reversions found stable
2. 291A-2 ① 10 day as 291A-1. NFT, spotted for reversions
 ② 12 additional tested via HFT - none HFT
 ③ c. 8 found stable
 ④ 4 reversions found stable
3. 291A-3 ① 8 day as 291A-1. NFT, spotted for reversions
 ② c. 8 found stable
 ③ 4 reversions found stable

HFT strains 2-
 202-16 } local found stable
 241-14
 241-19

Effect of λ-2 Resistance on adsorption

811 vs. 1439

1. Sept. 0.5 ml K12 λ (7/27/54 Pop 1) + 0.5 ml 10x conc. cell from non-adsorbing cell.

2. 1 prep tube 1578

3. Adsorb. cell tubes
 10^7 dil. = $\frac{113,66}{66,14} = 41 \times 10^7 = 4.1 \times 10^8$ (with recovery plaque/ml)

4. Adsorb. tubes contains
 811 10^7 dil = 927,793 = $860 \times 10^7 = 8.6 \times 10^9$ cells/ml
 1439 10^7 dil = 961,967 = $964 \times 10^7 = 9.6 \times 10^9$ cells/ml

811 4.3×10^9 cells + 2.1×10^8 λ
 1439 4.8×10^9 cells + 2.1×10^8 λ

5. Array after 15 min adsorption: 370, 10' centrifugation

811 10^7 dil = 0,1 = 0.05×10^8 λ
 1439 10^7 dil = 85,56 = 70×10^5 λ
 est. $\frac{5}{270}$ λ remaining

w2888 (292.3 ready?)

10/10/54

① Reexamination of 257C-6 ^{4-2+S} its segregant. to study crossing over - 29 colonies picked and checked out from original streaking of stock. Single (+) seg selected.

	HFT 2	HFT 4	4 ⁺	Genotype	2 ⁻	4 ⁻	4 ⁺	Genotype
292-1	1. 0	+	R	2-R 16	0	+	R	2-R
	2. +	0	S	4-S 17	0	+	R	2-R
	3. 0	+	S	2-S 5(4)18	0	+	S	2-S
	4. 0	+	R	2-R 19	0	+	R	2-R
292-3	5. 0	+	R	2-R 20	0	+	R	2-R
	6. +	0	S	4-S 5(1)21	0	+	R	2-R
C10- 4p5	7. 0	+	S	2-S 5(0)22	0	+	partial S	2-seg? 292-22
	8. 0	+	R	2-R				
	9. 0	+	S	2-S				
	10. 0	+	R	2-R				
	11. 0	+	R	2-R				
	12. 0	+	R	2-R				
	13. 0	+	R	2-R				
	14. 0	+	R	2-R				
	15. 0	+	R	2-R				

Soliotype 4-S = 2 ^{one} 1/2 + S161
 Attotype 2-R = 15 1/2 + S161
 Attotype 2-S = 4 1/2 + S161
 Attotype seg? 2-? = 1 ? not done

CONTINUED 292A

See also 29F

② Transduction of 4^r - 1924 x gal²⁻ → gal²⁻ 4^r Do this culture heterogenic ~~meaning~~ in fact v. 4^r? Plated out on B gal and reversion examined. 12 picked

- 6 obviously slow (+)
- 6 fast (+) of race 1 appear to be seg.

292-3

3. The seg done of 2 above - 3 col examined for *Samboda* reaction

- seg +
- pure +
- pure (-)

2279 All non lysogenic

Attempt to isolate $\frac{2-4^r}{2-4^r}$? and see character of phage

③ 1503 ^{128.5} ~~128.5~~ - Inoculation of gal 4 - After c. 100 colonies examined from single colony experiment (cells + HFT 4 plated out) - failure to find (-) spotting on B(0) of cells and HFT - After about 30 streakings out - 2 (-) observed along a segregating (+), perhaps $\frac{+}{-}$ stock of (+) made ~~(-)~~ contained

~~Fact of the seg from 292-1~~ 292-1
 transduced by HFT 1, 2, 4 - seg (-) is spontaneous gal²⁻
 N16 11/53 292-7

292-1, 22

292-1 - The original part of this in association with the trans. test was pure (-), no suggestion of papillates, (+)

- on streaking out for colony test, to see if $4^p/4^p$ segregating - streak found mixed (+), (-). Pure (-) colonies tested against Δ and 1 (+)

- ① The (+) - weakly sensitive (segregating probably $4^p/4^p$)
- ② 1, 3, 10 (-) colonies sensitive $4^p/4^p$ #6 (+), #7 from beginning
- ③ The above (-) spotted ~~against~~ on Gal to check reversin stable

6 reversin obtained: 2
Probably unstable when may be streaks incubated too long

292-22 - The case obviously segregating $4^p/4^p$

① 10 (-) colonies tested 7 out of 10 4^p #1, 2, 5, 6, 7, 9, 10

② The above (-) spotted to check reversin stability.

Reversin found seg = #8, #3, #4
Am. seg = #1, 5, 6, 9, 10
no reversin obtained in others (2, 7)

Apparently two cultures of 292A-1503 retained and pure of the trans. gold of 292 original = W2733

2734 ETL released a better growing isolate for this and labeled 2734

292A-1503

1503 In search of gold Het 1503

From the offspring of HET 4 or 1503

9-22/54 HET 1, 2, 41 apparently sensitive to Δ (less or than usually found) a (-) obtained not trans. ϵ

Cross with 2279 - Gal, 4^p to see 1. non-gol, 2. diploid obtained

- A. cultures on B Gal. after 3 days both plus occurrence of reversin (papillates)
- B. The cross of Gal.

	(+)	(-)
1.	12	121
2.	276	57
3.	7	81
4.	17	127
5.	4	56
6.	23	160
	69	592

Actual no (+) may be higher since small colonies appear to be unformed in appearance and may be immature (+), counted here as (-).

$$\frac{592}{514} = 11.4\% (+)$$

12(+) obtained and found stable for Gal

10/17/54

Search for HFT 4 (After many unsuccessful attempts)

2175 ECU - trans. plate = 40% (old lysate)

of 15 segregants tested (transd. saved also to see if only certain transd. give rise to HFT, as distinct from a rare segregational event ~~found~~ from any transduction - two found to be HFT in 2175 stocks mixed and also streaked out for purification (results could also be certain E (+))

Stocks: 293-11 = transd., 293-11S = HFT 4 }
 293-13 = " , 293-13S = " " }

also found 293-12 - lysates prepared 12, 13 (from lysates once cultures were vacated 56 hours before inoculation),
 - unutilized lysate streaked against 2175 ^{lyse 12} ¹³
 811 0 0 ??

SEE 365

11/27/54 293-1, 2, 3 all transductions from which HFT seg were obtained
 Behavior of transductions in giving HFT or NFT - Separate seg. from seg. (1) ↓

Seg. #	293-1	293-2	293-3	293-11	293-12	293-13	293-12A
1	0	0	0	0	0	0	0
2	0	1	0	0	0	0	0
3	2	0	0	0	0	0	0
4	0	1	0	0	0	0	0
5	0	0	0	0	0	0	0
6	0	3	0	0	0	0	0
7	0	0	0	0	0	0	0
8	0	1	0	0	0	0	0
9	0	1	0	0	0	0	0
10	0	0	0	0	0	0	0
11	0	0	0	0	0	0	0
12	0	0	0	0	0	0	0
	1 HFT 2	2 HFT 2		1 HFT 2 (pur plate)	7 HFT 4	3? HFT 4	293-12A
	SS = solid smear						substituted for W12431 as HFT 4 stock
	293-1A	293-2A	293-11A	293-12A-G			80% mixed and streaked out and found HFT on 12/15 and all 8 lysates
Repeat only	1	1	0	0	0	0	
	2	5	3	1	11	0	
	3	4	2	2	6	0	
	4	6	3	2	640	0	
	5	SS	0	1	3	1	
	6	4	1	1	0	1	
	7	SS	0	2	630	0	
	8	4	0	0	2	3	
	9	9	2	SS	0	2	
	10	7	4	0	5.50	0	
	11	3	1	0	1	1	
	12			0	12		