(49) Total Review of pedigree expts. APR 18 1956

Exp.		1954	SW666
1137	✓	2-15	12
2	✓	16	12
3	✓	17	12
4	✓	18	12
8	✓	3-1	967
41	✓	9	12
42	✓	11	960
43		16	60
44		17	60
47		30	60
49		4 2	60
50		12	60
51		14	22 × 578
36		2 23	12
37		26	× 967
40		3	92
46		3 19	9 × 967

Tabular resume of pedigrees.

APR 19 1956

a = minimum interval of branching.

Exp. ☐ Initial

Interval of latest branching | interval of termination

a b c d

how many intermediate branches.

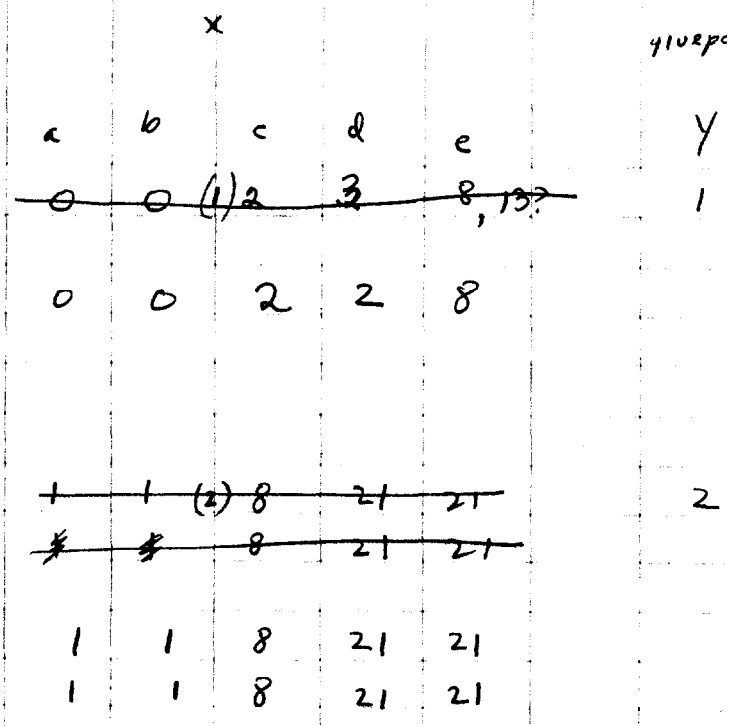
max. yield per branch

APR 19 195

#			a	b	c	d	e followed to	Yield
1131	CD	followed 3 generations. All cells became immotile. Separated at n_3, 1 cell gave a swarm; D3. the others non motile clones. On further isolation, D3 was mixed → 12⁻ : 8⁺, the latter all i, the former x — FA10 → b.	0	—	—	—	1sw/n_4.	1s
	C5	all nm at n_{13}	0	13 (0)	13	13	13	1
	B5	" "	0	13 (0)	13	13	13	1

APR 19 1956

1132 E5 Unicat. 2 divisions.
8: all⁻ at n_3, and at n_8



D5. dicat, branch at n_1.

APR 19 1956

1133 E5 pure swarm / late isol.; 1/4 inv. 3/4 → H_1^i

	a	b	c	d	e	Y.
D5	1	11	11	11	11	
C5	0	10	10	10	10	1
B5	0	10	10	10	10	

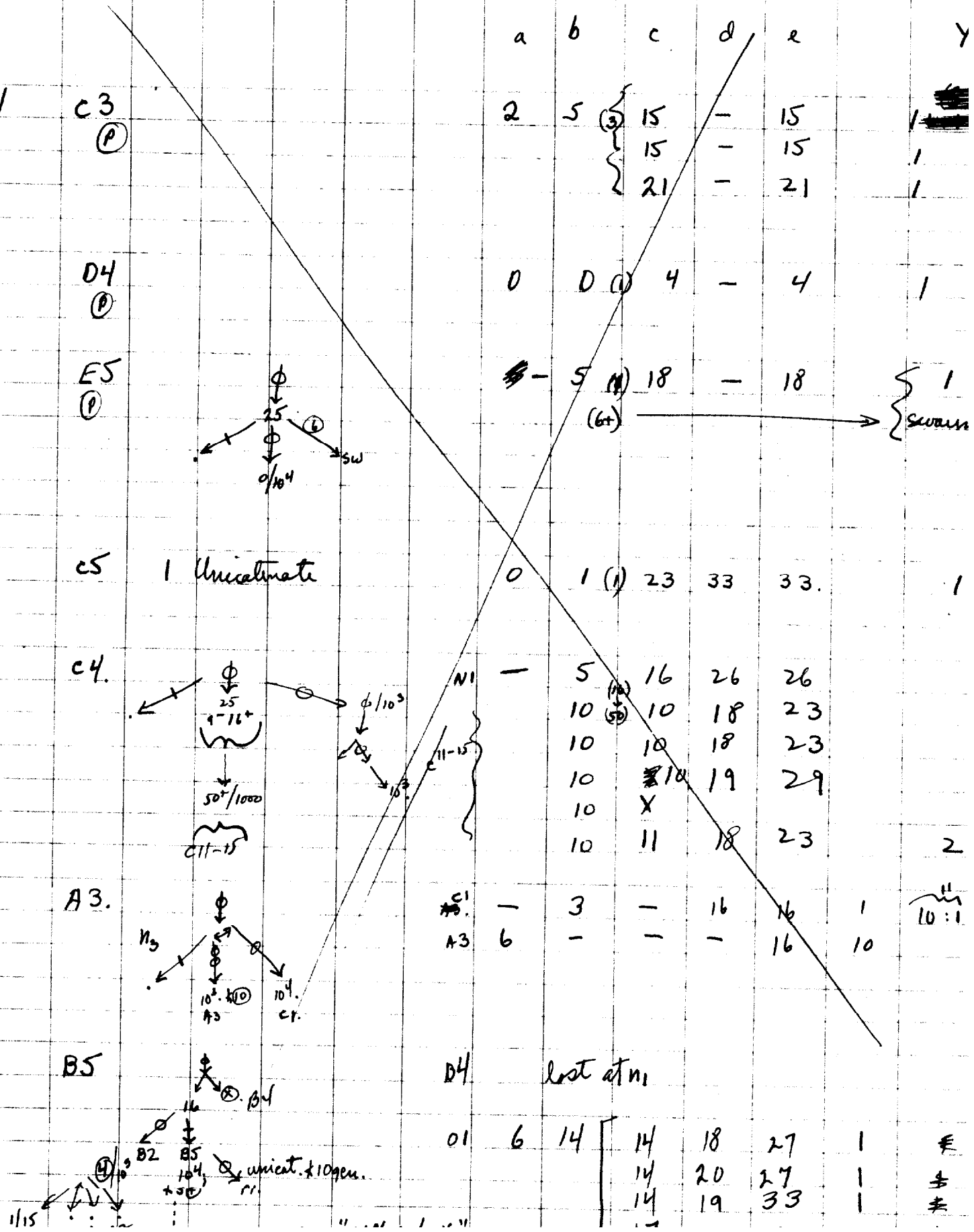
test for swarm essentially

(P) in notes

APR 19 1956

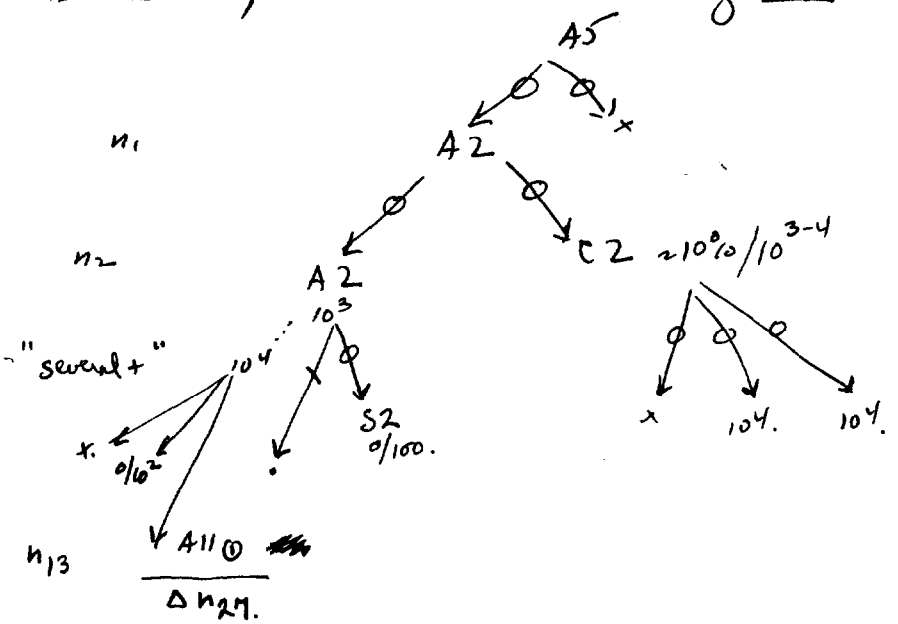
how many + at this time

		a	b	c	d	e	y
1134	c3 (P)	2	5	(3) 15	–	15	1
				15	–	15	1
				21	–	21	1
	D4 (P)	0	0	(1) 4	–	4	1
	E5 (P)	–	5	(11) 18	–	18	1
				(6+) →			scram
	c5 1 Unicellmate	0	1	(1) 23	33	33.	1
	c4.	N1	–	5	16	26	26
				10 (5)	10	18	23
				10	10	18	23
				10	10	19	29
				10	X		
				10	11	18	23	2
	A3.	c1.	–	3	–	16	16	1	11 10:1
		A3	6	–	–	–	16	10	
	B5	D4		lost at n1				
		01	6	14	14	18	27	1	✱
					14	20	27	1	✱
					14	19	33	1	✱



Note: 55 was mycoplasmadiagnosed as a
sarcoma as a positive ex. of a pluricatenate clone.
Exact number not specified. 5 isolated.
Pl was taken as an early sample

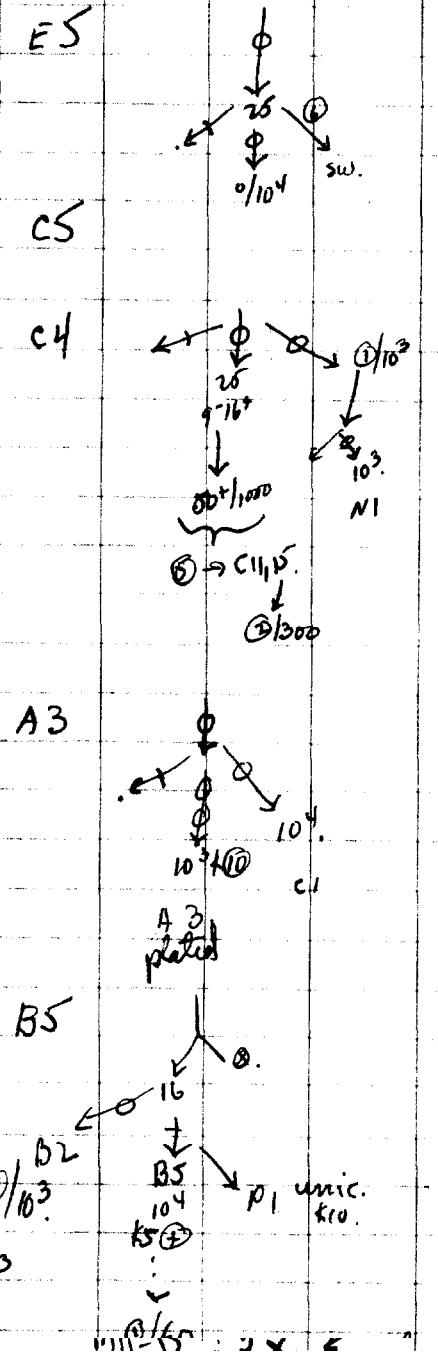
- Hare state Pl. cell isolated at n_{22} was a chain for at least Δn_4 and seen again at $n_{39,442}$ as no motiles. The chain had sibbranches between n_5 and n_2.
a should read, ~~earliest~~ latest time that branching must have ceased for that chain.



No
evidence.

4/21/5

		a	b	c	d	e	Y
1134	C3 (P)	D1	3	5	15	—	15	3
		D2	3	5	15	—	15	
		D5	3	5	21	—	21	
	D4 (P)		0	0	~~5~~ 4	—	4	1
	E5		1	4	5	17	17	1+x
			2	→			swarm	
	C5		0	0	23	33	33	1.
	C4		4	5	16	26	26	
			6	10	10	~~18~~	24	
			6	10	10	18	24	
			6	10	19	27	27.	
			6	10	~~19~~ x			
		C16	~~11~~	~~18~~	~~18~~ 18	26	26	
	A3		3	3	3	17	17	1
			6	13	—	—	13	10
	B5	B4	1	x				
		P1	5	13	23	33	33	
		01	6	14	18	—	25	
		91	6	14	14	18	27	
		2	6	14	14	20	27	
		3	6	14	14	19	~~33~~	
			5	13	—	21	21	



4/21/56

1134 A5

see for partitions @ '10:10?

terms observed.

	a	b	c	d	e	x	y
A5	1	—	—	—	1		
S2	6	12	12	19	22		
A11	6	15	42	~~42~~	~~47~~		
12	6	15	x.				
13	6	15	—	22	22		
* D11	6	14	x				
12	6	14	—	27	27		
13	6	14	—	27	27		

~~+ 5 12 —~~

A4

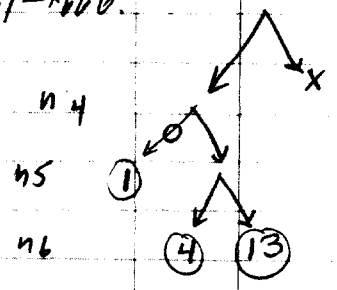
see @ for 4 pt sets and terminal patterns

	1	5	12	—	12
	1	5	20	—	20

4/21/56

1138
B4

967–666.



	a	b	c	d	e
F21	5	5	44	45	51
			19	28	28
D22	8	15	[illegible]	[illegible]	[illegible]
23	8	15	[illegible]19	[illegible]24	[illegible]31
24	[illegible]8	15	32	44	44
E24	8	15	28	28	38
A21	10	16	18	—	18
22	10	16	19	32	32
23	10	16	17	29	29
24	10	16	32	-	32
25	10	16	31	35	42
B22	10	16	32	—	32
23	10	16	17	29	29
24	10	16	19	26	26
25	10	16	33	38	38
C21	10	16	20	20	30
22	10	16	19	32	32
E21	10	16	29	[illegible]34	39
22	10	16	44	50	54

4/21/

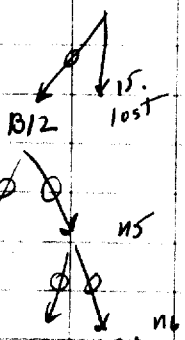
V.S.

1138

	a	b	c	d	
F21	5	5	44	—	44.

967 + 666. most lost.

B4:



-chain
$\Delta n = 39$
=

D22 A 21-25
23 B 22-25
24 C 21-22
E24 E 21-22.

E21 n_{13} → B32 term.
E22 n_{13} → B33 n_2; n_8 → E31 → n_1, n_1, n_1 E32 $n_2 < n_6$ can't to n_{10}.
E23 n_{13} → B34 $< n_5$.
E24 n_{13} → C31 term.; to minimize.

A21. uncat n_2, ⇒ C23
A22 " n_3 C24
A23 " n_1 ; term $< \Delta n_{13}$
A24 " n_3 ⇒ C25.
A25 n_2 → D25
B22 n_3 ; n_{13} → A31 ×
B23 n_1 ; term $< \Delta n_{13}$
B24 n_3 $< \Delta n_{10}$
B25 n_2 n_2 E23
C21 t.f. n_4:
C22 n_3 ; $< n_{13}$
C23 ① — lost
C24 ① $< n_{13}$
C25 ① n_{13} → A32 ×
~~D21 : E24~~
D22 n_4 ; $< n_{13}$
D23 n_1 → A31 term $< n_6$

recover for tabulation.

1141 — 13 init. 1E (33)

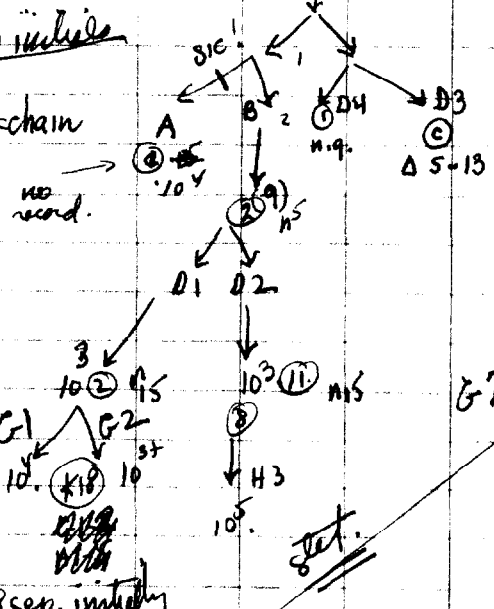
3 died
1 lost
8 E.

4-22-5

1141.

13 include

ⓒ = chain

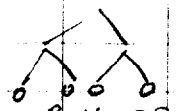


		a	b	c	d	e
A4	A	2	2	15	—	15
	D4	2	—	—	—	2
	D3	2	2	7	13	13
	G1	16	16	16	29	29
G2	1	19	27	28	32	32
	2	19	27	28	37	37
	3	19	27	28	40	40
	4	19	27	28	32	32
	5	19	27	27	32	32
H3	1	19	27	28	32	32
	2	19	27	40	47	50
	3	19	27	29	29	× 3 nuclei
	4	19	27	28	41	41
	5	19	27	28	41	41
A3		8 cells.			29	29

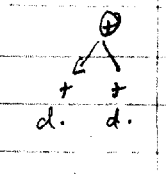
persisted G

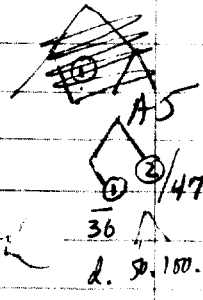
pulsations

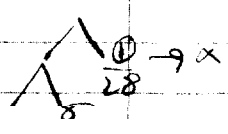
C2



a		2	2	46	6	11
b		2	2	7	7	20
c		2	2	5	5	18
d		2	2	2	2	15
A1	② d. after 1 division	1	—	—	—	1
B5	① n.q. — persisted 48h. comatile.	0	—	—	—	0.
A3	① died overnight	0	—	—	—	0
A5		1	5	—	—	5
		2	5	—	11	12
		2	5	—	12	18
B1		0	0	1	2	18
B2.		no data				
	①	0	0	1	1	15







APR 24 1956

1141A4.

		a	b	c	d	e	
	A4A	2	2	15	—	15	
	D3	2	2	7	13	13	
	D4	2	—	—	—	2	persisted
H3 -	8 cells D2	8	15	—	29	29	
	G1	6	16	—	29	29	

a	b	c	d	e
19	27	28	32	32
19	27	28	37	37
19	27	28	40	40
19	27	28	32	32
19	27	27	32	32
19	27	28	32	32
19	27	40	47	50
19	27	29	27	x sm.
19	27	28	41	41
19	27	28	41	41

G2 K18 (+) H2

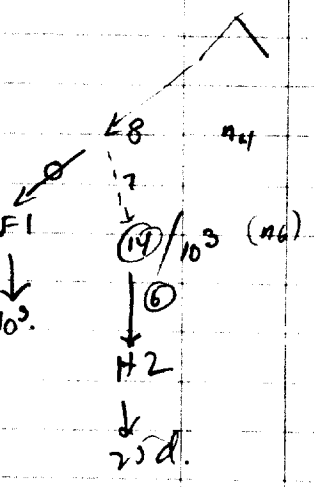
---

			a	b	c	d	e
Or alt. interpretation,	1141A4 for G2 subset →	H2.	6	16	31	—	31
	1141B4 :	G2 series	13	21	etc.		

Subtract [6] from above.

APR 24 1956

11/4/B4.



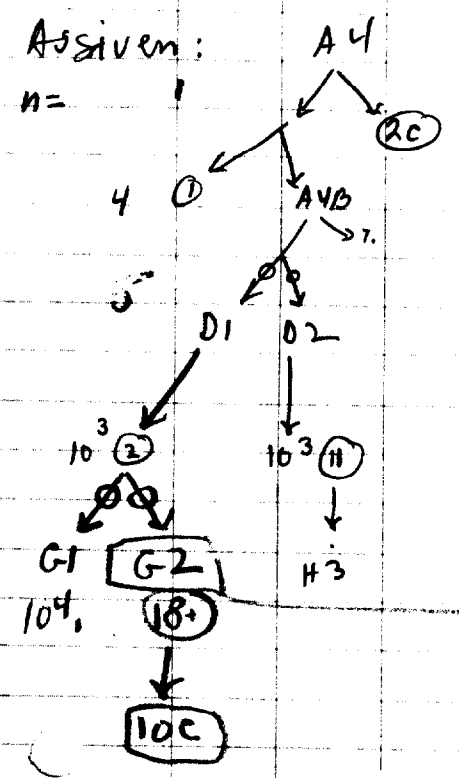
	a	b	c	d	e
F1	4	—	—	14	14

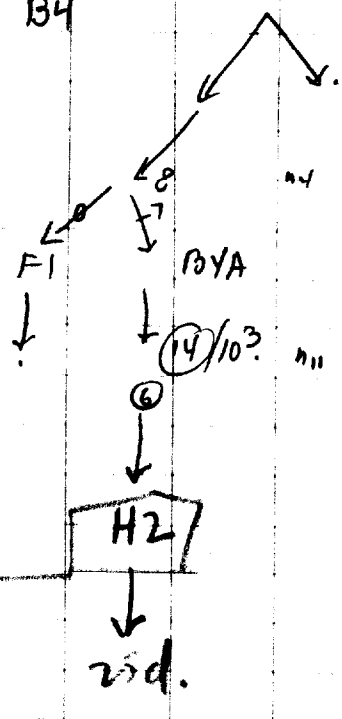
Note: notes are confusing on the C2 - H sequence of A4:

Note 14: partition.

H2	6	—	—	—	6 (lost by dying).

A4 and B4 pedigrees may have been mixed at C2 ↔ H2 so following interpretations are ~~possible~~ indicated. This seems





An account of H1, the given interpn. seems most likely, but it should be recorded with reserve.

APR 28 1956

1142
60 → 666
1/2 for plating

No early separations: survey of yields of ①.
A1-5 B1-5 ⑩ were ess. inoculable.
C-D followed ERG-plated.

Yields at first examination	+	Tot	$\log_{10}$
c2	2	2⁻	
c3	>50	3	
c4	10	3	
c5	1	4	
D1	>100	4	
D2	3	3	
D3	0	3	
D4	0	3	
D5	1	3	

		a	b	c	d	e	
C3	c3 c4 c1	6	10	10	23	23	…
	c2	6	10	23	29	33	
	c5	6	10	31	37	41	
D1	(3)	7	13	—	26	26	
	(1)	7	13	14	—	14	
		7	13	26	—	26	
		7	13	26	28	36	
		7	13	26	31	36	
		7	13	26	29	36	
		7	13	26	31	36	
		7	13	26	31	36	

C3 → ⑩/10^3

⑤ → 105: C1-2-3-4-5

10⁴. ①/10^4 10⁴. 10⁴. ①/10^4

106
C1 → 50.

C2 → ①/100 → ①/2 → 50., 10^3.

D1 → >100/10^4 ⑩

105 D,E (1-5): 3: 10⁴ / 1: 0 (5²⁴) / L. 10^4/①

D2 C5 0.
D5 B5 4.
F1 → B1 28.
C2 → B2 8.
F3 B3 50.
C3 nil →

108

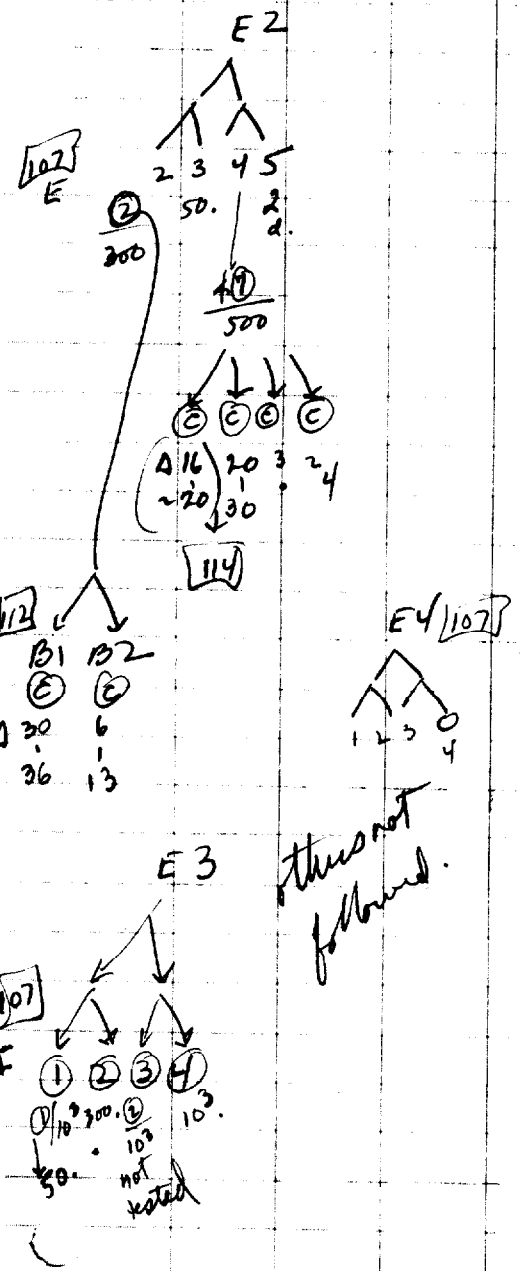
Platings: 16 "calls" plated → 9 grew (1 was 3 mixed, 5 reduced)

↓ residues

1143

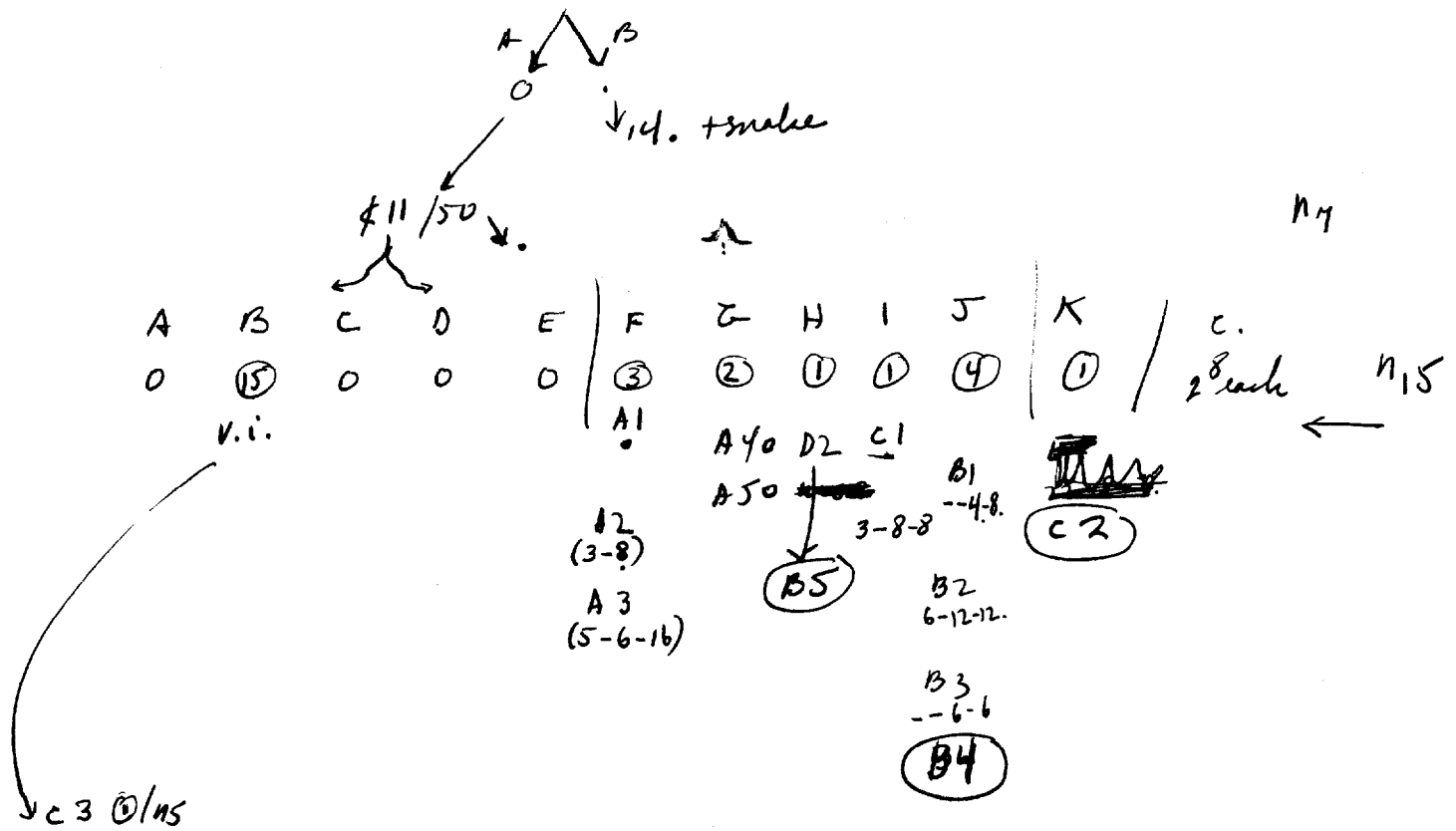
Plating: ~~4~~ 14/16 hd dup colonies; no Tox S.

110 E1-5 → 107



	a	b	c	d	e
	~~10~~	~~40~~	~~46~~		
	~~40~~	~~16~~			
{	3	10	40	46	46
{	3	10	16	23	23
E3	2	—	—	8	8
E4					
E5	2	—	—	3	3
BB	~~3~~ 3	11	27	31	31
4	~~3~~ 3	11	34	41	41
5	~~3~~ 3	11	14	17	~~19~~ 19
C5	~~3~~ 3	11	—	15	15
F	4	2	—	—	12	12
	~~2~~	2	—	—	10	10
	3					
	1	2	2	12	18	18

1144 B4



C3 ①/n5
C4 n6.
C5 ①/n5
D1 71.
D2 n5 n8.
D3 — n6 n10
D4 - n5 n10
D5 — n4 n10
E1 - 22-28-28
E2 · n5 n5
E3 · n4 n7
E4 · n4 n7
E5 · n5 n9
F2 · n5 n9
F3 n5 · n11 n11

$[\bar{x} = ①/2^x]$

Σ 34-38-39 C2 $\bar{n}_6$ $\bar{n}_1$ $\bar{n}_6$ $\bar{n}_1$ H3 (n32) # $\bar{2}$ $\bar{7}$ $\bar{5}$ H1 $\bar{7}$ $\bar{7}$ $\bar{4}$ G2 $\bar{2}$ $\bar{2}$ F2 $\bar{7}$ $\bar{7}$ 4 5.

Σ 21-23-31 B4 $\bar{10}$ $\bar{7}$ H2 (n32) $\bar{5}$ I2 $\bar{7}$ $\bar{4}$ H2 n2. n_{10}.

Σ 37-37-37 B5 $\bar{5}$ $\bar{4}$ $\bar{3}$ H1 (n36) $\bar{9}$ I1 $\bar{3}$ $\bar{3}$ H3 $\bar{7}$ $\bar{10}$ G3 0

But see notebook for missing items: now K → C2

But acc to notes, {C2 is n_{27}} {H3 = n_{32}}

Then does OK; there must have been an n_3 virtual between K → C2, H3 ; H → B5 (n5)

114434 any other summaries?

5/2/56

			a	b	c	d	e	
(11) isolat n_7			7 – 4 cells O.					
	H	B5	5	7	44	44	44	duel at n_{44}
	I	C1	5	7	10	15	15	
	K	C2	5	7	41	45	46	
(27)					both d.			
(2) isolat n_{15}	G	A4 A5	9	15	–	–	15	
(3) isolat n_{15}	F	A1	9	15	–	–	15	✓
		A2	9	15	18	23	23	
		A3	9	15	20	21	31	
(4) " " n_{15}	J	B1	9	15	–	19	23	✓
		B2	9	15	21	27	27	✓
		B3	9	15	–	21	21	✓
		B4	9	15	36	38	46	✓
(15) " " n_{15}	B	C3	11	15	27	32	32	
		C4	11	15	–	21	26	
		C5	11	15	20	25	25	
		D1	11	15	–	21	25	
		D2	11	15	–	20	23	
		D3	11	15	–	21	25	
		D4	11	15	–	19	25	
		D5	11	15	–	19	25	
		E2	11	15	–	20	20	
		E4	11	15	37	43	43	
		E3	11	15	19	19	22	
		E4	11	15	–	19	22	
		E5	11	15	–	19	24	
		F2	11	15	–	20	24	
		F3	11	15	20	26	26	
Isolated at n_8, n_{15}	H	B5	6	7	59	59	59	
	I	C1	5	7	10	15	15	
	K	C2	5	7	52	56	57	

3/30/56.

11/4/4

8 clones started

109

A2

① 52

a.

A5

① 26

10².
10⁴.

B1

⑩ 50

⑯ /100.

not pursued!

B2

⑩/25 ns

↓10

0/10³

*8 10³

*6 10³

*1

not pursued.

A3

⑧/64

Note partition

① 300

② 200

D1 D2

114

G1 G2

150. 60.

10⁴. 10³.

B5

③/5.

lost.

B4

see next page.

	a	b	c	d	e	
A2	0	0	6	—	6	
A5	0	0	5	12	17	
B1	3	—	—	—	—	
B2	5	15	—	—	10	① → 0
	6	15	-	-	15	① → ①
	8	*	not isolated			① → *6
	8	—				① → *8
B3	Pure swarm		Pure gm		saved.	
A3	1	6	14	21	27	
	1	6	14	20	24	
B5	2	6	—	—	—	3 cells lost.

5/2/56.

1149

This expt. designed for swarms; did not search deliberately for oligocalenates.
• = 0 or 1 or 2 Fla+.

35 Fla+ [118] 3 n.g. 4 swarms; 1 $\frac{\text{KB}}{200}$ G3; 19 few ⊕
8 few or no ⊕.

SWARMS: (2 hours)

B4: pure motile
G1: pure motile variable expression. Question of phenotypic variability.
H2:
H3: ± ± both pure S. low manipet

H2 × B → F1 — 2 cells { a 2, b 14, c –, d 24, e 24
slb { about 1–10% Fla+ swarm.
Conf 1/5 – key progeny test
∴ ± 20% + in H2B3
at least 2 genetics segregation. c/ B4, G1, H31
not segregating,
esp H3.

F1 → ① 2/10⁴ → [119F1] 10³.
F1 → ② 100/10⁴ → 20 → sw. not yet
F2 → 10³.
some +, some – definite

Note: some late isolates had 60%+ but were lost. These probably showed incomplete expression but late segregation not excluded.

F2 → ① → 10² → [119E3, E4]
F2 → ② → .10³

Expt. monogr? (esp. H3)

118 G3 48/200 → ③ 750/10⁴ → 119F4 → 20 → K2/10⁴ → 121F3

typical incomplete

1148

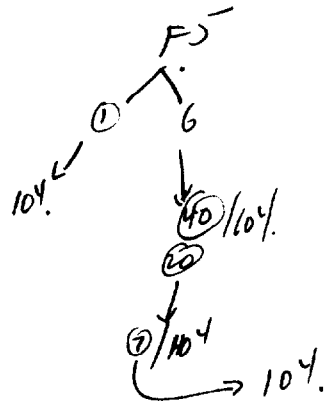
10 cells: 2 ng
2 ⓪
4 ~~2~~ ①
2 plus cat

B1. ⑩/300

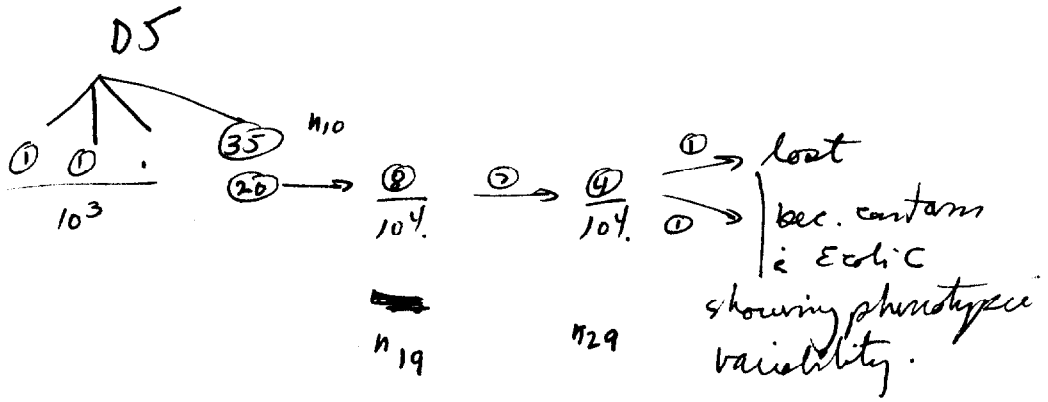
B4. ~~39~~/10^3. no pattern data.

Cat
120

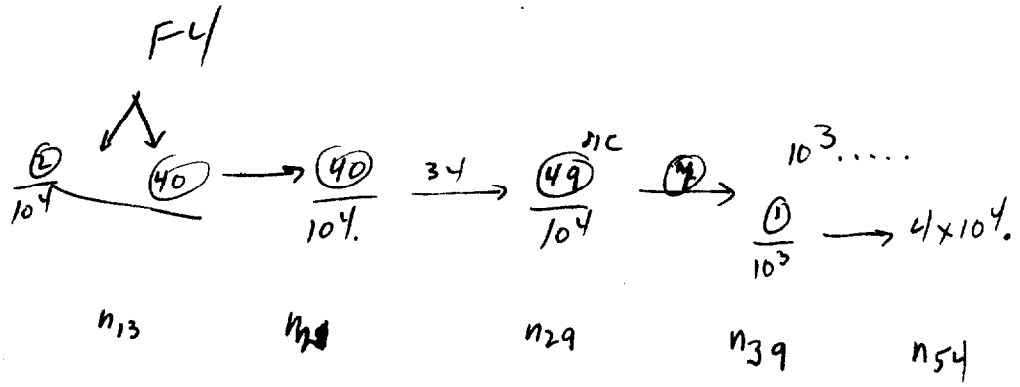
1149



Not certain when single chains



Remarkable problem of variability in Flat clones



1149

122 → 123

37 ① → 2 swarms B5 F5 pure pure

⑪ → swarms 71. (too few?)

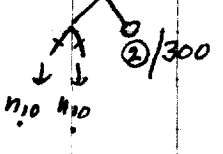
intended to separate early characters to make it easier to handle. primarily for segregating clones.

gave no ⊕ at n_{10}: A2, A3, A4, A5 B1 B2 B3 B4 C1 C3 C4 C5 D2 D4 E1 E2 E3 E5 F1 G1-G2 G3 G4 H2 H3 H4 (mislabeled = "uninteresting") = 26 clones.

These pedagogies generally are not useful except for partitions as no intermediate isolations were made. Note 2 pure swarms. Some late isolates were lost.

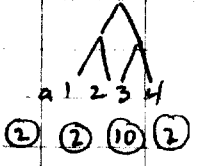
No indication of late replications.

A1



B5 ⊕

C2

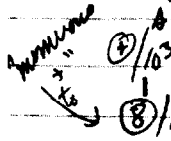


Partitions: at 1st div.

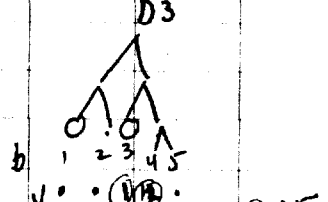
2 : 0
2 : 2 : 2 : 10
0 : 0 : 1 : 12
1 : 1 : 0 : 35
1 : 40
2 : 20
2 : 40

sw: pure
sw: pure.

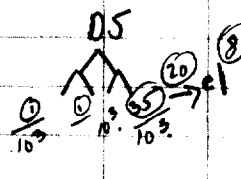
D1



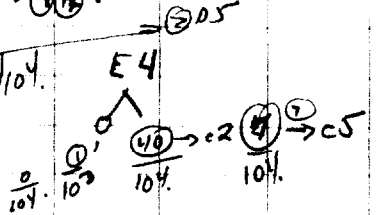
D3



D5

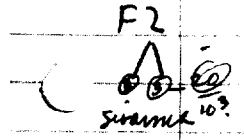


E4

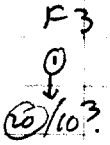


(natural F4 a 6 c d e / 6 39 — 54 54)

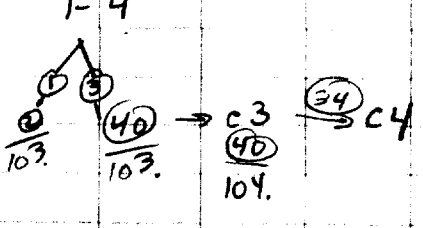
F2



F3

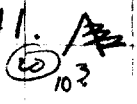


F4



F5 sw. ⊕ pure

H1.



11/9/13 Crossovers among tail ends.

35 from 11/9 x-FA60., x-FA90 → only b

F1 H5 D3 F2 all ++.
but 60C dubonic lysate as it transformed 967!

Found that SW967 is lysogenic on SW661 for a → x phage

1150

60—×666

1 solutes at 2½-3+ hours. 40 insolutes

(134) ↓ 126 A2 < pneumaum. Divide at n_1, n_2 when possible. Patterns or guides.

	at n_{10+}		
A3	1:1		
A4	1:0	obj not	
A5	0 ①		
B1	0 2		
~~B2~~	~~④~~		
B3	① ①		
B4	① ①		
B5	①		
C1	③ ①		
C2	① 0	obj nm	
C3	③ ①		
C4	③ ①		
C5	① ①		
D1	① 0 0 0		
D2	①		
D3	0 0		
D4	0 ①		
D5	① ①		
E1	① ①		
2	① 0		
~~3~~	~~④ ⑤~~		
~~4~~	~~4 ①~~		
5	① ①		
~~F1~~	~~① ④~~		
~~F2 3 0 a~~			
~~3~~	~~① ①~~		
4	① ⑤		
5	⑤		
~~G1 .... ① ++++~~			
2	① 0		
3	① ②		
4	①		
5	①		

H1 ① ①
0 ö obj ①
① ①
① → H3 126
①

Intermediate patterns

E3
④ ⑧

E4
⑦ ⑪ 1266 ⑬

A1
① · ㊸

F3
㉚ · ⑥

E5
① · ⑳

H4
㊿

C1
㉖

~~㉗~~ F1
④ · ㉗

B2
⑫

No further pedigree.

Are plus cotyledons delayed?

5/16/56.

→ × paralytics

TM 2 → × SW578. Most ⊕ proved unstable, and motility sluggish..

~~4 ± 5~~ 16 INV.
10 → 140 ⊕/10^3.

3: 2 clones

B1 ~~3~~ 5 /10^3.

C1 ~~3~~ 7 /10^3. These ⊕ proved very sluggish + could not readily be followed.

E2 ~~8~~ /10^3.

F1 2/10^3.

also remarks on stability of $F1a^+$ in E coli C.

at this point these expts. were interrupted for pair isolation.

5/22/56.

1212
Resumé

Representations and splits, early.

	Splits	at n		n		n	Σ n
A1.	3, 3	1					6
B1	1, 1	1					2
C1	7, 6	1	7, 0	2			13, 7, 6
			4, 2	2	3, 1	3	
B3	6, 3	1	6, 0	2			9, 6, 3
			3, 0	2			
D1	7, 4	1					11
E1	1, 1	1					2
F1	1, 7	1					8
G1	1, 1	1					2
H1	4, 2	1	3, 1	2			6, 4
C3	lost						—
D3	nf						—
E3	2, 0	1					2
F3	swarm	save. (prob prime)					sw. no seq?
G3	1, 1	1					2
H3	4, 1	1	2, 2	2			5, 4
B5	1, x	1					—
C5	24, 2	1	21, 3	2	20, 1	3	26, 24, 21, 3
					2, 1	3	

17 12 deletes.
A - x.

5/16/56.
5/17/56.

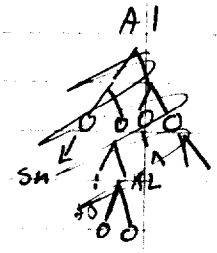
notes

Recapitulation

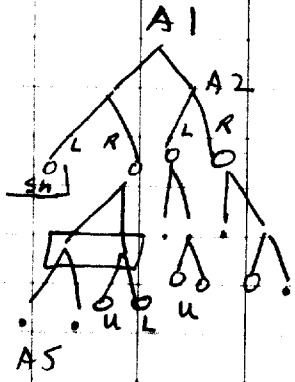
(18 real from 22 → 666 (3 hours)

37 → 666

count subclones at n_{10}.

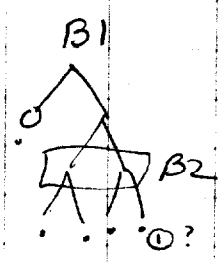


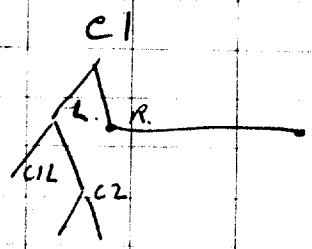
no + at n_{10}.



F3 swamp.

prob pure.





ch.

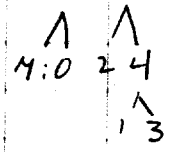
See protocol sheets.

Significant patterns.

A1: (3) (3)

B1: (1) 1:1

C1: 4:6

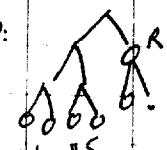


at n_1 unless indicated.

B3: 6:3

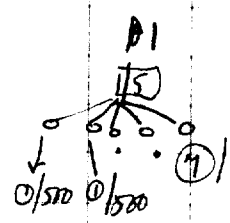
H1: 4:2

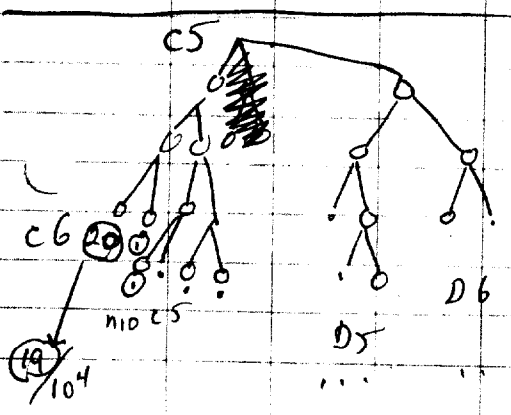
H3:



4:1

C 3 1:1 C





24:2 n_1

(21:3) n_2

(20:1) n_3