

5/22/53.

- A SW666 x SW1049 i / plates no swarms. vth. 5/27 6/9
- B " " " / b tubes →  $\frac{1}{2} i: = \dots = ? a$
- C " " 1,2 plates:  $\frac{1}{3} i: = \dots$  } 16 (pl. x c i on 1, 2)
- D " " " / b tubes  $\frac{1}{2} i: = \dots$  } 10 ++
- E SW967 x ~~SW1049~~ / gmtubes / SW1031 a:b  $\frac{1}{2} i: = \dots$  } 3 ++
- F " " x ~~SW1049~~ / gmtubes / SW1031 a:b  $\frac{1}{2} i: = \dots$  } 4 ++, dnx i, b.
- G SW1053 a:c x 15C (abny ene) / abc → enx  $\frac{1}{2} i: = \dots$  } 10 ++
- H " " c:a x 15C " " / abc → enx  $\frac{1}{2} i: = \dots$  } see below
- K SW1053 a x 666 / b  $\frac{1}{2} i: = \dots$  } 5/26 6/5
- L " a x 967 plates 1 sw, gm: → gm / gm  $\frac{1}{2} i: = \dots$  } a: c?
- M " c x 666 / b  $\frac{1}{2} i: = \dots$  } c: → b?
- N " c x 967 plates 1 sw, gm: → gm / gm  $\frac{1}{2} i: = \dots$  } Rev annot.

Note: in above experiments, SW967 nurseries (and transdominants) were very poor. SW666x yields were a few swarms/plate (.1/ml)

	isolates	a	b 5/24	6/3	c	d
G-H	1 enx	a	enx/a enx	48h →	a/a	a/enx.a
G	2 enx	a	wh. enx	18h →	enx	c? 3 permot.
	3 enx	a	magg. (off not)	magg.	enx	magg.
H	1 enx	c	enx/c enx		c/c	c/c enx
	2 enx	c	- enx		enx	c (wh.)
	3 enx	c	magg.		enx	enx
			magg. → magg.		enx	magg. <del>no</del>
						no swarms

save some maggotable derivatives.

No evidence of a:c (:enx) variation.

Perhaps 1053 insufficiently variable to start. Review H<sub>1</sub>:H<sub>1</sub> available

Re-oid after mot; s.e.i.

(over)

Q SW1054 (Lwoffate) enx → TM2 1. ph1 still i  
 2. ph2 enx :

R " 1055 " " → 1. ph1 still i  
 2. ph2 c: 1,2

SW105

After re-motility (second passage)

Q2 still enx+++ , immobile in enx screens.  
 However, reacts slowly + lightly i: i, not b, a, 1, 2  
 Compare SW986. (Edwards calls this i: enx)  
 1041-7

1041-7 (abony x TM1 enx: i) moved over night through enx  
 Q2 - immobile !! smatic?

WALL PAPER  
 Spot photo

1051C	1051M
1	12
2	13
6	15
9	16
11	17
12	19
16	20
18	21
19	22
20	23
	24
	25
	26
	27
	28
	29
	30

6/18/53.

Q SW1054 enx → SW1046 / i: 1, 2 3 tubes  
 R 1055 " " " "

B3. (4, 5 no swim) → still i+1, 2.

R3 ( " " ) → enx or rough. S.O. file southwest along and retreat  
 T.O. other (no swim) 6/26.

Q2 repeated had not swam in > 1 week. T.O.

→ enx+++ i++ a-c-. Remotely and pass through enx, i...

single column i; enx esp. check as photos from these.

Thus SW1055 is (at least)  $H_1^c H_2^{enx}$  and SW1054 is enclosed tentatively as  $(H_1^a) H_2^{enx}$

Rechecked: R3 s.c. (by side appl.)

R3	i	enx	b	-
	++	-		
	2	++		
	++	+++		
Q2: not	++	+++		
stake	++	+++		

∴ confusion of phases may again be involved. (cf SW 986)

R30 moved rapidly through i, largely blocked in enx. T.O. in view of evidence of mixture. enx series may contain secondary applications: wait for shipment of numerosa series

of SW1061!

5/30/53

31042

Received from Edwards + Moran:  
ca 24 hours swarming.

+ = 1cm and density

P.A.E.

A	WH2	vacc.	147	++	+
B	Perm	818	← SW1058	++	
C	"	837		+	
D	"	830		+	

A.M.

E	1967			++	
F	1255				
G	715B			+	+
H	Petes 7-404			+	+
I	Thames			±	
J	McLellan			+	
K	M66			++	
L	H1			+	+
M	1966			++	
N	CoE			-	
O	Schofield			-	
P	763B			±	
Q	C			+	
R	D2			±	
S	Perm 2			+	

curf : b w. not ab-equi  
" " "

also save

T	WH1 Rough
U	Platt "
V	PM764R
W	D7 Rough
X	D "
Y	Bonnetini 556 "

∴ choose among A, E, K, M. Melt off and re-inoculate; streak out.

In second run:

	"Smoothness"	Colonies NA	14.12 14.07	Aggl. (curf)
B	Hot tube	±	+	++
E		±	+	++
K		+	+	++
M		++	++	++
N	±	++	+++	++
O	-	-	SIC.	-

(726 AM) N - m plates } microscopically almost immobile  
O - " } (occasional spinner)

Use M particularly. s.c. after mat = SW 1058

22 x 0 dense slow swarms; fast: immobile  
(no trails seen initially)

still slow after 24-36 hours

Hard O are essentially  
O-forms (but why do not  
occ. motile cells swarm?  
Temp?

Fixm

(over)



PA22 x 0 20 swarms: all exp

No swarms on controls (2 plates, 48 hours)

Purify 4 for diphasicity test. <sup>exp</sup> → Absolutely immobile  
to 6/9/53.

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N gave one. late, very rough, slow swarms spent  
+ PA22 → 10.

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M taken as SW1056 for later study.

However, in exp serum → b! Repeat with  
single colony resolution! (and verify somatic antigen)  
PRE

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In recheck, of Edwards' cultures, B = Perm 818  
seemed best. After motility test, use as SW1058

SW1056b in b serum 1. → —, later rough  
2 → z33.

∴ exp → b: —

iva. (N97)

1. 1046 various single clay isolates for consistency in extent of variation: of B1-B38.

But s.c.i from B1, B2 (which had gone b:1,2:- and b:1,2;b: - respectively) → 1,2 all: - (1? b233).

46 FG. Some misinterpretable phases. B2-1 is only example of b:1,2:b in this series.

of 1046 B1 = SW1043 and SW1007; former is b:1,2:(-)  
latter is b:-

Suggests possibility of "exhaustion" of variability.

46 C : FA 22 → x SW1043 → 2 :  $\begin{matrix} i:1,2:i \\ i:1,2:- \end{matrix}$  SW1049

DE SW1031 b → x TM 1046 → 2 cases  $\left. \begin{matrix} b:1,2:b \\ b:1,2:b \\ a:1,2:a \\ a:1,2:a \end{matrix} \right\}$  not tested further

a → "

∴ SW1031 is interpreted as  $H_1^a H_1^b$

JK. Fr. hypothesis or homology test

J. abny<sup>2</sup> → x SW1049 ( $H_1^i H_2^- H_1^{1,2}$ )  $\leftarrow$  2 b:1,2 not test

K " → x SW 1042 B2.2 ( $H_1^b H_1^{1,2}$ ) → K1 enx:1,2:enx

not tested in 1,2:enx

Misc. tests on S. wren; ~~del-er-schem~~; estimates measurement.

A-B FA3 (attendaf c)  $\rightarrow$  x SW1031 a:b

SW1052 c:b: not test.

$\rightarrow$  SW1053 c:a: ht

49 G abny <sup>enx</sup>  $\rightarrow$  x SW1049 / i; b; 1,2  
see 465

$(H_1^i H_1^{1,2} H_2^-) \rightarrow H_1^{1,2} H_2^{enx}$

5 isolated enx : 1,2 : enx

Either phase in 1,2+enx  $\rightarrow$  either enx, or magghet.

Some are still being rechecked for i. enx: c: enx <sup>sw</sup> 1054

1 51. G-H. abny <sup>enx</sup>  $\rightarrow$  x SW.

1054-3  $\rightarrow$  G-1-3  
 $\rightarrow$  H-1-3

enx: a: enx <sup>sw</sup> 1055

Efforts to demonstrate c: a: enx, a: c: enx resp.  
in c+enx, a+enx have given inadmissible forms. cf. 498.

Q SW1054  $\rightarrow$  x TM2 ph2  $\rightarrow$  enx: monoghamic!

1055  $\rightarrow$  x  
(c: enx)

$\rightarrow$  c: 1,2

$\therefore$  While H, H, structure of java seems to be justified by SW1052-3 (c:b c:a), the final proof of a triphasic is not settled. Homologies of (enx  $\rightarrow$  x H, H, )  $\rightarrow$  still to be settled. (Q-R large scale).

(over)

A more readily variable example of  $H_1, H_2$  would be desirable.  
(SW1031 is perhaps the most distinctive).

Tests 1050 perhaps should be repeated. Try on a:enx.

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heritage  $H_1 - H_2 \rightarrow P/a_1^-$

1051 Tests very limited  $\therefore$  Total actually only 5.

SW1049 ( $i:1,2 H_1, H_2, 1,2$ )  $\rightarrow$  SW666. 2  $i$ : -

ph)1,2  $\rightarrow$  : b: 26 1,2: 7

SW1031  $\rightarrow$  SW967  
1053  $\rightarrow$  SW967 ) note: very low yields +  
apparent lysis why?

1053<sub>a</sub>  $\rightarrow$  SW666 a: - q: -

1053<sub>c</sub>  $\rightarrow$  " c: -

abutus-egypti and paratyphi A.

1052. Screening of Moran strains leaves doubts on several of these concerning identity, as some are  
ex: b.D E (1967 - relation to 1966) not yet  
tested.

Fix on 1052B (Edwards) as authentic strain.

1045. Several attempts  $\rightarrow$  a: - unsuccessful.

However, SW1048 seems more transducible (fr  $F/a^+$ )  
by FA(PB), though not by TM.

Note: F.R. records SW1048 as I<sup>-</sup>, 948 as I<sup>+</sup>.

(104551)

SW1047 as I<sup>+</sup> SW694 as I<sup>-</sup>.  
Transduction?

Perhaps should check other transductions  $\rightarrow$  1048 for restoration of I.

6/3/53 (2 of E4L)  
 W2281x H245

40 colonies streaked out., replicate to pick 1 apparent Gal<sup>-</sup> to EMS  
 Gal. Replicate to EMBS Gal, Lac, Mal, MH.

Pick 2 as mostly likely still Mal<sup>+</sup>, lac<sup>+</sup>.  
 of original 40, all were lac<sup>+</sup> (v) exc. 2, 8, 32, 40.

	lac	Mal	MH	Gal	
4	v	+v	+v	+v	} not - . not Gal <sup>-</sup> .
7	v	v	v	+v	
14	v	-v	+v	+v	
23	-v	v	v	+v, -?	
27	v, v	v, v	v	+v, -?	
28	v (+/-)	+v, -?	+v	+v, -?	
29	v	v	v	+v	
39	-v	+	+	+v	
	-v	+	+	+	

14, 23, 27 must be ~~rechecked~~ rec to verify whether Gal<sup>-</sup> or Gal<sup>+</sup>  
 (latter is assumed Gal<sup>+</sup> or +, modified by other reorganizations)

Esther later isolated H324 from this cross. T.O.  
 This material

6/6/53.

by PDS - W2284 from W1895 / motility passage.  
 kirk's hands, F<sup>-</sup> and not re-infected by W1817 or by W1895.  
 (However, controls not certain).

1. W2284, W1802 grown with W1305 3 hours. Mixture (F)  
 then streaked out (→ F1) and plated directly with W677, W1896:

		D/O 48h.	EMSlac
1	W1802 x W677	-	-
2	W1802 1896	++	++ +? -
3	1802 F 677	2	3 -
4	" 1896	+	
5	2284 x 677	-	-
6	" 1896	++	++ +? -
7	2284 F 677	+±	+±
8	" 1896	+	->+
9	W-6 677	+	+ ->
10	" 1896	+	

This mixture of W2284 (or W1802) with W1305 is clearly F<sup>+</sup>

F1. Isolate Lact<sup>+</sup> from initial mixtures with W1305. (ca 10-20 colonies from  
 EMSlac pooled in broth + used after incubation overnight).

W1802' x 677	+
W2284' x 677	4

Restrict on EMSlac.  
 lact<sup>+</sup> check purity  
 2284 F1 - 2 pure Lact<sup>+</sup>  
 1802 F1 - 2 some? muc.  
 May have some  
 Lact<sup>-</sup>

F2. Lact<sup>+</sup> from mixture overnight. Pool incubated 4-6 hours.

1802' x 677	+
2284' x 677	+

Re-purify and  
 check single colonies  
 of 2284 F2

(over)

55A

W2284 + W1941 in both overnight.

S.O. EMBlac ca = isolate, test, rep. lact

sterile x W677

B. to W1802. Came out ca 100:1 - :+, rep. lact and test as above.

Repeat 2284 ... F x 1956 re-purified (pool still)

1802	x 1956	—
1802F1	"	++
1802F2	"	+++
2284F1	"	—
2284F2	"	+(few)

maybe utter sterile or unstable

Try passing back to W677.

10552:  
6/10/53.

2284 x  
x  
2057

prot. test as Hfr, F<sup>-</sup> ...

4 Malt<sup>+</sup>  
3 Malt<sup>-</sup> test by SRP

On retest 6/14/53

2284F2 pool

S.C. 1  
S.C. 2  
S.C. 3  
W6

W677  
—  
—  
—  
++

W1896  
++

∴ pool has become F<sup>-</sup>!  
Test for transfer to W677!



6/19/53...

H. W2284 "F2" + W1956 overnight, 20. in EMS lac sm to recover "exposed" 1956 as lac-S<sup>R</sup>. No pool to Livenessay.

I. (W2284F2 + W1305) x W1956 10 prototrophs  
2284F2 " 0.  
∴ 2284F2 does not become grossly infective after losing F<sup>+</sup> quality.

JK J (W2061 (MTL-Hfr) + W1956) x W1802 40 prototrophs

K. (W1305 MTL-Hfr + W1956) x " 80 "

J. Transfer from Hfr? Temporary or mutable? or recombination.  
lac<sup>+</sup> v<sup>R</sup> Mal<sup>+</sup> S<sup>3</sup> lac<sup>-</sup> v<sup>R</sup> Mal<sup>-</sup> x lac<sup>+</sup>....

no other matrices at hand.

Repeat: (W2061 + W1607) x W1956  
lac<sup>+</sup> S<sup>3</sup> (lac<sup>-</sup> Gal<sup>-</sup>)<sub>SR</sub> lac<sup>-</sup> S<sup>R</sup>

or (2061 + 1956) x 1607!

Reinoculate K.

L. (W1956 + W2061) x W1607 D/O heavy sweep → v. low yields.  
Test all 8 prototrophs in EMS lac.

M. (W2284F2 pool + W1956) W1956 recovered, + W1607 no prot.  
see infer wOP<sup>+</sup> in 2284F

Controls W2061 x 1956 } no prototrophs  
1956 x 1607 }

H245...

6/9/53

A. H245 from S. Z'om EM (lac in Penassay (TLB, -)

x 2. gave ca 50 prototrophs mostly lac+ v strds: H245 ca 1% lacv.

6/9/53 B. acrated in D(lac, HC) } 1. x W1321 H-F S^A Lac^+ Gal^+ lambda^S  
C. standing in " " } 2. x W1486 F+ " " "

B1 no prot. - -

v strds: B 90% v. Rare +  
C 70% v 1% +

B2 ++ -> high prot. no -

C1 ca 10 prot. Lac - prot. on EMS Gal. all Gal -

C2 + < lac+ to EMS Gal  
lac- to " "

This stock therefore seems to behave as F-, especially when acrated.

Lac+ should be uniformly lacv. C2

Lac Gal		Lac Gal	
1 V	+V?	17 V	V+V?
2 V	V?	18 V	+V
3 V	+V?	19 V	+V
4 V	+V	20 V	+V
5 V	V H327	21 V	+V
6 V	+V	22 V	+V
7 -	+	23	(over)
8 V	+V	24	
9 V	+V	25	
10 V	+V	26	
11 V	+V	27	
12 V	+V	28	
13 V	+V	29	
14 V	+V	30	
15 V	+V	31 -	V+V?
16 V	+V	32 -	-

C1. 3 Gal? 9 Gal - lambda^S Restrict 3.  
phenotyped on 1.

C2. Restrict possible lacv or Galv.

Lac Gal	
33 -	V?
34 V	V+V?
35 V	+V
36 V	+V
37 V	+V
38 V	+V

(over)

459:	Lac	Mal	Gal	Rev single cols.	Lac Mal xyl Gal MH D(0)
1	V	-	-		V - V V +
2	V	-	-		V - V V +
3	-	whq?	-	Lp 2/3	V - V V +
4	-	-	V	#3 is Lacv Gal- (i.e. prot)	V - V V +
5	-	+	-		V - V V +

Assume 5 as significant. Save #3 as Lacv Malv Gal = Lp^2 H326  
#5 may have been uninfected or from Gal = Lp^3, or may be crossover.  
(over)

Next experiments are

- ① Hexotypic rearrangement
- ②  $lp^s lp_2^s$  rearrangement. (perhaps better from a  $Gal^+$  revision ??)
- ③  $Gal^+$  revisions (cis and trans)
- ④  $Gal^+$  transductions ....

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Most lacv from C2 are  $Gal^+$  mother. H327

Retain #5 as  $Gal^+$  lacv and check other mothers  
also check "Galv" from 1, 2, 17, 34.

H327: Lacv Galv Mal<sup>v</sup> S<sup>s</sup>  
           $lp_2^s lp^+$

(S<sup>s</sup>/S<sup>R</sup> still on hand)  
by i<sup>+</sup> suppressor  
not relevant

No individually orient factors: does this segregate  $lp^s$ ?

↳ either finds  $lp^+ / lp^-$ ,  $lp_2^R$ .

?  $lp_2$  status must be checked.

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no S<sup>R</sup> with  
heavy molecule  
in EM103 lacv

C. 1-5: #1,4 are evidently lac<sup>v</sup>; maybe useful later as Hfr IF<sup>-</sup>  
 #2,3,5 show peculiar mottling; definite evidence of segregation.  
 Could look for evidence of mutational segregation. Store in NA stubs.

		lac	Mal	
A	1	+ <sup>v</sup>	+	E1
	2	+	+	
Restreaks	3	+	+	
of parent		+ <sup>v</sup>	+ <sup>v</sup>	
s.c.i	B1	+ <sup>v</sup>	+ <sup>v</sup>	
	2	u. light + <sup>v</sup> ?	-	
from plating	3	+ <sup>v</sup> or v?	+ <sup>v</sup>	E2
on EM3lac	4	+ <sup>v</sup>	+ <sup>v</sup>	
to EM3lac.	5	+ <sup>v</sup> and -	+	E3

presumably lac<sup>v</sup>.

C	1	+ <sup>v</sup> and -	+	E4	have presumably been reisolated
	2	+ <sup>v</sup>	+ <sup>v</sup>		
	3	+ <sup>v</sup>	-	E5	
	4	+ <sup>v</sup> +, -?	+ -?		
	5	+ <sup>v</sup> -?	-		
	6	+	-	E6	
	7	+ <sup>v</sup>	-		
	8	+ <sup>v</sup> +?	-	E7	
	9	+ <sup>v</sup> +, -	-		
	10	+ <sup>v</sup> +	- occ +	E8	
	11	+ <sup>v</sup> +, -?	-		
	12	+	-	E8	
	13	+	+		
	14	+ <sup>v</sup> -?	-	E8	
	15	+ <sup>v</sup> -	+		
	16	+ <sup>v</sup> -	+	E8	
	17	+ <sup>v</sup> -?	-		

Remarkable mottled appearance of all of these. Save EMS originals protum; choose for further study:  
 A1; B5; B3; C1; C16; C15; C8; E5.

Repick single colonies and spot EM3lac, brush EM3lac / sur; streak out EM3lac again. Handle as E1-8

check 56 & 2, 3, 4

2 karu Melv bal +  
3v " " +  
" " " +



6/9/53

A. H310 (from 2.2. EMBlac to Penassay)

X W1801 EMBlac Prod. lac -

Pool plated on EMBlac, Mal. ca 5% Mal-, 70% lac-. Made possible lac, Malv for later picks. Plates may have been moist. Growth very rough & spreading.

H: mostly lacv.

B. X W1802 EMBlac. Prod. lac +

Pool: almost pure lac +

see 57E

spills very high. >> 1000/plate  
pool growth and natural for sampling.

C. H310 x W2209 (V-G-F-) Dilute plating. >> 100/plate. Strains seem weaker lac+ on EMBlac. 24 tested: 16 Mal- 1+, - 7 Mal+ Repick 4 Mal+ which might be Malv. (2 from sect+ among -; 2 Malv+). Replicate to check lac poss.

Transfer possible to EMBlac

	Mal	lac
1	+v?	+, +v?
2	+v	+ +v?
3	+ and -	+ and -
4	+	+

Replicate streak plate (24 cols - C):

lac	Mal	lac S
1 +		S
2 +		S
3 + only sect	+,-	-
4 +	(+ colony foot)	R
5 v?	-	R
6 +	-	R
19 +	-	S
20 +	+	S R
21 +	-	R
22 +	-	R
23 +	-	R S
24 +v?	+	S

lac	Mal	S
7 +	-	S R
8 +	+	S R
9 +	-	R R
10 +	-	R R
11 +	-	R S
12 +	+	S

No Malv indicated.

lac	Mal	S
13 +	-	
14 +	-	
15 v?	-	
16 +	-	
17 +	-	R R
18 +	-	R

Recheck #1, 2, 5, 15, 24 on

EMBlac, Mal, Mtl; spot on Shac.

Remember C1-5.

Some lac+ might be v.

C: Repeat: 20 tested streaked in Blac; have 12. = 6-17

lac	Mal	Mtl
1 v	+v	v?
2 +v	+v	+v
3 +	-	-
4 v	+v	+v
5 +v	-	+v

D H310 x W194/1 v. heavy yield. Numerous spotted lac - TR!

7/8/53. Replate H328:  
all colonies prototrophic, including  
c100 lact, few lact - segregants  
on two good plates.

SRP crosses: 1057E (J.L.)

Penassay (7.5 ml) cultures of the following were made and grown up overnight:  
1817                      57E 1, 2, 3, 4, 7  
1177

Half a ml each of the 1057E and 1817 or 1177 were mixed in Penassay (7.5 ml), incubated 8 hrs, centrifuged, washed twice in centrifuge, resuspended in 0.5 ml water, diluted on 5 lac sm<sup>7</sup>/17153

	X 1817	1177	control	S.O. on 3 lac
57E1	lact±	lac-	—	lact+ (light + dark)
E2	lact, few lac-	lac-, few lact+	—	lact+ (light + dark)
E3	lact±	lac-, few, against solid-background	—	lact+, lac- (fewer-)
E4	lact, few lac-	lac-, few lact+	—	↓
E7	lact±	lac-	many cols, all lac-	
control	—	—		

E7 unreliable; E1-4 evidently Hfr. Test lac - signants.



6/16/53.

		Restricta s.c. EMBlac	MH from rough	Mal (from rough)	SM
x1801	1	+ (rough) only	+	+	S
x1802	2	+, sectored or mottled +, and occasional -	(similar to H310)	+	S
x1802	3	" v. rare -	-	+	S
x2209 =1678F-4	4	" occ. -	+	+	S
H328 ←	5	" , 1 definitely sectored +/-	+	-	R
	6	app. pure +	-	-	R
	7	+ , s, m, occ. -	-	+	S
	8	pure mottled +.	+	+	S

all prototrophic on EMS lac

dependence of R/S.

#2,3 2n from H310; Mal, S from F<sup>-</sup>

4,7, 2n from H310; Mal, S from F<sup>+</sup>

5 2n from H310, Mal S from H310 (TL<sup>+</sup> from F<sup>-</sup>)  
 mutations; hemizygosity. H328.

Replicate each to D/O for nutritional segregation.

No auxotrophs noted in direct replicates (very few segregants checked)  
 Are these diploid or unstable lac?? (H310 itself?)

5 Mal<sup>+</sup> from H328. Struck on EMBlac, Mal. of H328.

All appear to be lac<sup>-</sup> Mal<sup>+</sup> ( ). Suggests H310 itself is Mal<sup>-</sup>.

Attempts at H310<sup>+</sup>: plate on EMS Mal<sup>+</sup> 100 hydrof. casein: ca 20 papillae after 3-4 days, but these are not Mal<sup>+</sup>!

Repeat - now smaller papillae: all Mal<sup>-</sup> -! Replicate papillae (over)

Proof that H310 is diploid; lac is only segregating marker!  
(pure TLB, -!)

a) Hfr 2n vs. F<sup>-</sup> 1n.

b) On a single occasion x 1895 gave Lac<sup>+</sup>/Mal<sup>+</sup> Su.  
(cf. 1059.) H313

---

for test fruits our population:

---

1057E1.-2 papillae of H310 in EMSMal HC.

↓  
Mal<sup>+</sup> Lac<sup>+</sup>. same 1057E2 = Mal<sup>+</sup> Lac<sup>+</sup>

∴ Mal hemizygous

Reduces ~~57E21~~ 57E21 as Hfr.

and segregants.

57E21 x W1607 → disequilibrium of recombinants  
parent controls =

6/10/53

A. (acrated H290) x } { 1. W1394 } mEMSlac  
 B unacrated " x } { 2. W1918 }  
 (D(lac, M))

yield  
 A1 6 colonies: 1 lac+ ; 1 lac+  
 2 10 " nonlac+ ; 20 sec.+ )  
 B1 2 -  
 2 sec -,+ ; 0.

1058A. Skelton A1 (1-5) B1 (6-9) mEMSlac

In repetition, none were lac, Malv. T.O. for now

6/9/53.

H290 from <sup>6/9/53</sup> <sub>(7/30/51)</sub> yeast: direct streaking → to EMS lac. grow in D(lac HC...)

H302:  $\Sigma O_2$  tube 1 - → growth slowly: pure lac-  
tube 2 → " " " " invariable

lyophil → heavy immediate growth, ca 50% lacv.  
→ 90% lacv ✓

H313: resp. invariable

lyophil: (3/53 → found all segregant)  
6/53 → EMS Mal: → mostly Mal-, Mal+ some V.  
EMS Mal

H226 occ. lacv from viral 9/52 → ✓ (see (lac+))

H267 Almost all lac- " " Restrict on EMS lac; back from EMS lac to D(lac) →

318 Invariable

319 Viable; rare lac+ (v. light, maybe v.)

213-14 <sup>13</sup> pure lac+ 14. Restr.

H304-5 mostly +, - T.O.

W1940 x 1956  
Hfr F<sup>-</sup>

1059

6/11/53.

A mEMSlac

B lac together in broth; plate EMBlac., EM15lac

??

A: 30 lac+ from EMS lac to EM15lac. Re-possible lac<sup>v</sup> others are + (1-). Restraints 10: all are Mal<sup>-</sup>, #14 lac<sup>v</sup>? others lac<sup>v</sup> may be enough!

B. Papillae eventually noted.

lysis in stalks?

Not scoreable for lac<sup>v</sup>. Sample  $\frac{1}{2}$  noted in each stalk, probably reversions.

None show lac segregation. Repeat expt.; also check for l.

59BB: Investigate lytic appearance in B:

59Z 6/16/53 -x- EMS, EMBlac after growing together 6 hours is aeration. T.O. review of inquiry

BB: W1940 proved to be mixed Mal<sup>+</sup> / Mal<sup>-</sup>

In plating of W1940 + W1956, Mal<sup>+</sup> and Mal<sup>-</sup> left. are  $\frac{1}{2}$   $\frac{1}{2}$  = W1940  
all lac<sup>-</sup>. Mal<sup>-</sup> normal are  $\frac{1}{2}$   $\frac{1}{2}$  = W1956.

=W 2302 isolate Mal<sup>-</sup> as presumed mutant of W1940 (This stock had been recovered from an old, discarded stock which grew out slowly). Recheck lyophil and current cultures of W1940. Reserve W1942 also for further experiments.

Renew W1940 stock from lyophil (pure Mal<sup>+</sup>) and throw out others.  
(over)

D	W 1956	x W 1940	EMB Lac
E	"	"	S Lac
F	"	W 1942	B Lac
G	"	"	S Lac

after  
growth,  
overnight

Lac<sup>+</sup> seen as papillae in streaks of D, F.

ca 1% Lac<sup>+</sup> colonies in E, G.

Hold suspicious colonies for later retest

7/4/53:

abandon in view of Cavalli's findings re Hfr - Gal linkage



$\lambda$ -2 tests on A1, 3, 5  
 EMBS/Mal D/O)  
 1 R  
 3 Mal<sup>+</sup>S Mal<sup>R</sup>  
 5 Mal<sup>+</sup>S and R?  
 (can't background  
 action) S!

$\lambda$   
 S H332  
 $\therefore$  presumably s/R  
 but Mal<sup>+</sup>/+

H324. In EML test segregation #1-7 appeared to be  
 segregating Mal<sup>+</sup>/-, but #8-55 were all Mal<sup>+</sup>  
 (including many Xyl<sup>-</sup>). Ferrous confirmed 6/17!

Check with T6: #1-10 ~~Mal<sup>+</sup>~~ were T6<sup>S</sup>. H324 T6<sup>S</sup>  
 Gal- H325 T6<sup>S</sup>/R

H324 as now available is Mal<sup>+</sup>/+.

The bulk of EML's data must pertain to this "secondary".

Except for auxotrophy, H332 is most suitable for  
 infection of  $lp^s/lp^s$  ..... Now need to obtain  $\text{Gal}^+$  sources.  
 H331 also OK; not segregating Mal



G1 H325. v. slow lac<sup>v</sup>? — neutral + test prototrophic  $\lambda^-$   $\lambda_2^R$  Gal<sup>-</sup> Mal<sup>-</sup>

H H329A 1 Hal<sup>v</sup> lac<sup>v</sup> Gal<sup>v</sup> lysogenic auxotrophic  
 2  
 3 " " "

G1 might be Gal<sup>-</sup> lac<sup>v</sup>, but is apparently  $lys^R$  and presumably unsuitable for present purpose. H1-3 might be used by crossing to Gal<sup>-</sup>  $lys^S$ .  
 On the whole Gal<sup>-</sup> (H326-331-332) seem technically most suitable.  
 Check G1 by transduction to Gal<sup>+</sup>; if unsuitable, probably unusable.

✓ H325 appears to be as given. (See 1062) Grow in D (Glu) Don't bother.

6/15/53.

- A H302 grows in D(M, Lac) x Y10  
 B " - aerated, into D(M Lac) 7 hours not aerated x Y10  
 C A aer. x W1918  
 D - not aer. x W1918

Plate on EMS lac.

A. 2 plates No prototrophs

- B 20-30 lac<sup>+</sup>/plate 1 or 2 ? lac<sup>-</sup>/plate (H-302 paralytic)  
 C 3-400 lac<sup>+</sup>/plate 1-2 ? lac<sup>-</sup>/plate (H-302 orthostrophic)  
 D very heavy background 5-6 ? prototrophs/plate.

B. Strain 72 lac<sup>+</sup> on EMS lac for lac v.

C. Hold!

Pick to EMS lac. Not all grow out.

picked as probably lac v

	Lac	Mal
1	v <sup>+</sup>	v
2	v	v
3	v	v
4	v	v
5	v	v
6	+	+
7	v	+
8	+	+
9	+	+
10	v	v
11	+	+
12	+	+
13	+	?
14	v	v
15	+	?
16	+	v
17	+	+
18	v	v
19	v	v
20	v	v
21	+	v
22	+	+
23	+	+
24	v	v
25	v	v
26	v	-

∴ in sum: 1 Mal  
 23 probable lac v  
 (not fully characterized)  
 33 lac<sup>+</sup> (22+)  
 (6 were not prototrophic)

27-56 picked as exp. lac<sup>+</sup>  
 lact: 27, 28, 29, 30, 31 32 33 34 36  
 Mal<sup>+</sup> or - + - + + + + + + + +  
 lact 37+ 39 40+ ? 41-44 46-48  
 Mal<sup>+</sup> or - + + + + + + + +  
 (lac 36? 37? 38 v 45 v 54 v  
 Mal 36 v v v v  
 lact 49-52 53, 55, 56  
 Mal<sup>+</sup> or - + + + +

Re-collect possible v and  
 shake with son, etc.

Mal v usually scored on basis  
 of +, - being present.

Elimination evidently does not occur in  
 renumber 30, 45, 54 as 6-8-11

2uP<sup>+</sup> x 1uF<sup>-</sup>

Conclude

1-23 are likely lacv. (Malv).

Re-~~organizing~~ the following

	Lac	Mal	→ smMH	<del>smMal</del>
1 ✓	+	+	+	+S
2 ✓	+	-	+	+S -R
3 ✓	-	+	+	+S -R
4 ✓	+	+	+	+S
5 ✓	+	-	-	-R
6 ✓	+	+	+	+S
7 ✓	-	+	+	+S
8 ✓	+	+	+	+S
9 ✓	+	+	+	+S
10 ✓	+	-	+	+S -R
11 ✓	-	-	-	-R
12 ✓	+	+	+	+S
13 ✓	+	+	+	+S
14 ✓	+	+	+	+S +R
15 ✓	+	+	+	+S -R
16 ✓	+	+	+	+S -R
17 ✓	+	+	+	+S
18 ✓	+	+	+	+S -R
19 ✓	+	+	+	+S +R
20 ✓	+	+	+	+S -S
21 ✓	+	+	+	+S -R
22 ✓	+	-	-	-R
23 ✓	+	-	-	-R

Restraints entire series of 23.

EMBLac, Mal

Bush or EMBMal

EMBMH

SM.

Edmispider may have segregated.

A) lacv Malv Sv MHLv number : 2, 3, 10, 15, 16, 22

B) lacv Malv Sv MHL+ " : 14, 19,

C) (Lac+ Mal+ MHL+ S<sup>s</sup>) : 4, 6, (7), 8, 9, (12), 17, 18

D) Lac+ Mal- MHL- SR : 5? 11, 20.

E) Lac+ Malv MHL+ S<sup>s</sup>? 1 20

F) lacv Malv MHL- SR 18

G) lacv Mal- MHL+ S<sup>s</sup>

Recheck: ~~for~~ for MHLv. B)

c): heptoid: find deplord D.

G) Lac.. Malv MHLv S<sup>s</sup> 21

6

2

8

3

Do H302 MHL-  
MHLv?



*Enchocoris* from retreats

Comments.

	lac	Mal	MAL	S	
1	.	v	+	S	
2	.	v	v	v	
3	.	v	v	v	
4	+	v	v	v	
5	v	v	v	v	
6	v	v	v	v	
7	v	+	v	v	
8	.	v	+	v	
9	+	+	+	v	haploid
10	.	v	v	v	
11	v	v	v	v	
12	.	v	+	v	
13	+	+	+	v	haploid
14	.	v	+	v	
15	.	+	v	v	
16	.	v	v	v	
17	+	+	+	v	haploid
18	v	v	-	v	
19	.	v	+	v	
20	.	v	+	v	
21	.	v	v	v	
22	.	v	v	v	
23	+	-	-	R	haploid

lac+ matted!

Mal+ S<sup>R</sup> / Mal- S<sup>S</sup>

some probably lac<sup>+</sup>

18 seemingly Mal<sup>-</sup>, Mal<sup>v</sup> no Mal<sup>+</sup> segregants. Retreats apparent Mal<sup>v</sup> for more material: 4 colonies gave same pattern. Most Mal<sup>-</sup> are S<sup>S</sup> Mal<sup>+</sup> probably include pure +.

Summary: 56 prototrophs. 19 ultimately diploid. (lac<sup>+</sup> or lac<sup>v</sup>)  
Most probably lac<sup>+</sup>1- rather than +-1-+.

lac <sup>v</sup>	MAL	Mal <sup>v</sup>	S <sup>v</sup>	8	2	3	5	6	10	11	16	22
+	v		S	4	1	8	12	20				
-	+		S <sup>R</sup>		haploid only (9, 13, 17, 23)							
v	+		S	1	7							
v	v		S	2	4	21						
+	v		v	2	14	19						
v	+		v	1	15							
-	v		v	1	18							

1/2/53

Recap. of types (provisional class. of 13, 15, 17, 27 as  $S^V$ ).

- A. 16 lacv (+/-+) MH-Mal-S<sup>R</sup> : 2 3 4 5 6 9 11 12 16 20 23 24 29 30  
31 38
- B. 1 lacv (++/--) V V V~~R~~ : 13
- C. 10 lacv (+/-+) V V V : 7 19 21 22 25 26 32 33 34 39
- D. 1 " MH-Malv Sv : 37
- E. 1 lacv (++/--) Mal-MH-Sv : 15
- F. 2 lacv (+/-+) Mal-MH-Sv : 17, 27

31  
 haploid: 5  
 aneuploidy: 4  
9

1 28 35 36 40  
 8 10 14 18

36 prototrophs tested.

31 diploid lacv : 29 (app.) lac+/-+  
 2 lac++/-  
 (may be incorrect).

Malv : 12  
 MHv : 11  
 Sv : 15.

#2 maybe ++/-+ #3 +/-+

Compare:

(Major types only, Mal - S).

		B		C	
1	Malv Sv	11	58	12	39
2	Mal- SR	0		16	52
3	Mal+ Ss	1	5	0	
4	Malv Ss	6	32	0	
5	Mal- Sv	0		3	9+
6	Mal+ Sv	1	5	0	

19/47

31/76

$$B = 2nF^+ \times 410$$

$$C = 2nF^- \times 11918$$

$$H302 = H-Locv MH-Mal-SR$$

Inferences:

1. In parent functions assume whether  $F^+$  or  $F^-$ .  
If  $2nF^+ \times nF^-$  depended on polarity reversal, there would at least be some Mal-SR. Split (B:C)(1:2) is extremely significant.

2. "No" Mal+S<sup>s</sup> in B, suggests that "none" of these diploids are hemizygous in this region. If elimination occurs from the  $F^+$  side, it must involve only 1 of two strands, and the eliminated one may be discriminated against in zygote formation. It is possible that elimination does not occur at all from the  $F^+$  side of n. Note high incidence of BH. (Mal/S) crossovers!

3. High incidence of C2 suggests usual elimination from  $nF^+$  side.  
? are these homo- or hemi-zygous? Need tests on these random CS.  
Nearly half are S<sup>v</sup>. Note: in B, only S<sup>s</sup> and S<sup>v</sup>; in C only S<sup>s</sup>, S<sup>v</sup>.

61C hemizygosity tests: (EM5Mal)

17: numerous Mal+ (purity???)

2: 2 papillae

3: 5 papillae

} → Malv.

15: 2 plates no Mal+ 7/10

2 plates: 3 Mal+ 7/18: → Malv.

#17 initially a few Malv. Associate lac<sup>+</sup> Mal - and  
(mutants?) test



None of this is inconsistent with pre-elimination, if it occurs. Would need a Mal/S crossover (Mal-S<sup>2</sup>) or a Mal-hemizygote in P<sub>3</sub> to substantiate it. Presumption of elimination is based on polarity differential.

Immediate requirements: hemizygosity tests in C<sub>1</sub>, C<sub>5</sub> classes.

Later: look for Mal-diploids in P<sub>3</sub>. (Should occur by crossover).

→ x Gal<sup>-</sup>

1062

6/19/53

A. NI-x H326 in EMS Gal

B. NI-x H331 " "

Several thousand est. Gal<sup>+</sup> from 1 pop. Pool and streak out on EMS Gal background ca 4-5 papillae

check for lac<sup>v</sup> Gal<sup>+</sup>: pick lac<sup>v</sup> to EMS Gal. mostly lac<sup>-</sup> but most Gal<sup>+</sup> are definitely variegated.  
2 lac<sup>v</sup> Gal<sup>+</sup> secured from A, B resp. Signs of lysis in seg. from each. Further screening needed.

C. NI-x H325A (Gal<sup>-</sup> lac<sup>v</sup>) → 40 Gal<sup>+</sup> (4<sup>+</sup>!)

of EML

6/16/53

A. SW 1058 (mot.) in cerv serum.

See 1042, 1052

Prickin tube had not moved in 4 days.

6/16 1/2  
3

7/9. All immobile! T.O.

6/17

B. FA 18 → X

1/2

C FA 22 → X

1/2

6/26. no swarms in any of above! (Try IV serum!)

6/26 SW 1058 mot (ferulant) = D and TM 2 = E.

1. mot

2. IV

3. IV V XII (Typhlo)

4. IV XXVII XII

24h.

D 1 ++  
2 + (definite inhibition)

3 ++

4 +++ stimulation? or fresher medium

E 1 24h.

2 +++

3 ++

4 +++

→ still IV XII III

Reover O2 after 48 hours. Shearist and test for transd., /env etc.

More & purification 7/1/53. /env is FA 18, FA 22

6202 / ~~pepas~~ FA 22

no mot. 7/9/53 T.O.

2/2/53

Cross-studies on EMPB tac, i stock & as received from WBC.

B. &	FA10	PB-1	PB-3a	PB Taunton	PB Dundee	TM & PRE
TM2	++	-	-	-	-	++ lytic
100DA (PB1)	++	++ seen v. fine plaques	-	-	-	-
SW730	-	-	-	-	-	- (budding stocks)
SW887A1	-	-	-	-	-	+ not lytic (napoli stocks)
SW957	±	-	-	-	-	+ inh? (0901)
Ordering: 1024E1	-	-	-	-	-	-
3	-	-	-	-	-	-
4	-	-	-	-	-	-
5	-	-	-	-	-	-
6	-	-	-	-	-	-

occ. shagreened plaques?

titer?  
should test  
mPB1

& perhaps should first be grown to higher titer. Try TM & PRE / TM2

SW887  
SW957.

PB1 / PB1.

Try other & on napoli, Dundee, etc.

Grow Taunton &, ~~etc~~ PB2 &.

Test & pups → SW666

FA22 +  
control -

TM / TM2 -  
/ 887 -  
/ 957 -

PB2 -  
PB1 -  
PB Taunton -

} Use

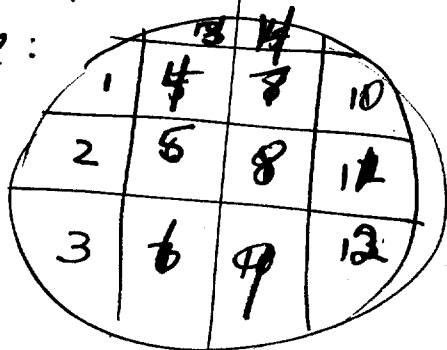
& maybe n.g.

Use PB2  
PB BAOR  
PB Dundee

2/5/53  
From results over,  
&'s have not been made to  
adequate titre.  
as basic stocks. (over)

# Test loopfuls for activity.

PB4:



B = act.

(PB1  
PB2  
666 (Jussey)  
Taunton)

For later study,  
2  
BAOK  
Dundee  
represent types of phase.

	PB1	<del>TAUNTON</del> Taunton	SW666	<del>TAUNTON</del> PB2	TM2
1. PB1 (Ch.)	++	-	-	-	
2. 2 "	<del>++</del> ++	-	-	++	
3. " Taunton	-	isol. pl.	±	- sic.	
4. " BAOK	-	-	-	-	
5. " Dundee	+	fine phages?	++ fine phages	±? v. fine	
6. FA10	++	♦♦	++	+ 0.	++ <sup>0</sup>
7. 41/1 J.L.	+ isol pl	-	-	-	
8. 42/2 J.L.	+ isol pl	-	isol pl.	fine pl.	
9. 4T/1 J.L.	-	isol. pl.	isol pl.	-	
10. Jussey (Ch.)	++	-	++ ✓	-	
11. 3a	++	-	-	-	
12. 3a1	++	-	-	-	
13. 3b	++	-	-	-	
14. Dundee	? fine pl.	-	+	fine pl.?	

Ch. pups??

Q is not prepared to adequate titer.

In first preparation, PB2, PB BAOK cleared; Dundee did not. Re-grow Dundee on 666, PB2 and Dundee

7/5-6/53

Cherry phages.	Ch:	TM2	SW688	SW666	PB1	SW730	SW887	SW957
	1.	-	-	-	±	-	-	+++ <sup>lytic</sup>
	3.	-	-	-	-	±	-	-
	4.	-	+	-	-	-	-	+ 1 pl. lytic.
	5.	-	-	-	-	-	-	-
	7.	± dead	-	-	-	-	-	-
	8.	-	-	+	-	± sh.	+	- ± 6?
	9.	-	-	+	-	+	-	-
(12)	10.	++ <sup>T</sup>	-	-	++ <sup>T</sup>	-	++ <sup>T</sup>	-
	17.	-	-	+	-	+	-	± lytic +

From point of view of survivorship, #12 seems most likely. Label on this vial reads: S. luciana #12. Cherry's letter refers to S. luciana #18 as v. specific for group XI. Presumably not this v.

Grow #12 on TM2, SW887 to test her reduction. Also grow #1 / 957; 9/SW730; #4/957; #17/666, 957.

Need to find better hosts for 3, 7: try homologous strains (cuticularis, ruber, virid and yellowium).

#1 seems to differ from 01 in host range. Called para A phage!

Storlas 6/5. Ch2 grown on TM. spot: OK. / TM ✓ v. small plaques. Titer not too high. but worth trying on 09010; SW967 etc.

m SW887 " " ✓

m 730 " " ✓

PB2 1/2 large + small plaques. Moderate #

BORR 1 ✓ small plaques. " "

Dundee 1 ✓ " " " "

Regrow for higher titer.

Ch Storlas repeat spots: Taunton, BORR, Buelbo: not discernable / PB1. May not have source.

#7 SW764 (centrifuges) - +++ (lytic) and matted survivors  
 967 (dublin) -  
 957 (0901) + (overgrown)

BAOR PB1 + shaded  
 666 ± shaded  
 730 -  
 887 ± shaded  
 957 ± shaded.

Transd. BAOR x 666 →  $\frac{b}{-}$ , Cal -  
~~= 1063 AT~~ in serum b: -

Repeat 7/8: BAOR x 666 → b  
 967 → (gm) +  
 x H901 → -

Repeat, looking for tracks on plates → x 666

Uprigs: test ster. 7/764, 1/H901 ster.  
 (checked) PB1 - under BAOR (nw) } but - x 666 possibly tracks. ++  
 7/8/53. 12/TM2, 887, P01 } not sterile .. ① melt off and test  
 4/H901 } ② chloroform & amp  
 17/H901 } retest.

7-x 666, 967, 957 all - in 24h. Titers?  
 ch2 - x 957 (ch2/TM2, 887, 730) all - in 48h.

B1 (CH12/PB1 not str. → x sw600 not) → Gal+ doubtless.  
 B2 CH12/TM2 " " " → Gal+ contamination  
 B3 " 1887 " " " → Gal+ factors in &  
 7/8. Repeated after CHCl<sub>3</sub> treatment → 666, 957, 967 all —

A. BARD → x SW666 / motility → v. numerous swarms + tracks  
 B. = FA90 / EM/ocal → ca. 100 Gal+ / .1 ml  
 Purify. controls: each 0, 0.

(A) ① Pool motile transductions (2730) all Gal-.  
 ② Individual " " : all Gal- . Pick to broth for  
 subs. lysogenicity test.

(B) 1. 20 streaked out: none unstable, all pure Gal+. Test pos. for  
 lysogenicity, motility.

∴ as before, no  
 Gal: H<sub>1</sub>, F<sub>1</sub>, linkage

(C) FA90 → x SW900. / Gal same (ca 5-10 plate).  
 / D (Meth) ± FA. no evidence of transduction).

CH2/TM2 → 666 }  
 1730 → } =  
 1887 → }

also obtain spint H<sup>+</sup> SW150.

B2 → 666 -  
 Jumboe → 666 -

(over)



lysogenicity tests on 63 A-B.

A: 1-8 all are lysogenic on BAOR; sensitive to FA10;  
SW666 not " " " " ;  
SW928 " " " " ; resistant

all are resistant to FA90.

no control, BAOR / FA90. By 90 x SW666 these have  
evidently become lysogenic for BAOR indicator. In transduction  
plateings, only rare plaques were seen. Save A1, B1 for  
further study.

---

B: 1-20 15 lysogenic on BAOR. Suggests that  
BAOR might also be lysogenic on many of these.  
5 may be sensitive. In initial screenings a  
few were noted as possibly mixed typ/sens.

7/18/53. *Bacillus lysogenicis* etc...

	BAOR	PB2	SW666	FA90.
1063 A1	++	-	-	-
B1	++	-	-	-
SW666	-	-	*-	- (shaded)
SW928	-	-	+	-
BAOR	-	-	-	+
PB2	-	-	-	+

note difference of 1063A1/PB2 and FA90/PB2. Suggests 63A1 may be lysogenic for a phage other than BAOR!!

- v. small plaques in A1, B1.

BAOR lysogenic for  $\phi$  carried by 1063A1 = 63C1.

streak out BAOR/90, PB2/90; and 63A + BAOR.

BAOR may be lysogenic for another phage which attacks PB2, C66..., but this is probably distinct from the lysogenizing phage in 63A-B.

Q on bacteriophage SW1060 (Cherry, 305-50):

cross str. on EM13 Lac

	SW1060	PB1
Taunton (Ch. str.)	-	-
Bucles (" ")	4 plaques	-
Dundee / Dundee	11 plaques	ca 30
Dundee / SW818	-	++

After 48 hours, SW1060 reexamined in 1, 2, 3 series  $\rightarrow$  SW1060'

Try to grow Dundee, Bucles on SW1060.

63C1 FA90 x 63B1 / mot  
63C2 FA10 x 63B1 / mot

Hold 63 strains for later lysogen. test

Tom's ?? Malv.

7/1/53...

TEN 174-95 for further study. From W1325 x W1394  
 streakout 2 colonies from EMS lac. 1 → pure lact<sup>+</sup> mal<sup>+</sup>; poor growth on Stac

2 → lac, Mal<sup>+</sup>, - and? ↓

on Stac: many satellite, pinpoint colonies

Restreak 4.?

1, 2, 4: variable colony size on EMBStac; no -

3: Mostly lac<sup>-</sup>, Mal<sup>-</sup>. Occ. small +<sup>s</sup>

7/4.

A. Restreak + colonies from 1, 2, 4 on EMB, EMS Mal / ser.

→ no overt sign of sensitivity in EMB. Edges of many colonies look lysed (or lac<sup>v</sup>?)  
 of EMS in 9.0m EMS!

→ all intact!

Test parents, intact +, - for sensitivity to "e".

B. Restreak 6 colonies from 3 (pres. + or v, not -).

all mottled + on EMB lac → pure lact!

Replis of 1-2-3-4 above: 1, 2, 4 and lact<sup>+</sup> 3 are S<sup>R</sup> Mal<sup>+</sup>

lac<sup>-</sup> 3 = S<sup>S</sup> Mal<sup>-</sup>

C. Tests for  $\phi$ : using parents, 2+ 3- and 4+ as indicators, and parents, mottled + as sources: no signs of  $\phi$  in cross-bush on EMB 0

maybe diploid or more likely, contaminated protoplast is performed substitution.

Try Mal<sup>+</sup> prototypes on EMB Mal / ser.

Mite p.v?

205

7/8/53.

N97 o.c. to both

A1. — in b plates (jars b, and mummata b)  
B  $\Delta$  UV 30 sec (washed cells)  
C  $\Delta$  - 48 ~~sec~~ 10 min.

Numerous swarms - more in B than A? - n b (jars Eudiale)

None on b (mummata)

note A4: b, 1, 2 ++? Rubble per. cool.

7/11/53. (UV - 40 sec.) standardize mummata.

# swarms/plate 24 hours (two clusters)  
A 8, 3, (15)  
B 2, 10, 2  
C 1

D:  
after 48 hours, A and B  
1 each  
2 on mummata.  
b + 1, 2 ++? vs.c.

Too variable for any clear result!

Inoc 7/14 in 1, 2.

~~7/18~~ 7/19. a few beginning to swarm, very weakly

A1, A4  
B1 B2  
C1 C2  
D1 D2.

7/20. C1 }  
D1 }  $\rightarrow$  magg.  
D2 }

Inoc 7/19 in 1, 2

No percept swarms through 1, 2 serum in all tests!

7/26 Profig.

serine essay for Singer.

1068

6/18/53. W-1975, 1977 in D/O) + glycine or serine  
24 hours P17.

r/ml in 10 ml tubes

	0	1975	1977	
L-serine	1	+	+	
	5	++	++	
	10	++	++	
	100	+++	++++	+++ est 10 <sup>7</sup>
glycine	1	+	-	
	10	++	-	
	100	+++	-	(+++ est. ca 5 x 10 <sup>3</sup> )

confirms previous conclusion that W1977 is specific for serine.  
In both cases, 100x serine showed early inhibition - serine itself for  
multiplicity?

6/20/53. W1977

x in sol 2mg/ml  
10x/ml in 10 ml tubes  
= .05 ml

x. ~~100x~~ . . .  
140x . . .  
40x . . . +  
L-serine 1x . . .  
10x . . .  
20x . . .  
100x . . .  
DL-serine 200x . . .  
- .005

Later: Singer's compound has about 10% activity of L-serine.

200x dl-serine = 100x l-serine.

Inhibition of early growth at serine > 50x/ml.

Might be possible to select resistant mutant, not found.

Occasional (ca 1/10) tubes at low serine adapt. (W1977)

Small samples

100x (X)  
50x (X)  
10x (X)

serine 50x  
L 10x

(over)

Full wild growth not achieved for either W1975 or W1977 with 100x serine, or 100x glycine. Mixtures not tried in this expt. Recrystallized L-serine used for these expts.

Suzgi later stated his

compound was mostly

aspartic acid! (Activity parameters?)

~~asp decarboxylase?~~

TM2 morphologic ph 2

7/17/53. SW1061 received from Edwards 7/16.

Both it and slant on tube TM2<sup>2</sup> did not migrate in 1, 2, 3.  
 Serum. = 67A1

cf also Tube-slab "TM2" ph 1, 2 5/12/53., and TM2 phi  
 = 67A2 = 67A3

67A1 }  
 67A2 } movable in 1, 2 serum!  
 SW1061 } all 1, 2 ++ i -

Save SW1061 for future study.  
 Prepare FA 22/1061.

67A probably came from  
 1039-1; confused history, but  
 likely a single USA colony  
 i → 1, 2, while looking for  
 absence of cross reaction.

67A3 = TM2<sup>i</sup> : i ++ 1, 2 -

67A3: 1-4 (s.c.) all react 1, 2 +++ i ++ promptly!  
 i serum Repro 67A3-1 in i

Part 1 → 3 s.c. from NA to with { all 7 1, 2 +++  
 from NA with i - or ± 1 ++ -  
 1, 2 +++ b -

1061 sterile → 1, 2 +++

67B7 = 22/SW1061 → SW666. No ~~react~~ or non b in tubes  
 though numerous swarms in plates and not in  
 tubes to be reported! ∴ H<sub>1</sub> H<sub>2</sub> 1, 2

In 1, 2 serum, A1, A2 and SW1061 are each microblyd.  
 TM2 (A3..) passed readily through both i and 1, 2 → i ++ 1, 2 +++  
 22 x 1061 also microb. → i ++ 1, 2 -

S. benzenoides trans ductan

1068

7/18/53.

SW 1060 from Cherry reported to be susceptible to PB phage  
~~Beules~~ Beules, Taunton. As my hands, Dundee ++.

① in 1,2 serum → c phase. This was tested as follows:

- FA10 (2273) - 1 pl?
- FA90 -
- FA18 (2274) + spreading of plaques
- Dundee +
- Dundee/618 -
- PLT7 +

680 1 Dundee/1060c  
 2 Taunton 1 "  
 3 Beules 1 "  
 ↓ T.O.  
 no activity on 1060c.

This stock seems to be ~~not~~ susceptible to PLT22 and PLT7  
 as well! Attempt to grow these phages on it.

Slide agglutination SW 1060 (from EMBS agar, dust):

-	-
IV V XII	+
IX XII	-
VI VII	+++
VI VIII	++
II XII <sub>2</sub>	-
II XII <sub>3</sub>	-
IV XVII XII	-
C	-
1,5	-
P	+ gran
Vi	-
V	-

definite reactions with *S. paratyphi* B.  
 penum  
 cf. other benzenoides and cholerae suis.

No overt lysis of PLT7/1060c; PLT22/1060c. "lysed" still active  
 on T012 (carryover?). Try -x SW666.

7/20  
 68A1 7/... -x 666  
 68A2 22/... → 666  
 ↓  
 not! → c (probably carryover of FA) type proceeding  
 SW1060c itself partly motile: motility



SW961 also reacted slightly with B serum, C, ++  
SW853, 732, 737 reacted poorly with both in first test.  
853 also showed plaques (?) vs. P7, P22.

T.O. and start fresh after summer.

68B.

in C<sub>1</sub> (smeta) or C (H) serum as indicated  
7/21 - 7/26. Initial results not checked. Initial checks  
in C<sub>1</sub> (.01 ml or .02 ml / tube), about same.

7/26.

1. SW1060C 1C → 1,5
2. " X PA22 1C → 1,5
3. SW1060C 1C<sub>1</sub>
4. " X PA22
5. SW1060 1C<sub>1</sub>
6. " 1C<sub>1</sub> 2x
7. " X
8. " X

Vi + Typhi x-x.

1069

Misc.  $\varphi$  -x 0901

9-10/53. See 1071.

ca 9/10. Motility SW 759, 760 through plates of semisolid.

after 2-3 passages, both cultures were ~~actively motile~~ but reacted Vi + d-, except for an original SW 760 which was Vi-. Passage strains of SW 759 M2 tested ~~x~~ in semisolid tubes  $\pm$  d antiserum.

- A. 1 no serum +++ in 24-48 hours.
- 2 d " 2 tubes Complete inhibition.
- 3 FA10-x, " NO outgrowth
- 4 PA22-x, " " "

resaid after two weeks.

B. Use A1 as medium. - found not very motile. Repeat passages.

9/15. C. Various det't phage from Anderson, rather low titer. Test x SW 957 in tubes: (820, 63, 1, 4, 6, 25', 28', 26', 30', 12) all negative. Refer to fresh tubes for results. 29' swarmed. Repeat test.

10/19. However, "control" PA22, FA85 gave no swarms: former a few tubes? In checks, 28' and 820 swarmed in second tube but not the 1st. Appearance of multiple sites of infection??

Try other  
mediums + Vi.

Hfr - Gal.

Hfr - wj 28.

7/27/53.

W1895 x W1321 in Penassay, then streaked on EMBS Lac, EMBS Gal <sup>sm.</sup> 5 plates each.  
controls OK.

1 full lact 1 full Gal+ (ca 1% est. of all lact+<sup>sm</sup>).

Restreak on EMBS Gal <sup>sm.</sup> Hides 1070 A1, A2 - to HLLS to test.  
(should be M-) <sup>sm.</sup> Hfr ... 2

H303	x1177	control
1070A1	800-1,000 cells/ml	-
1070A2	"	-
control	-	-

10/6/53. W1895 + W2333-8 in broth overnight. 8 streak on EMBS lac <sup>sm.</sup>: All showed lact+<sup>sm</sup> ca 1-3%.

- W 2333
- ~~2334~~
- 2335
- 2336
- 2338

Use 2333 for further expts.

2337 gave fur or name Mutant streak does +.

September 5, 1953.

A. SW957x

B. SW666.

- 1. Control
- 2. FA22
- 3. R (Anderson stroke)
- 4. ♀'s.
- 5. FA84.
- 6.

Plates

Tubes.

A 1 —  
 2 no sw on plate  
 3 ✓ numerous sw. all d.

no sw.  
 no sw !!  
 1 +, 1 -

B 1 —  
 2 ✓ numerous sw —  
 3 ✓ <sup>3x possible.</sup> ~~numerous~~ sw on rec. tracks.

3: <sup>all 3:</sup> <sub>2-d (not depropane)</sub> <sub>1 b (not 51 R)</sub> <sub>bal-</sub>

✓ no sw  
 ✓ sw  
 1 +, 1 -

4 k

✓ sterile

422

✓ sterile

5. 5-6 swarms ca 100 tracks. P. cl. and islets swarms = 71 A5.

∴ R functions as transducing phase. FA22 appears to have lost titre in part. Use 71A5 for x-gallinaria in d serum. (cf. H901)

1071A6. 957/k (not motility). Tested and ✓ resistant to FA51, k.

9/9/53. R-X ... sw.

n motility agar

1 plate each

- C.
- 1 550 —
  - 2 1063 —
  - 3 1064 —
  - 4 1066 —
  - 5 1067 —
  - 6 1068 —
  - 7 71A6 numerous T+S.
  - 8 1072 +

- 1. FA51 (01) -x 71A6
  - 2. 53A " "
  - 3. 53 " "
  - 4. \_\_\_\_\_ " "
- 1 sw ?  
 2 sw

swarms: significance ?

1, 2 +++ i++ (2)++ (5)++ !! (1071)

Note reactivity is ⑤ as well as ②.

Note ⑤ = S. Berlin/S. para B. X-R may be related to ③.

September 15, 1953.

58-161 from store (MLM) streakout. Pick one colony = 1072-01 (EM/3lac)

9/14/53. Restreak of for reisolated, single ~~side~~ clones.

9/15/53. Save samples; inoculate motility tubes and plates.

	A:	P19.	N20	AB	B:
9/18.	1	✓		+++	
	2	+++			
	3	+++			
	4	✓			
	5	+	?	++	
	6	+++			
	7	+++			
	8	+			
	9	+++			
	10	-			
	11	-			
	12	-			
	13	+++±			
	14	++			
	15	++			
	16	+++			
	17	++			
	18	±			
	19	+++			
	20.	+++			

agar probably n.f.

P23+++  
+++.

++ to +++ isolate.

+++ = nearly cover plate or through tube.

→ streak out +++.

Remove + to fresh plates. all to SP18.

hold tubes all day.

P23: A10 only unmotile culture. Reincubate from top.

store streak plates to 10/5. Pick single colonies at A, B.

(inoculum and after growth of second motility selection) for compatibility tests. Note sectoral microids at ~~16~~ 16, 19 A-B, 20 B (same A) with ~~11, 17, 18~~ 11, 17, 18 B. Relationship to selection? Restreak variegated colonies from 19 B. Note noticed in series A.

10/8. Test large loops of concentrated mixtures

2 W1177, W1896 in EM 8lac.  
for compatibility now

10/10. Most X ~~W1177~~ overgrown! check this point!

W1177 (contaminated ??)

	x W1896 yield	predominant hex or -
1	F	
2	F	
3	F	-
4	+	+
5	++	ca = sic
6	++	+
7	F	-
8	F	-
9	F	- fur +
10	F	- "
11	++ ?	-
12	++	- sic.
13	F	-
14	++	+
15	F	-
16	F	-
17	++	+
18	++	+
19	F	- fur +
20	++	+
W1607	++	all -
78A1	F	

XX enrichment  
all X 1177 are XX.

Repeat 11/1/53. (test - B)

	W1177	W1876
A1	+ heavy backing.	+
B1B	2?	+
2	4 5?	+
3	-	+
4	-	++
5	-	+++
6	-	++
7	+	+
8	-	++
9	-	+++
10	-	+++
W1607	-	++
-	-	-

note ratios might be distorted by growth

of 1-10, 1, 2, 7 maybe F+ still. Recheck 3.

Some platings may have been too heavy. unless off!

5, 12 should be subsided after compatibility is contained.

# Motility selection.

1072  
summary.

11/7

8-161 stocks streaked out. Individual colonies = 72A (1-20) grown (at least) in motility tubes (1-10) or plates (11-20). Mass inoculum for second passage. Also streak out these inocula and save s.c.i. = 72B(1-20)A.

Streak out second passages and s.c. = 72B<sup>(10)</sup>B<sup>(10)</sup> for F test.

Tests on 1-10 by JZ (ullogram separately, washed and plated D(0))

11-20 by TCN (grown together, plated on EM5bac Th SM; DO Th sm').

Results on BxB series: All F- except 1, 2, 6, 7, 14, 15.

14 may be feroxi Hfr (cf. W2206 = 1022 C3). Showed up by reduced fertility & F+ as well as ~~incompatibility~~. Few prototrophs were rechecked.

on EM5bac: 1: 1<sup>+</sup>/2      2: 3<sup>+</sup>/5      6: 1+1      7: 3<sup>+</sup>/3<sup>-</sup> and

highly likely bonafide. Should be rechecked, and further selected.

Swiss A should also be renewed.





10/10/53.

A. Test SW1028 = <sup>orig.</sup> N97b → S. marini, b: 1,5 in b, 5 serum for occurrence of 1,2 phase. 1: sterile culture 2: virility passage

B. <sup>1043 FA866</sup> SW1027 (~~FA866~~) → S. marini (Loria kind) / a, enx [for b(12): enx]  
1 } sterile culture 3/4 migrated → b (w/ growth), #4 late - still enx  
2 }  
3 }  
4 }  
no s.c.i. / b. benx  
vb ✓ → enx  
see orig.

C. <sup>FA15</sup> SW803 enx (abony<sup>2</sup>) → ~~S. marini~~ N97 original / b; 1,2 for (b:12) enx

6/6 tubes migrated in 24h. 10/11 PM streak out.

all enx: no s.c.i. in enx, benx

(poss. contamination abony in phase? Re-FA15B <sup>in</sup> sterility. → all b.  
15c ✓

(cf. SW426  
1051-  
1055

P11. Purify B1-3 C1-6.

A17. A1 → 1, ... → s.c.i. 1, 2, 5, 2- 1, 5, ... Probably had some 1, 10  
A2 none yet

C2, 3. slow buds. Others still almost stationary.  
in b, enx

B, C / b, env.

10/16 - C3 b+? z33+? 1,2- env- somewhat rough.

C4 still env+, b- ~~z33~~ 1,2-

These were enlarging buds,  
not swarms.

rechecked on s.c.i.

10/17

swarmed to  
bottom

B3

C1

Retest single colony isolates:

10/19 C3 : b. z33? + 1,2- env-

C7 : env+ b; 1,2; env-

C4 : env+.

B3 : Rough. b?

benötigt.

1074

A 976  $\rightarrow$  main  $\rightarrow$  b; 1,5 / b, 5  $\rightarrow$  0 ✓

B) SW1043  $\xrightarrow{S:}$  lambda bridge  $\rightarrow$  b  $\rightarrow$  cur ✓  
a: cur

C) along cur  $\rightarrow$  N976  $\rightarrow$  cur  $\rightarrow$  b  $\rightarrow$  cur ✓  
b: cur  $\rightarrow$  0.  
~~not no, 2~~  
(higher)

---

proof cur of cur  $\rightarrow$   $H_1, H_2$  <sup>1,2 cur</sup> is still  $H_2$ ?

---

ca 10/10/53

Note: 1043: pullorum was tried extensively on SW1040/a and failed.  
Other pullorum FA should be tested → 957 for highest activity.

- A.  $\swarrow$  → 957 Test n.g. 3/4 owing to overgrown contamination. All others gave numerous tails. Also swarms in P2, 9, 12. P2 probably most active. (Test each on 1/2 small plate)
- X B. P, G1 → <sup>long lines</sup> 874 / 9, env no swarms. test 2-P and 1-G anyhow 10/18: still env
- X C. P, G1 → <sup>astroware</sup> 770 / chx; 1,5 2 each. No swarms.
- 13 D P, G1 → SW967 (dublin 0) G1: few long tails, 1 swarm. P1 <sup>singly</sup> ~~no~~ tails!
- X E P, G1 → SW989 (TM 0) no T or S in either
- X F P, G1 → SW991 (dublin i) / i 2 each
- X G. P2 → SW1004 (summit) / 9, 1,5 ca 10/14. } 3 each 10/19.  
X H. G1 → " " " } no swarms
- J P2 → SW1040 / a No swarms 10/19. Hold to..

C, F poor vegetations, rough. C still 1,5... F still in.

K ① SW1028 / b+5. → slow but: b- 1,2+++ 1,5++  
② (noted) v. slow but not 10/19. 2- 5:.

1,10??

(on next transfer → 5++ again)

S. typhi seemingly only receptor for phage from gallinarum. Results of pullorum equivocally negative.

X. not finished.