

Composition
Graph Ruled

Carle's Notebook

M52-
protein coat ~ 180 coat protein subunits

coat protein subunits - A, B, C

The FC loop interactions play a complex role in
RNA particle assembly and stability

→ mutations in the FC loop → failed to form particles

↳ Some form - but phase not well defined
lines not visible

A contains coat protein dimer (CCPD) can be formed by
protein a coat protein gene about the same
amount as in other regions

↳ protein dimer look at β subunit

Insertion Pr. B - into the CCPD @ 72-75 d.d.
↳ 54 d.d.

Protein - at the β -terminal
Site - sites A + B

Insertion Protein - same into the gene - same protein
- ~~app~~ - copy gene into the protein site

M52 - β insertion in coat protein @ 1380T + 1383C

$$\frac{400 \text{ (5x)} (L)}{5} = \frac{(5x) 1000}{5}$$

unit = trans. DNA dig. 1 hr

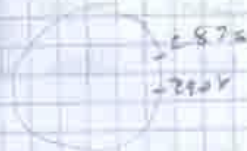
1 ug DNA plasmid

1 - 10 mins longer to lig

$$660 \text{ cells} \times 5,206 \text{ bp}$$

$$3435,960 \text{ genes/mole}$$

$$58 \times 660 = 33,660 \text{ genes/mole}$$



2.8 AT

Ligation

DNA Fragments	100-1000ng	200 (100ng/200)
cloning vector		100
low ligase activity		100 (100 u/200)
ligase (T4 DNA)?		
H ₂ O		
		<hr/> 1000

control - no DNA insert + with known blunt-ended sites

- removed 5' phosphate from all the fragments - control
- phosphatase treatment of plasmid digest - prevent self-ligation



- RT digest
- Buffers
- Purified all the RT and phosphatase. (DNA) Purification

Ligase requires a 5 phosphate + 3' hydroxyl - on digested fragment

Insert = Vector ratio

minimum 3:1 to 10:1

plasmid DNA can be used directly for ligation

Note

but if more than 1 μ g of DNA is ligated then
or if ligation time is from
a long gelting/boiling step afterwards -
use to dilute the ligation mix.

10x Buffer	Making	10x TBE Buffer	5x TBE Buffer
(5x) base	Tris-HCl Base	108 g	54 g
	Sodium EDTA	7.5 g	3.75 g
	Boric Acid	55 g	27.5 g
	dH ₂ O to	1L	to 1L

Making 1% Agarose Gel (500 ml)

1) 500 ml 0.5x TBE Buffer

$$(5x)(L) = (500ml)(.5x)$$

2) 5g agarose powder

$$L = \frac{250}{5} = 50 \text{ ml}$$

3) heat up in microwave

4) 25 ml EtBr

5)

1. to 6.5

Allyl Waters

required to purify oligos longer than 50nt
by gel electrophoresis.
↳ false not being fixed

Annealing oligo protocol

1 μ g of each oligo

mix in K₁₆ buffer (50 mM NaCl, 10 mM Tris-HCl,
10 mM MgCl₂, 1 mM DTT, pH 7.9)

- heat to 95°C - 5 min

- cool to Room Temp for 30 min

Subsequent ligation rxn

2 μ g of the annealed oligonucleotides

50-100 ng of BstI-linearized AAV vector plasmid

Standard rxn conditions & ligation enzymes (T₄ DNA ligase)

SHV AAV vector plasmids are selected & grown in

DH10B bacteria, using ampicillin-containing LB medium

the is a really small piece

the host is small

the is not a piece

→

JS AAV-ESVc GFP-U6 Restriction Digest

with BBS-E

	(-)	Exp 1	Exp 2	(+)	(-)	plasmid = 1.0 µg/µL
JS AAV-ESVc GFP-U6	1 µL	1 µL	10 µL			
10x Buffer 12	2 µL	2 µL	2 µL	2 µL	2 µL	2 µL
BBS-E	—	2 µL	2 µL	2 µL	—	[BbsI] 5 units/µL
H ₂ O	17 µL	1 µL	6 µL	11.5 µL	17 µL	
	20 µL	20 µL	20 µL	20 µL	20 µL	

incubate @ 37°C for ~30 min.

Need 50-100 ng of digested linearized JS AAV plasmid for
ligation Rxn.

Add phosphatase to remove 5' P_i - ask Jennifer or Andy

• check digest on gel

• Purify the Plasmid DNA (same as PCR purification)

♡ GABE

5:37 PM

Ligation (Insert: Plasmid) 20 μL rxn.

	#1	#2	#3	#4	#5	#6
	3:1	5:1	(-)	10:1	20:1	50:1
SHRNA Insert	1.195 μL	2.245 μL	-	4.495 μL	8.99 μL	22.475 μL
	1.95 μL (1:10)	2.95 μL (1:10)		4.95 μL (1:10)		
dsAAV-EGFP	1 μL	1 μL	1 μL	1 μL	1 μL	1 μL
	(100 ng)	(100 ng)	(100 ng)	(100 ng)		
ligase	2 μL	2 μL	2 μL	2 μL	2 μL	2 μL
Buffer						
DP (50/50/0.5)						
T4 DNA ligase	1 μL (100 U)	1 μL	1 μL	1 μL	1 μL	1 μL
water	13.94 μL	13.44 μL	15.89 μL	10.44 μL	14.94 μL	15.44 μL

Incubate for 10 min

Transformation

Note: after ligation you have to transform it!!!

competent cells (100 μL cells)

60 μL

need 3 μL of dDA ligase (no more than 5%)

1 μL

1 μL of 1B

Making 500 mL of standard buffer.

1M NaCl, 0.5M EDTA, 1M Tris pH 8.0

$$(50 \mu\text{M})(500 \text{ mL}) = (1000 \mu\text{M})(x)$$

1000 μM

NaCl $\frac{2/500}{1000} = 25 \text{ mL}$

50 mL

$$\text{EDTA } (1 \text{ M})(500 \text{ mL}) = (500 \text{ M})(x)$$

500

$$\frac{500}{500} = 1 \text{ mL}$$

$$\text{Tris } (10 \text{ M})(500 \text{ mL}) = (1000 \text{ M})(x)$$

10 mL

$$\frac{5000}{1000} = 5 \text{ mL}$$

935 mL

1000 mL

Standardization Protocol 1/8/2011

Alga #1 0.3137 M (100) 3.1 \times Alga A (1:10)

Alga #2 0.3127 M (100) 3.1 \times Alga B (1:10)

20 \times 5x standard buffer

73.8 \times H₂O
100 mL

- low added mineral oil - 60 mL

• Heat @ 55°C in 60K plate for 5 min

• Let film to cool down in boiling water for 1000 times for 40-60 min

8/21/00



The PCR of control did not show up in the gel.

- Might have failed to ~~start~~ ^{work} as the ~~gel~~ ^{gel} or might have ~~not~~ ^{not} failed anyway. All.

- Need to do the PCR positive (+) and negative (-) control again.

Run gel again 8/21/00

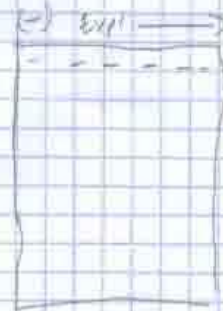
- buffer
- ADV EXP1
- ADV (-) control (uncontaminated)

come out good! see pic

2 ml. buffer 1000x → 200x

Load for 1st isolation

1 tube	5x tubes
1.5 ml (-) ADV	3.5 ml EXP1
1.7 ml dH ₂ O	1.7 ml dH ₂ O
1.8 ml dH ₂ O	1.8 ml dH ₂ O
15 ml ALLIS.	



Fuse 55
MC1061

Tetracycline resistance ?

MC1061 - warm, uninfected female.

using the DH5 α - already got many chromosomal competitors.

make LB fix plates.

500 μ l dH₂O
+ 17.5 g LB Agar

• no clare @ slow for 20 min.

Rapid Gel solution protocol

$$\#1 = 576.8 \text{ g} \left(\frac{100 \mu\text{L}}{1 \text{ g}} \right) = 576.8 \mu\text{L}$$

$$\# = 316.4 \text{ g} \left(\frac{100 \mu\text{L}}{1 \text{ g}} \right) = 316.4 \mu\text{L}$$

GeneClean
Turbo Silt
Solution

total for
100 μL
316.4 μL

- 1) incubate @ 55°C for 5 min.
- 2) transfer 200 μL @ 200 to column and let for 5 seconds.
discard waste.
- 3) add 200 μL of GeneClean Turbo Wash to flow
column for 5 seconds
discard waste.

centrifuge for 4 min.

- 1) New collect tube !! transfer centrifuge
add 30 μL GeneClean Turbo Elution.
incubate @ room temp for 5 min.
- 2) centrifuge for 1 minute and cap collect tube.

Standard Transformation
protocol for the DH5 α

Pre-warm plasmid to 22°C before starting

Single Tube Aliquots

- 1) Add 1-5 μ l plasmid DNA to a thermal cycler tube on ice.
Mix gently for a few seconds.
Try to keep the volume of DNA less than 5% of total.
- 2) Incubate on ice for 2-5 minutes (don't bother)
- 3) Spread 50-100 μ l into a pre-warmed culture plate. Incubate the plate at an appropriate temp (37°C) for the culture to grow.

2 ml DNA

1 ml DNA

1.66 ml dya

1.66 ml dya

11.33 ml

~~11.33 ml~~ 12.33 ml

$$(10x)(L) = (1x)(15) + L$$

$$9x = 15$$

$$(375) = (500 \text{ ng}) \cdot 1x$$

375x 15 mg/ml Tet stock
40 x/ml plates

$$\frac{50 \text{ ml}}{375x} = 0.133 \text{ ml} = 133 \mu\text{l}$$

$$= 1.33$$

us/m = 8

NEW ENGLAND BioLabs $\frac{133 \mu\text{l}}{50 \text{ ml agar}}$

Introducing Crimson™ Taq

133 μl
50 ml
m/plates

10.66 ul

9/25/07

Bedding system with fresh RE digest of $\text{C}_2\text{H}_5\text{OH} - \text{P}_2\text{O}_5 - \text{GF}$
of BSE and fresh digest $\text{C}_2\text{H}_5\text{OH}$.

5/25/09

Amended the digest

100 x 100 Volume

3.2 x digest A (1:100) x 100

3.2 x digest B (1:100) = 100

30.2 Six samples buffer

73.6 x H_2O

100 x

PH $^{\text{H}}$ buffer water @ 3.08 pH
and 4.08 pH

5/25/09

get some more colonies, making whole eggs.

Reference digest
 for ADV - RSVa - GFP - US

	(-)	(E)	(+)	(-) RSVa	total lanes with BASE
ds ADV-RSVa OFF-US	1ul	10ul	-	-	
Reference	2ul	2ul	2ul	2ul	new format of bitter #2 open @ 1/25/09
RSV	-	2ul	2ul	-	
RNAse inhibitor	-	-	4.5ul	1.9ul	
dH ₂ O	17ul	6ul	11.5ul	7ul	

incubate @ 37°C for 60 min.
 spin 135g x 2
 @ 2:55 p.m.

check digest with a Gel.
 from Gel to isolate

isolation

Plate Counts
for E4 on 8/21/09

Ex 55 plate
1/2 of LB from TET

Good + Amp 500 \rightarrow colonies \approx 100

10:1 dilution

3:1 $\sim \times 2$

10:1 $\sim \times 3$

20:1 $\sim \times 2$

50:1 ~ 1

Good plate

LB + Amp

each tube ~ 3 ml
of LB + Amp.

Amp [100 mg/L]

need 600 μ g of Amp for 30 ml of LB

[100 mg/L]

30 ml, 600 μ g

$$(100 \text{ mg/L}) = (30 \text{ ml}) \left(\frac{20 \text{ mg}}{30 \text{ ml}} \right)$$

3 ml each need ~ 30 ml

10

600 μ g

$$3 \text{ ml} \text{ (LB)} = (20 \text{ mg}) / (10)$$

Note You can see the LB colonies plate for about a month

You can grow the LB liquid cells for about 1-2 ^{generations} ~~days~~

Protocol for Mincip

- From what I read into a $\approx \frac{3}{20}$ L of LB with amp
which temp to be $\approx [100 \text{ mg/L}]$
since is 500x.
So have to 30 ml of LB.
Incubate in shaker for 16-18 hours
started around 3:00 pm.

@ 2:40 PM

FUSE 515 (conting)

- in 3ml of LB plus tet. Some want tet to be [] as plates
pH 2-3 colonies.
I need a \approx ~~10~~ 10 mL LB + 26.667 mL of tet.

Incubate @ 37° 16-18 hrs

Start @ 3:49 pm

@ 2:40 PM

Next day - ^{cells} centrifuge $\approx 7000 \text{ rpm}$ for 3 min, to get a pellet.
Use both 1.5 tubes and transfer the amount
one by one.

- Note: when tube is frozen, it's a good sign that
your cells grow.

8/25/08

Agar Gel isolation

$$\#1 = \frac{.4813}{.15} = 480.3 \text{ mL}$$

$$\#2 = \frac{.5657}{.15} = 565.7 \text{ mL}$$

of Gene Chem
rebo
See 9/10

$$\#3 = \frac{.3818}{.15} = 381.8 \text{ mL}$$

1) Incubate @ 55°C for 5 min

2) Transfer 1000 µL to column - centrifuge for 5 seconds.

MiniPrep Protocol 8/26/09

1) Resuspend pellet bacterial cells in 250 μ l Buffer P1

- Need to add Lysozyme to buffer P1
- No cell clumping - should be visible after resuspension.
↳ pipette up & down
- If Lysozyme storage has been added to P1
make sure it's well shaken.

2) Add 250 μ l Buffer P2 & mix thoroughly by pipetting
tube 9-6 times, (close bottle immediately)

- Mix gently + don't vortex - until it becomes
slightly clear or viscous.
- Don't allow cells to lyse for more than 5 min.

- Should turn blue if Lysozyme was added to P1

3) Add 350 μ l Buffer N3 & mix immediately & thoroughly by
inverting the tube 4-6 times

- If Lysozyme was used, mix until it's colorless.
Suspension should be colorless.

4) Centrifuge for 10 min @ 13,000 rpm

- Compact white pellet will form

5) Apply the supernatant from step 4 into the QIAprep

Spin column by decanting or pipetting. (750 ml)
don't get greedy
spin from top → bottom

6) Centrifuge for 30-60s (13,000 rpm). Discard the flow through.

7) Wash the QIAprep spin column by adding 0.5 ml Buffer PB
and centrifuging for 30-60s. Discard the flow through.
(Not needed for DH5α)

8) Wash QIAprep spin column by adding 0.75 ml
Buffer PE & centrifuge for 30-60s. (13,000 rpm)
(Need to add Ethanol)

9) Discard the flow through & centrifuge for an additional 1 min.
to remove residual wash buffer

10) Place the QIAprep column in a clean 1.5 ml tube.
To elute DNA, add 50 μl Buffer EB (low salt Tris) or water to the center of each QIAprep.
Spin column, let it stand for 1 min,
and centrifuge for 1 min.

Note: Lys. Blue to Buffer PE

1:1000 dilution

8/26/00

Note: Accidentally added twice as much of CB buffer (solution)

to the 20:1 mix (AAV ^{100%} ~~100%~~) mix.

lanes get for mixtures:-

1.0L of each (except 20:1 tube 2.0L)

5.0L total

2.0L total

1.5L total

12.5L total

15.0L

latter	500	500	500	300	300	100	100	200	500
21	22	23	24	25	26	27	28	29	30

100g, 100g, 100g, 100g, 100g, 100g

8/27/09

Checking FUSSES and ADV format
with a digester

FUSSES format

- going to identify it again (FUSSES wrapper) with
a limbath letter

Alpha

FUSSES		
#1	#2	#3
Sak	Sog	Soo
1.5dga	1.5dga	1.5dga
B:5:u:u:u:u	B:5:u:u:u:u	B:5:u:u:u:u
1.5ak	1.5ak	1.5ak

BASE GAMA digester

with NdeI and MclI to check for stLNA

160
200

insert
about 900 bp
+ 165bp band for no insert

Table

	5.1.11	3.7.11	10.1.11	10.1.13	24.1.12	30.1.12
	112 bp	122 bp	120 bp	120 bp	60 bp	60 bp
	(-) Ew	(-) Ew	(-) Ew	(-) Ew	(-) Ew	(-) Ew
Insert	100	100	100	100	100	100
BaseGAMA	200	200	200	200	200	200
NdeI	-	-	-	-	-	-
MclI	-	-	-	-	-	-
11.1.12	170	170	170	170	170	170

Note

(2017-2018) Micrograph -
20% glycerol for 10^6 - 10^8 cells

Since glycerol is very dense ex. used for all for that
use round 2.25 ml of glycerol and 10 ml for 10⁸ cells

↑ total sample = ~~200~~ 100 ml

Since 14 glycerol for future work

Store @ -80°C

↓

Then centrifuge the whole 500 ml
in the jumbo centrifuge

8/25/04

Final SS (Mimicry digital 6)

Year = 4.44 dpc
Total = 7.22 dpc

total = 1.5 dpc
(1) ~~to~~ attach
12.5

	#1	#2	#3	#4	#5
1/23/04	(1) Ep	(2) Ep	(3) Ep	(4) Ep	(5) Ep

8/25/04

Yesterday these 5 tubes Final SS involved last gel was
incubated because the gel was probably made (well where some)

• you want of Final SS #1 & #2 for comparison

total of both (from previous experiment tubes)

Time of LB with top (young/old)

• ~~to~~ ^{to} attach JTC

Today - you ~~will~~ stopped incubation in Final SS #1 & #2

incubated for 16 hours

Results: do comparison and figure out see if I get

with Final SS

• this when Ready = good with gel growth

8/28/04 FUSESS Classification
of stimp 8/28/04

using the formula below and Sub of each
stimp on a 19. August gel

FUSESS - stimp (15 ul total)

	#1	#2	#3.1	#3.2	#3.3	#3.4
DATA	Sub	Sub	Sub	Sub	Sub	Sub
10V	1.5 ul	1.5 ul	1.5 ul	1.5 ul	1.5 ul	1.5 ul
11V	8.5 ul	8.5 ul	8.5 ul	8.5 ul	8.5 ul	8.5 ul
	→ 500 ul	2.500 ul	500 ul	2.500 ul	500 ul	500 ul
	200 mg	500 mg	200 mg	100 mg	100 mg	200 mg
	100 mg	100 mg		100 mg		200 mg

9/3/09

Success digestion w/ NsiI and NdeI done on 7/1/09

Success of NsiI

Positive controls w/ PfuII plasmid = 1500bp

	#1		#2		#3.1		Positive controls w/ PfuII plasmid = 1500bp	
	(-) EXP	(+) EXP	(-) EXP	(+) EXP	(-) EXP	(+) EXP	(-) PfuII	(+) PfuII
plasmid	4ul (100%)	16ul (100%)	8ul (100%)	32ul (100%)	4ul (100%)	20ul (100%)	1ul	1ul
Buffer M3 (20x)	2ul	4.5ul	2ul	4.5ul	2ul	4ul	2ul	2ul
NsiI (100x)	—	.18ul 1.8ul	—	.18ul 1.8ul	—	.2ul 2ul	—	.3ul 2ul (1:100)
dH ₂ O	14ul 14ul	2.7ul 16ul 20.7	12ul 12ul	2.7ul 19ul	14ul 14ul	2.7ul 19ul	17ul	14ul 14ul

Expected sizes = 7517 bp
1684 bp

Start 4pm

Success of NdeI

Positive controls w/ PfuII plasmid = 1500bp

	#1		#2		#3.4		Positive controls w/ PfuII plasmid = 1500bp	
	(-) EXP	(+) EXP	(-) EXP	(+) EXP	(-) EXP	(+) EXP	(-) PfuII	(+) PfuII
plasmid	4ul (100%)	16ul (100%)	4ul (100%)	16ul (100%)	4ul (100%)	20ul (100%)	1ul	1ul
Buffer M4 (20x)	2ul	5ul	2ul	5ul	2ul	5ul	2ul	2ul
NdeI (200x)	—	.07ul .7ul	—	.07ul .7ul	—	.18ul 1.8ul	—	.15ul 1.5ul (1:10)
dH ₂ O	14ul 14ul	2.3ul 14.7ul 24.1	14ul 14ul	2.3ul 14.7ul 24.1	14ul 14ul	2.3ul 14.7ul 24.1	17ul	14ul 14ul

Expected band sizes = 4300, 3150, 1281, 675

8/31/09

Strep. Avidin - IGen Zoon Box well 9P
transformant (536816)
in DHS dx cells

100 ul of DHS dx cells

3 ul of DNA

plate on Amp LB plates, incubate @ 37°C

incubated @ 6.10 pm @ 37°C

Colony 9/1/09 - many colonies 1,000's !!

~~Start~~ and ~~start~~

~~later~~ ~~start~~ ~~start~~

Colony on plate - 8 for cultured in grow

exactly on LB + Amp

Start @ 5 pm

9/2/09 - nice cloudy tubes - good - for the strep Avidin

inhibition tubes.

- doing stripup of dem 9/2/09.

val. day

7/2

9/2/09

- did Miniprep of the strip Anilin.
- Ran gel to check Fusc SS w/ NSEI
 - seems that NSEI did not cut
 - but it did cut controls, - DNA might be too strong?

- Running gel of Fusc SS w/ NSEI

- Running gel to quantify the strip fig
- mini prep - running lab of each sample

Strip Anilin

#1

#2

#3

$\frac{20}{4} = 2.22 \text{ dpc}$

50 ng/ul

100 ng/ul

↑ 100 ng/ul

$\frac{95}{4} = 5.16 \text{ dpc}$

Tamara - do several samples of the sheep adhesion cells.

9/3/09

Agarose to do to ^{the} original dsDNA (5200 bp)

reads the ^{the} what w/ (+) reads
to try further +2 and +4.

- do ^{table} digest with ^{the} what and what - should digest - ^{the} what band.

- do ^{the} the same Agarose - Same I and Kpn II to

check FFP's

↑
integrity of ^{the} what

Swit should be incubated @ 25°C !.

Expected bands

w/ Kpn I

- 3,062 bp

- 153 bp

- 13.71

Sam I

9,495

220

W & what be the as see it

what

FW 17,510

1.82

padding

74

200

FW 9,300

2.22

400-400

9/3/14 Single digest of dsDNA (original)

w/MS Sma I ~~and~~ ~~PCDN4~~

Sma I - incubated @ 25°C

PCDN4 = 110 ng/ul

	(-) dsDNA	Exp Sma I	(-) PCDN4	(+) PCDN4
Plasmid	1 ul	2 ul	1 ul	1 ul
Buffered H ₂ O	2 ul	2 ul	2 ul	2 ul
Sma I	—	.02 ul [.4 ul (20)]	—	.011 [.22 ul (110ng)]
H ₂ O	17 ul	15.8 ul	17 ul	16.89 ul

~~20 ul~~

20 ul rxns

Expected band sizes

dsDNA

4,475

720

11 ~~6~~ ~~kb~~ ~~bp~~
with 20 rxns

PCDN4

~~98~~

~~98~~

98

22

incubated @ 25°C for 1 hour

start: 3:03 PM

end: 3:24 PM

9/3/04

Proble Lösung mit Adel und Xhol
 als ADV (Lsgung)

positive Werte

	ADV				positive Werte		
	(-) als ADV	Exp. Wert	Exp. Xhol	Restwert	(-) P1	Wahl Puff II	Xhol Puff I
Adel	1ul	1ul	1ul	10 ul	1ul	1ul	1ul
Yffer	2ul	2ul	2ul	2ul	2ul	2ul	2ul
Xhol	1ul	1ul <u>1ul (1:10)</u>	—	1ul	—	1ul <u>1ul (1:10)</u>	—
Xhol	—	—	0.5ul <u>0.5ul (1:10)</u>	0.5ul	—	—	0.75ul <u>0.75ul (1:10)</u>
H ₂ O	17ul	16ul	16.5ul	6.5ul	17ul	15.5ul	16.25ul
Bla	1ul	1ul	1ul	1ul	1ul	1ul	1ul
Zug							
Wahl							

Wahl 1. Zugwert = + 110 kg

in 29 gel.

Xhol Puff I - exp. = 9,300

2,400

600

Wahl II = 2,5 kg gel II

Proble e 370 für 1600

9/3/09

Single digest of λ DNA (orig) of KpnI

	(-) λ DNA	Exp KpnI	(-) λ DNA	(+) KpnI
Phosmix	1 μ l	3 μ l	1 μ l	1.5 μ l
Buffer #1	2 μ l	2 μ l	2 μ l	2 μ l
BsaI	2 μ l	2 μ l	2 μ l	2 μ l
KpnI	—	3 μ l	—	2.25 μ l
		3 μ l (1/10)		2.25 μ l (1/10)
Water	10 μ l	12 μ l	10 μ l	13.25 μ l

20 μ l rxn

Expected bands	λ DNA	KpnI
	5,682	10,500
	153 (avg)	4,600 (20 ng)
	13.71	

incubated @ 37°C for 1hr

9/1/09 Friday - Check double plotted IPR's
 again w/ Sam I - this time 200
 m. Little more and hope to see the 700' dip bands.

- do - double dipping at the ADU airpops
 with N40E and S40E

Small dipper dddv (9/1/09)

	(-) dddv	Small dddv	(-) pcdv	(+) pcdv
Plasma	1UL	1000UL	1UL	1ab
* Butler & H	2UL	5UL	2UL	2UL
Sam I	—	1UL 1UL (1200)	—	2UL 2UL (1200)
dlho	17UL	34UL	17UL	16.7E46

Start 1:30 pm
 end 2:50 pm

(4)

9/9/09

Sitosterol α and β

where transfection in DM5.0X cells

Seal cells for each to 50 cells for each no plate

- where plated @ 4700 pm

incubated @ 37°C

9/9/09 - 10:10 AM

• Transfer 3 volumes of each plate (sitosterol α & β plates) in CB

• Run gel of double marker double digest in 2.5% gel

• do mapping of the fuses

Sitosterol α and β

• 3 volumes of each plate

in 3 ml of CB and Amp

$$\begin{aligned}
 (30 \mu\text{l}) \times 1 &= (1 \mu\text{l}) \times \frac{20}{100} \times 100 \\
 &= \frac{20}{100} \times 100 = 20 \mu\text{l}
 \end{aligned}$$

incubated @ 37°C for 1 hr 1:30 pm

Note - result gel if you're going into the next day

- did mapping of fuses 9/9/09 → 2:00

- for double digest of DM5.0X mapping 9/9/09

11:26 17:26

1/1/16

UG

Things to do today

- Mumpage & Silvestri at end B

- Loose bits of LADAV on SHANA waives again

↳ this is a list of things LADAV original found
and make the same
25-30.06 20, time

- to figure out (GROSS) waives w/ what soil and RICE

LADAV - SHANA Waives Report on - 4 p. gal

	1	2	3	4	5	6	7	8	9	10
	2.1	2.1	2.1	5.1	5.1	5.1	10.1	20.1	20.1	20.1
	Exp1	Exp2	Exp3	Exp1	Exp2	Exp3	Exp1	Exp2	Exp3	LADAV
found	17.5g	20g	5.5g	10g	10g	10g	10.2g	10g	10g	15g
difficult	7.5g	10g	10g	7.5g	7.5g	7.5g	7.5g	7.5g	7.5g	7.5g
before										
3/10	2.5g	2.5g	2.5g	2.5g	2.5g	2.5g	2.5g	2.5g	2.5g	2.5g (1.100)
1/20	1g	1g	1g	1g	1g	1g	1g	1g	1g	1.5g
2/1	.5g	.5g	.5g	.5g	.5g	.5g	.5g	.5g	.5g	.75g
1/10	1.25g	1.25g	1.25g	1.25g	X	1.25g	1.25g	1.25g	1.25g	1.25g

Master mix (10.22g)

BA 2.5g

difficult 7.5g

1/20 10g

1/1 5g

4.25g for each tube

except for 10.22g

Experiments

11/24/04

Original dHAP

pl	Standard	Final total dHAP	100% total dHAP
plasmid	1ul	1ul	1ul
DNA	2ul (100)	2ul (100)	—
Buffer 10x	2ul	2ul	2ul
H ₂ O	X	X	1ul (100ul)
total	X	5ul (100ul)	X
Alto	15ul	14.5ul	17ul

Start @ 3:10pm @ 37°C

end 4:10pm

Get wells on 4th Agmax gel 15x100, 1.5/ul E+8x

- total dHAP
- 3.1 ul
- 3.1 #2
- 3.1 #3
- 3.1 #4
- 3.1 #5
- 3.1 #6
- 3.1 #7
- 3.1 #8
- 3.1 #9
- 3.1 #10



plasmid

Buffer 10x

H₂O

Alto

9/18/08

- To do
- check for SS λ map
- check the fixation of and selection β with digesting for conversion line
- repeat fixation the subHA-delta get again

70.5

gross map date by Terry & Justin

7/24/08

digest w/ HpaI

	(-) gross	gross λ map	- PCR	(+) PCR
plasmid	1.2 ul (20 ul)	1.2 ul (20 ul)	1 ul	1 ul
Buffer #4	2 ul	2 ul	2 ul	2 ul
HpaI	—	0.4 ul 1.4 ul (1'10')	—	0.3 ul 3 ul (1'10')
Algo	16.5 ul	4 ul	17 ul	17 ul

7/24/08

incubate @ 37°C for 1 hour

digest w/ HpaI

	(-) gross	gross λ map	- PCR	(+) PCR
plasmid	1.5 ul	17.5 ul	1 ul	1 ul
Buffer #4	2 ul	2.3 ul	2 ul	2 ul
HpaI	—	0.5 ul 3.5 ul (1'10')	—	0.3 ul 1.5 ul (1'10')
Algo	16.5	—	17 ul	17.5

incubate @ 37°C for 1 hour

Part of design and construction

low to high for POK

low - 25 %

allow for delays - 20

to miss allow

5% → material of work

4% of the program

in the 5% of the program

4% more BCB

material - in the program

allow for 3 days of material at work

in program

7/1/20

~~100~~ 100% - 100% of the ...
 - direct ...
 -

100% - 100% of the ...
 ...

100% - 100% of the ...
 ...

...

$$100 \times \left(\frac{100}{100}\right) \rightarrow \left(\frac{100}{100}\right) = 100\% \quad 100\% = 1.00$$

	100%
100%	100%
100%	100%
100%	100%
100%	100%

...

11/14/20

Let's look at the situation of the market → Early

Interest rate = 5% per year

Situation of the market

Present value = 110 per year

Situation of

Present value

	(+) PV	Exp PV	(-) PV	Exp PV	(-) PV	Exp PV	(+) PV	(-) PV
Present	100	2.706 (1500)	100	2.706 (1500)	100	2.706 (1500)	100	100
After 1st	100	100	100	100	100	100	100	100
2nd	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)
3rd	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)
4th	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)
5th	100	4.55	100	4.55	100	4.55	100	6.45

interest rate 5% per year expected value 5.5% and 1.5%

Situation of the market

Interest rate 5.5%, 70%, 1.5%

	(+) PV	Exp PV	(-) PV	Exp PV	(-) PV	Exp PV	(+) PV	(-) PV
Present	100	4.106 (1500)	100	4.106 (1500)	100	4.106 (1500)	100	100
After 1st	100	100	100	100	100	100	100	100
2nd	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)
3rd	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)
4th	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)	100 (1100)
5th	100	3.156	100	3.156	100	3.156	100	7.00

2/5/20

- Things to do
- report for 2020/2021
- list of 2020-2021
- ✓ - report of 2020-2021 & 2021-2022
- ✓ - list of 2020-2021

- 2020-2021 - get a loan - 2020-2021 - 2020-2021 - 2020-2021 - 2020-2021
to provide 2020-2021

- 2020-2021 - 2020-2021

Solution of P.P. data after first order

1. Jan
 #100 }
 #100 }
 #100 }
 #100 }
 #100 }
 #100 }

#100 }
 #100 }
 #100 }
 #100 }
 #100 }
 #100 }
 #100 }
 #100 }

Solution of
 under the first order

DATE	YAC	YAC
200	YAC	YAC
100	YAC	
100	YAC	
100	YAC	
100	YAC	
100	YAC	

7.500
 7.500
 7.500

Solution of
 under the first order

DATE	YAC	YAC
200	YAC	YAC
100	YAC	
100	YAC	
100	YAC	
100	YAC	
100	YAC	

6.600 to 7.500

1000 to 1000

4/15/20

Solution of #1 done at NAE and NAE

	C-101	#102	C-103	#104	#501	#105	FJ 106	NAE 107	NAE 108
plasma	1ul	2.7ul	1ul	2.7ul	1ul	2.7ul	1ul	1ul	1ul
plasma 2	1ul	1ul	1ul	1ul	1ul	1ul	1ul	1ul	1ul
plasma (100%)	1ul	1ul	1ul	1ul	1ul	1ul	1ul	1ul	1ul
NAE (100%)	—	1.7ul	—	1.7ul	—	1.7ul	—	1.5ul	—
NAE (100%)	—	1.5ul	—	1.5ul	—	1.5ul	—	—	1.1ul
plasma	7ul	3.05ul	7ul	3.05ul	7ul	3.05ul	7ul	6.75ul	5.9ul

include a 5% for loss

total 5.8ul and 1.7ul 5.8ul 5.8ul 5.8ul 5.8ul 5.8ul

NAE 100%

Solution of #2 done at NAE and NAE

	C-101	#102	C-103	#104	#501	#105	FJ 106	NAE 107	NAE 108
plasma	2ul	9.4ul	2ul	9.4ul	2ul	9.4ul	2ul	2ul	2ul
plasma 2	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul
plasma (100%)	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul
NAE (100%)	—	1.5ul	—	1.5ul	—	1.5ul	—	5.5ul	—
NAE (100%)	—	1.5ul	—	1.5ul	—	1.5ul	—	—	1.1ul
plasma	10ul	1.7ul	10ul	1.7ul	10ul	1.7ul	10ul	1.75ul	1.9ul

no need for 100% and 100%

to be done

9/19/18

- Tides in the Gulf
- Highs & Lows in the Gulf - again at different times
- ...
- ...

Principle #3 for deep-sea

- good one

#1 Soil - 100m

Soil - 100m

100m #1 Soil 100m

Deep-sea - 100m - 100m 9/19/18

#1 - 100m

#2 - 100m

#3 - 100m



Yes, the soil is good

8. Beginn mit Sonne → Zeichen der Zirkulation und ...

... in ...

q11/08 Energy 1100-1600-1600-1600 map

with 200 I and 400 I

400 I sign of 1100-1600-1600-1600

	#10	#100	#100	#200	#20	#500	#1000	#1000
planned	100	300 (1000)	100	300	100	300	100	1000
Butter #1	100	100	100	100	100	100	100	100
BSA (100)	100	100	100	100	100	100	100	100
400 I (100)	—	400	—	400	—	400	—	2000
also	700	4000	700	4000	700	4000	700	4000

more 400 I
1000

more 400 I
400

1000 - 400
1000 - 400
400 - 1000
1000 - 1000

1000 - 400
1000 - 400
1000
1000 - 1000

inward @ 300
e 2000
e 4000

add 700 to each
side

add 700 to
each side

11/1/00

Sum 2 Journ of ADA-RNA-GP-UK

	11/1	11/2	11/3	11/4	11/5	11/6	11/7	11/8
planned	1uc	1uc	1uc	1uc	1uc	1uc	1uc	1uc
buffer #1 all 7/8	1.5uc	1.5uc	1.5uc	1.5uc	1.5uc	1.5uc	1.5uc	1.5uc
sum 2 (11/1)	—	1uc	—	1uc	—	1uc	—	1.5uc
11/2	1.5uc	2.5uc	3.5uc	4.5uc	5.5uc	6.5uc	7.5uc	8.5uc

11/1/00
11/2

11/1/00

incubated @ 25°C
@ 3:17 PM
end 4:20 PM

Master mix S-L-S

11/1/00

buffer #1
5uc
sum 2
(11/1)
1uc
11/2
1.5uc

buffer #1
5uc
11/2
50uc

add 5uc to each tube

add 10uc to each tube

Silicon α & β alloys 1/1

Alloy and steel

9/14/00 - did not have time to finish silicon alloy work
found it interesting

β expand	α expand
5318	1174
7450	8215
	309
<hr/> 6765	<hr/> 6753

I only want 1000 to the
American

water with my own hands

7

00

1/1		00		00	
Alloy	Steel	Alloy	Steel	Alloy	Steel
Alloy	Steel	Alloy	Steel	Alloy	Steel
Alloy	Steel	Alloy	Steel	Alloy	Steel
Alloy	Steel	Alloy	Steel	Alloy	Steel
Alloy	Steel	Alloy	Steel	Alloy	Steel
Alloy	Steel	Alloy	Steel	Alloy	Steel
Alloy	Steel	Alloy	Steel	Alloy	Steel

Lead 1000 Lead 1000

4/10/88

High to Shaker at end of reaction

- on what and what else is

Start Shaker at end of what

	(-1) 42	#1 Eng	(-1) 42	#2 Eng	(-1) 42	#3 Eng	(-1) 42	(-1) 42	(-1) 42
1st	2ul	7.6 (200)	2.4	7.6 ul (200)	2ul	7.6 ul (200)	2ul	2ul	2ul
2nd	2ul	2ul	2ul	2ul	2ul	2ul	2ul	2ul	2ul
3rd	2ul	2ul	2ul	2ul	2ul	2ul	2ul	2ul	2ul
4th	—	2.5ul	—	2.5ul	—	2.5ul	—	2.5ul (100)	—
5th	—	2.5ul	—	2.5ul	—	2.5ul	—	2.5ul (100)	—
6th	10ul	4.75ul	1ul	4.75ul	1ul	4.75ul	7ul	9.5ul	10.5 ul

4.1.1 @ 1.75 ul

Substrate 2/10/88

Only added water to the reaction mixes
and 2ul to the 5th (100)

all tubes had DNA during

Shaker at end of what

	(-1) 42	#1 Eng	#2 Eng	#3 Eng	#4 Eng	#5 Eng	(-1) 42	(-1) 42	(-1) 42
1st	2ul	14ul	2ul	14ul	2ul	14ul			
2nd	2ul	2ul	2ul	2ul	2ul	2ul			
3rd	2ul	2ul	2ul	2ul	2ul	2ul			
4th	—	1.5	—	1.5	—	1.5			
5th	—	2.5ul	—	2.5ul	—	2.5ul			
6th	15ul	—	15ul	—	15ul	—			

100 100 100

Monday 11/11/04

311-1 100mg
311-2 100mg
1011-1 100mg
1011-2 100mg
1011-3 100mg
1011-4 100mg

2/3 8/16/04 skipped

311-1 50mg
311-2 100mg
311-3 100mg
1011-1 100mg
1011-2 100mg
1011-3 100mg
1011-4 100mg
1011-5 100mg
1011-6 100mg
1011-7 100mg
1011-8 100mg
1011-9 100mg
1011-10 100mg
1011-11 100mg
1011-12 100mg
1011-13 100mg
1011-14 100mg
1011-15 100mg
1011-16 100mg
1011-17 100mg
1011-18 100mg
1011-19 100mg
1011-20 100mg
1011-21 100mg
1011-22 100mg
1011-23 100mg
1011-24 100mg
1011-25 100mg
1011-26 100mg
1011-27 100mg
1011-28 100mg
1011-29 100mg
1011-30 100mg

11/15/04

only the green ones for
check the 10/11/04

10/11/04
10/12/04
10/13/04
10/14/04
10/15/04
10/16/04
10/17/04
10/18/04
10/19/04
10/20/04
10/21/04
10/22/04
10/23/04
10/24/04
10/25/04
10/26/04
10/27/04
10/28/04
10/29/04
10/30/04
10/31/04

✓

✓

✓

✓

✓

✓

Started at 4:00pm took 100mg of 1011-1
and 100mg

8/19/09

- slide with microscope & straight window - check
press

20 - 100

50 - 100

100 - 100

50 - 100

also found some

in glass at 20°C

place in the 2000 year steel
box

9/23/09 - day 10 of the for dithio - sheet

- state some more and that was the needed before

for the linkage on Sunday

- Need to check dithio sheet with slit

to check 100's

Jan 1 - 105 - 6:10

Jan 2 - 5000 5:11

Jan 3 1000 - 6:00 3:00 1:00

Jan 4 - 15.3 10:00 10:00

15

10

2:00 10:00

	(+) S.I.	Exp S.I.	(-) Tail	Exp Tail	(-) S.I.	Exp S.I.	(-) Tail	Exp Tail	(-) S.I.	Exp S.I.
Jan 1	105	105	6:10	6:10	2011	2011	2011	2011	105	105
Jan 2	5000	5000	5:11	5:11	15	15	15	15	5000	5000
Jan 3	1000	1000	6:00	6:00	3:00	3:00	3:00	3:00	1000	1000
Jan 4	15.3	15.3	10:00	10:00	10:00	10:00	10:00	10:00	15.3	15.3
Jan 5	10	10	10:00	10:00	10	10	10	10	10	10
Jan 6	10.5	10.5	12:3	12:3	10.5	10.5	10.5	10.5	10.5	10.5

10:00

Start at 3:16pm

end 8:16pm

9/2/08

Empty gal of the steel I digest

of the 1200V - 240V

in 100 gals

checking for 240V - 240V or in 1000 gal.

1000 - 1000
1000 - 1000

1000 - 1000 original 240V and

silicon K and B in glass for

burn out.

3000 - 6110

1.5 1000

100
50%

1000 5667524 to 9.1000 = 1.7502 x 10⁵ pules

1000 -

5.2750 x 10⁵ pules

1000 1675581 27.0000

2710.00 - 500

500.00

need 1000 of 1000

5.2750 x 10⁵ pules

need 1000 of 1000
= 1.503 x 10⁵ pules

need 80.52 mg of lead for

1.311 ton.

1000 mg



9/2/04

Basal

Basal magnet 20mg/ok

Basal with Magnesium - then add phosphate

	(-) Basal	Basal (10mg)	Basal (10mg)	Basal (10mg)	Basal (10mg)
Basal	2ul	10ul	10ul	10ul	10ul (10mg)
Basal + P	2ul	2ul	2ul	2ul	2ul
Basal (10mg)	2ul	5ul	2ul	2ul	2ul
Magnesium (10mg)	—	2ul	2ul	—	1ul
Basal	15ul	1ul	1ul	15ul	15ul

Start 5:15 PM
end 6:15 PM

9/6/04
start 10:30 PM
end 11:30 PM

Basal

9/6/04 - Magnesium

Basal @ 3% for 1hr

Basal with Magnesium - then add phosphate

	(-) Basal	Basal (10mg)	Basal (10mg)	Basal (10mg)
Basal	1ul	5.7ul	1ul	1ul (10mg)
Basal + P	1ul	2ul	2ul	2ul
Basal	—	2ul	—	1ul (10mg)
Basal	17ul	11.6ul	17ul	17ul

Start 5:15 PM
end 6:15 PM

Start 5:00 PM
end 6:15 PM

Let Atomic phosphorus and Atomic Phosphorus

	Exp #1	Exp #2	Observation
Phosphorus	20ml	20ml	
Test tube (100) → (10)	2.53 ml	2.37 ml of Atomic Phosphorus	
Alkaline Syringe	1ml (10)	1ml (10)	
Atomic phosphorus	—	$\frac{3}{8} \frac{1}{100}$	Wash for 15 min to 27°C

→ 0.27% 60 min

for digest on = 2.5% gel with/water bath

purify and get ~~100%~~ yield

expected bond = 2.2%

Expected bond for an atomic phosphorus is with P(VI)

→ $\frac{2.75}{100}$ and 2.53%

and the 2.75%

for 10% = 2.9% gel

gel
100%

9/23/07 Sun

Shrimp - start @ 1:15 pm

Start to
demon @ 1:55 pm @ 1

1.5 years of Aquaria tank

300 ml of .5% gel.

First digest @ 1:55
gel

Water	Clorox	(1) 1000 ml	(2) 1000 ml	Amia	Shrimp
		1	2	1	2

Get initial

SAP

Aquaria

#1 ~~402.5~~ grams #2
402.5 478

~~398~~ #1 #2
398 304.5

$$402.5 \left(\frac{10000}{1000} \right)$$

$$398 \rightarrow \left(\frac{10000}{1000} \right)$$

~~402.5~~

402.5 g

478 g

398 g

304.5 g

402.5 g
478 g
398 g
304.5 g

1.5 years @ .5% for 5 ml

7/15/18m. NMR analysis of pure ST ligand. (Blends with ligand)

3:1 Ligand

	6) SDR	SAR 2nd	R-Mutants	Anticore Count
Core SS ligand	100 mg 1.5ul	100 mg 1.5ul	100 mg 1.5ul	100 mg 1.5ul
DRM ligand	—	80.5 mg 5.31 ul	—	80.5 mg 5.31 ul
DM ligand Buffer	2.5ul	2.5ul	2.5ul	2.5ul
TM SDR ligand	1ul (100u)	1ul (100u)	1ul (100u)	1ul (100u)
DMG	6.5ul	1.19ul	6.5ul	1.19ul

@ 9pm

Neutral to form comp (25%)

DRM ligand

3/19/04

Tide - transfer liquid into test plates

1st time test plates

2nd time - need 132 ml of test

from 10-15 ml of test/plate per each plate

- Note liquid - 100 mg total DNA used!!!

used from (2) for each of 11 plates

500 - for each - total 9 - 2.5 ml of liquid

minimum amount for 500

add 200 ml of LB and include from

to show for each plate

start @ 3:02 pm

end @ 4:03 pm

plates ~~500~~ @ 4:00 pm @ 37°C

$$(1000) = (1000)(1000)$$

500

(500)

500

Stratigone - 1000000 - 1000000 42

PCP - 1000000

gestirmt PCP

Sauk 1000

gestirmt PCP		Sauk 1000	
	AS Linder Kategorie		Handelsweg T ₀ #1 Kategorie
100	PCP - 1000 - 1000 - 1000 (1000)	100	PCP - 1000 - 1000 - 1000 (1000)
100	5x Phasen HP Bitter (1000)	100	5x Phasen HP Bitter (1000)
Sau	PCP - 1000	Sau	PCP - 1000
200	Handel 1000 5' (1000)	200	Handel 1000 5' (1000)
200	Spot 1000 5' (1000)	200	Handel 1000 inner 5' (1000)
200	Handel	200	Handel
200	Phasen 1000 (1000)	200	Phasen 1000 (1000)

4) 1000 1000 1000 - 1000 1000

1000 1000 1000 - 1000 1000		1000 1000 1000	
	Kategorie		Kategorie
100	PCP - 1000 - 1000 - 1000	100	Sauk 1000
100	5x Phasen HP Bitter (1000)	100	5x Phasen HP Bitter
Sau	PCP - 1000	Sau	1000 1000
200	Handel 1000 5' (1000)	200	1000 1000
200	Spot 1000 5' (1000)	200	1000 1000
200	Handel	200	1000 1000
200	Phasen 1000 (1000)	200	1000 1000

9/15/16

- Get very little market return for
the fixed SS - shifting plans

1 volume per year

- Need to take leverage, also time delay

to higher concentration rights and

low O/S !!

- looking ahead to all work of fixed SS w/ impact

- reading protocol on options

- lat. index left points

- 1000 - 1000 - 1.5 p

- 1000 - 1000 - 2.0 p

1.5 p - 4.5 p for 300 us

SAT - Right of fixed SS - right @ 2.5 p
at 1.5 p

Right of fixed SS @ 3.0 p - 3.5 p

at 3.0 p

9/25/14

TV Light - how SE SAP & Air - the SKEL-ETP

1:5 Light (1000:1 Lux) Overlight

	←) SAP	SAP SKE	→) Acto	2. Light SAP	← 16°C
Person	500g → 1.4 m	300g (1.3m)	700g SKE	500g (1.5m)	
Skull-mount SKE	—	26.6g	—	26.6g	
		1.59 m		1.59 m	
Better TV Light	2m	2m	2m	2m	
TV DVD light (Multi Color)	1m	1m	1m	1m	
Other	900g	11.1m	900g	900g	

x 25.6
1500 3920

1000 = 1000

1:5 1000 = 1000 1500 = 1500

1:1 1000 = 1000 1000 = 1000

9/26/01

PCR - stop - Herz water dishes

9/26/01

✓

	Reagents	50 ul rxns
1ul	Roche Amplicore - inner primer 1' (10ng/ul)	
10ul	Six Primase Ht BstKw (10/1000)	
5ul	2 mM dNTPs	
2ul	Herz 1/3hr 5' (10000 uM)	
2ul	Herz 1/3hr 3' water (10 uM)	
24.5ul	H ₂ O	
1.5ul	phosiz DNA polymerase (1/1000)	

- larger - large - short - transfection serum @ 25%

incubate @ 37°C for 1 hour

before plating

added 250 ul of LB to each tube
of 50 ul of cells

used 2.5 ul of PCR

9/26/01

9/16/08 - get values of 2-p/lines with 1st point

#1 = 39.19 gms
#2 = ~~39.19~~ 29.50
#3 = 19.99
#4 = 56.86

9/16/08 - do 1st 5 steps - then into the comp

- 2 values in the 1st column
- 2 values in the 2nd column
- 2 values in the 3rd column
- 2 values in the 4th column

9/16/08 - 2nd - 1st 5 steps - then into the comp

- check ~~values~~ (sum of 5 + that) ~~miniprof.~~
-

9/16/08 - get values

#1 = 33.10 >
#2 = 35.20 >

500
 1000
 1500
 2000
 2500
 3000
 3500
 4000
 4500
 5000
 5500
 6000
 6500
 7000
 7500
 8000
 8500
 9000
 9500
 10000

	Aug 1	Aug 2	Aug 3	Aug 4	Aug 5	Aug 6	Aug 7	Aug 8	Aug 9	Aug 10	Aug 11	Aug 12	Aug 13	Aug 14	Aug 15
Temp	100	105	110	115	120	125	130	135	140	145	150	155	160	165	170
Humidity	70	75	80	85	90	95	100	105	110	115	120	125	130	135	140
Wind	1.2	1.3	1.4	1.5	1.6	1.7	1.8	1.9	2.0	2.1	2.2	2.3	2.4	2.5	2.6
Pressure	13.6	13.7	13.8	13.9	14.0	14.1	14.2	14.3	14.4	14.5	14.6	14.7	14.8	14.9	15.0

Time	Temp	Humidity	Wind	Pressure
10:00	100	70	1.2	13.6
11:00	105	75	1.3	13.7
12:00	110	80	1.4	13.8
13:00	115	85	1.5	13.9
14:00	120	90	1.6	14.0
15:00	125	95	1.7	14.1
16:00	130	100	1.8	14.2
17:00	135	105	1.9	14.3
18:00	140	110	2.0	14.4
19:00	145	115	2.1	14.5
20:00	150	120	2.2	14.6
21:00	155	125	2.3	14.7
22:00	160	130	2.4	14.8
23:00	165	135	2.5	14.9
00:00	170	140	2.6	15.0

10:00
 11:00
 12:00
 13:00
 14:00
 15:00
 16:00
 17:00
 18:00
 19:00
 20:00
 21:00
 22:00
 23:00
 00:00

Started @ 5:00 pm

PCR setting

- 96°C → 2 min
- 96°C → 30 sec
- 55°C → 30 sec } 10x
- 72°C → 1 min
- 72°C → 7 min
- 1°C → 10 min

Pract II - 100V - 100mA



5/10/20

100V
100mA

100V

100mA

100V
(100mA)

100V

1/30/04
 $\lambda_{max} = 667.574, 70 \text{ nm}$

- 100% - digest protein with Bgl I in MS buffer 3 250
- 50% - digest protein - stop - dig with Bgl II - 200 bp - 5' to 3' (0.06-0.1)
- 25% - digest pUC19 as a control DNA with Bgl II

3.1. 2000

$$\frac{1000}{1000} = 40.791 \times 10^{-6} \text{ parts}$$

$$\frac{1000}{5000} = \frac{10.791}{5000} = 0.2158$$

$$\text{need } 2.6375 \times 10^{-5} \text{ parts of DNA} = 13.70 \text{ ng of}$$

Insert

	EG (mass)	protein Exp	Hex (c) 500	Hex (d) 500	hex (e)	hex (f)
100%	100ng (3ul)	100ng (1.5ul)	60ng (1.5ul)	100ng (2.5ul)	60ng (1.2ul)	100ng (2ul)
50%	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul	1.5ul
25%	-	2ul	-	2ul	-	2ul
10%	10.5ul	6.5ul	13ul	9ul	12.3ul	9.5ul

stirred @ 2.25 pm
 incubated 37°
 and 3.25 pm.

- gel ...

(mass ...)

0.3361 g

... - ... - ...

... - ... - ...

3.1 (mass ...)

	(g)	... (g)
...	30.50 - 0
...	SA	13.75
...	5.232 - 3.08 u
... ...	2.5 u	2.5 u
... ...	1 u	1 u
...	13.5 u	-

... @ 25°C

1/5/92

- long message of the fractal - House (1-6)

the spin line for me

fractal - simple

Original $S_0 = 2^0$ and $N = 2^0$



1/5/92

total $S_1 = 2^1 = 4$ bp, 17 bp

count - points

total $S_2 = 2^2 = 16$ bp, 51 bp

1/5/92

fractal = $4^0 = 1$ bp, $2^0 = 1$ bp, $1^0 = 1$ bp, $0^0 = 1$ bp

fractal = $4^1 = 4$ bp, $2^1 = 2$ bp, $1^1 = 1$ bp, $0^1 = 1$ bp

	1	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
point	1	4	16	64	256	1024	4096	16384	65536	262144	1048576	4194304	16777216	67108928	268435968	1073743872	4295015680
bp	1	4	16	64	256	1024	4096	16384	65536	262144	1048576	4194304	16777216	67108928	268435968	1073743872	4295015680
total	1	4	16	64	256	1024	4096	16384	65536	262144	1048576	4194304	16777216	67108928	268435968	1073743872	4295015680
S_0	1	4	16	64	256	1024	4096	16384	65536	262144	1048576	4194304	16777216	67108928	268435968	1073743872	4295015680
Area	16.5	11.4	16.5	1.4	15.5	1.4	5.5	1.4	13.5	1.4	15.5	1.4	17	15.5	1.4	17	17

fractal = $4^2 = 16$ bp, $2^2 = 4$ bp, $1^2 = 1$ bp, $0^2 = 1$ bp

and A 5.48 μ

12/16/04

for 45-MHz and 100-MHz

with 100-MHz and 100-MHz

in PC/MAS 1.0 or previous version

- used to convert 100-MHz to 45-MHz -

d)

	Frequency	Frequency	100-MHz	PC/MAS 1.0	PC/MAS 1.0	PC/MAS 1.0
	Hz	Hz	Hz	(Hz)	Spec	Acq
100-MHz	100.0	100.0	100.0	100.0	100.0	100.0
	1.8 Hz	1.8 Hz	1.8 Hz	1.8 Hz	1.8 Hz	1.8 Hz
45-MHz	45.0	45.0	45.0	45.0	45.0	45.0
100-MHz	100.0	100.0	100.0	100.0	100.0	100.0
45-MHz	—	45.0 Hz (1.8 Hz)	45.0 Hz 1.8 Hz	—	45.0 Hz	—
100-MHz	—	100.0 Hz (1.8 Hz)	100.0 Hz 1.8 Hz	—	—	100.0 Hz
100-MHz	100.0	100.0	—	100.0	100.0	100.0

100.0 - 3.4 Hz = 96.6 Hz
 100.0 - 1.8 Hz = 98.2 Hz

1.8 Hz
 100.0 Hz
 (1.8 Hz) (2?)

$$= \frac{100.0 \times 1.8}{1.8} = 100.0$$

→ 51.5 Hz

$$K_{eff} H_{eff} = 6.17 \times 10^{-2} \text{ m} - 0.7 \times \left(\frac{1.4 \times 10^{-2}}{6.17 \times 10^{-2}} \right) = 5.17 \times 10^{-2}$$

6.17 m - 5.17 m = 0.1

$$\text{New } 1.552 \times 10^{-5} \text{ pounds} \\ \text{of water per sq ft} = 0.93 \text{ ft}$$

of water - of water per sq ft - 0.93 in sq ft

Radon Solution Pk

	<u>Radon</u>	<u>Radon</u>
Rad	Radon Radon	Radon
Rad	Radon - 1.552 - 1.4 - 1.4 - 1.4 (100%)	$R_1 = 0.3711$
Rad	5 x Radon - 1.4 - 1.4 - 1.4 - 1.4	$R_2 = 0.1728$
Rad	1 - M Radon	$R_3 = 0.1297$
Rad	Rad - 1.4 - 1.4 - 1.4 (100%)	
Rad	Rad - 1.4 - 1.4 - 1.4 (100%)	
Rad	Rad	
Rad	Rad - 1.4 - 1.4 - 1.4 (100%)	

10/10/00

Calc 55-2021 and solution spot and spot digest.

	(-) Lithium	Br (10.50g)	Ca (15.64g)	Fluor 21	(+) Sul	(+) KNO
10/10	100g (15g)	250g (10.50g)	100g (15.64g)	100g 100	100g 100	100g 100
10/10/2	200	200	200	200	200	200
10/10/3	200	200	200	200	200	200
10/10/4	—	250g	100g	—	100g	—
10/10/5	—	250g	100g	—	100g	—
10/10/6	—	250g	100g	—	—	100g
10/10/7	—	250g	100g	—	—	100g
10/10/8	14.2	—	—	15.64	13.4	13.4

10/10/8 5.51pm @ 37°C

For-202
10/10/8 - 22/19 g

10/12/03

Salt flat - water and frost-free

get groundwater

10/12

Frost-free - dry in sun

water - dry in sun

Windsor
frost-free and water vapor

	Frost-free in sun	Frost-free in sun	B:1 (in sun, ground)
Windsor frost-free	2.5m 1.8m 2.5m	1.8m <u>2.1.47 46</u>	@ 25°C
Windsor water	—	1.8m <u>1.8m</u>	sun @ 1.32pm
Windsor water	2.5m	2.0m	and 1.5pm
Windsor water	1.8m	1.8m	
Windsor	2.5	—	

sun in (in) 1 hour

in 2.5m @ 25

water vapor

ground

water vapor

in
(in sun)

water

2.5m

ground

water vapor

water

2.5m

water

10/12/03

Salt flat - water and frost-free

get groundwater

10/12

Frost-free - dry in sun

water - dry in sun

Windsor
frost-free and water vapor

	Frost-free in sun	Frost-free in sun	B:1 (in sun, ground)
Windsor frost-free	2.5m 1.5m 2.5m	1.5m <u>2.5m</u>	@ 25°C
Windsor water	—	1.5m <u>1.5m</u>	sun @ 1.5m
Windsor water	2.5m	2.5m	and 1.5m
Windsor water	1.5m	1.5m	
Windsor	2.5	—	

sun in (in) 1 hour

in 2.5m of 2.5

water vapor

ground

water vapor

in
(in sun)

water

water

ground

water vapor

water

water

water

10/15/02

Low SS - Avg 2.1 d₅₀

Disturbance (9/30/00) 50:1 #2 - Avg 2.1 d₅₀

Low SS - Avg 2.1 d₅₀

	SI (uvss)	SI (uvss)	SI (uvss)	SI (uvss)	SI (uvss)	SI (uvss)
Point	1000	1000	1000	1000	1000	1000
	200	(3500)	(3500)			
Water	200	1.500	1.500	200	200	
SS	200	1.500	1.500	200	200	
Disturbance	-	1.700	1.700	-	1.1000	(1.1000) 1.100
SS	1300	200	1.500	1500	1300	1300

Strat @ 3:25 pm @ 370
at 7:25 pm

Disturbance - Avg with Avg

	SI (uvss)	SI (uvss)	SI (uvss)	SI (uvss)
Point	1000	1000	1000	1000
		(16,66700)		
Water	200	200	200	200
SS	-	100	-	1000
Disturbance				(1.1000)
SS	1700	1000	1700	1500

12/12/19

Acidimetry of SAP

Wt 55 - divided into 10g

	Wt 55 SAP	Wt 55 Acidim
Wt 55	46.00	4.600
Wt 55	5.1600	5.1600
Phosphate	SAP .500	1.00 1.00

↳ @ 500g

↳ @ 500g

↳ 300g

for 1L

gel order

Wt 55	Phosphate (-)	Phosphate (+)	Wt 55	Wt 55	SAP Wt 55
-	-	-	-	-	-

gel

10/15/04

Longitude of

60° 55' and 110° 10' - 110° 15'

SAP & A-1

	(C) NO. UNIT	SAP 3:1	A-1 3:1	(SAP) 5:1	A-1 5:1
glass 10/13/04	30g 30g	17g 1.7g	17g 1.7g	17g 1.7g	17g 1.7g
insect 9/13/04	—	15g (18g)	15g (18g)	25g 4g 25g	25g 1.5g 25g
Bitter Tulsi	1.5g	1.5g	1.5g	2.5g 1.5g	1.5g
Excess leaves	1g	1g	1g	1g	1g
Alko	9.5	9.9	9.5	9.5	9.5

New total
18g

10/10/20

did revision ✓

for 55 - here - answer

21 16.5m

23 17.5m

- 20000 - here = 20000 - 10000

↓ 10000 at 10000 - 10000

for 55 - here = 10000

	(1)	(2)	(3)	(4)	(5)	(6)
Method	200	300	200	200	200	200
	2.77m	18.667m	2.77m	2.77m	2.77m	2.77m

10000	2.77m	2.77m	2.77m	2.77m	2.77m	2.77m
-------	-------	-------	-------	-------	-------	-------

500	2.77m	2.77m	2.77m	2.77m	2.77m	2.77m
-----	-------	-------	-------	-------	-------	-------

10000	2.77m	2.77m	2.77m	2.77m	2.77m	2.77m
-------	-------	-------	-------	-------	-------	-------

10000	13.2m	13.2m	13.2m	13.2m	13.2m	13.2m
-------	-------	-------	-------	-------	-------	-------

start @ 6.14pm

end 7.14pm



10/15/00

pelour 10/20/00

CRSR list of ASPATI

	INVEST	CRSR EL	INVEST H2	EL P2000 20	CRSR P2000 P2000
plura	6000 20	7000 1500	20000 1500	11000 1000	11000 1000
diff. 114	200	200	200	200	200
WADA					
BSA (1200)	200	200	200	200	200
ASPATI	←	.200	.200	→	1.000 (1100) 1100
ALGO	200	.200	.200	1.500	13.000

K-PAK 1200 -
 3000
 200 20000
 200 200 (1100)
 ↓
 200 20000
 200 20000

10/15/00 to 10/20/00

Start @ 3:30 pm @ 37°C

End @ 11:54 am

status 10/2/2008

ASNA-Swiss digest of PVF

	ASNA-Swiss (-)	ASNA-Swiss SIP #1	ASNA-Swiss SIP #2	PCNA (-)	PCNA SIP #1
Home	Home ✓ SIP	Home SIP	Home SIP	Home SIP	Home SIP
Project 2	SIP	SIP	SIP	SIP	SIP
MU	—	2.4K	2.4K	—	2.4K (1.1K)
WIND	—	—	—	—	2.4K
ASNA	1.7K	2.6 SIP	2.6 SIP	1.7K	5.9 K

ASNA SIP - SIP SIP

ASNA SIP

7.5 SIP SIP SIP

SIP
SIP + SIP SIP

SIP @ 3.1K @ 2.7%

SIP @ 12.23 pm

10/2/04

10/2/04 - routing protocol

diget fire 5000 spk + val 1000

	min 1000 20 (min)	min 1000 1500 1500	min 1000 1500 1500	1000 2000	1000 2000	1000 2000
planned	1000 500	1500 1500	1500 1500	1000 2000	1000 2000	1000 2000
Buffer 2 w/1000	200	2.500	2.500	200	200	200
500 (1000)	200	2.500	2.500	200	200	200
SPK	—	1000 1.500	1000 1.500	—	1000	—
Val	—	1000 1.500	1000 1.500	—	—	1000
1000	1000	1.500	1.500	1.500	1.500	1.500

1000 + 5000 @ 370

10/2/04 5000 @ 11:35pm

10/17/04

- LAST - 11/11/04
with info

Sales Report

WORKSHEET 2 - 11/11/04 - 7,624, 3,119, 1175

WORKSHEET 1 - 5693, 2110, 2004

WORKSHEET 3 - 9100, 2110

WORKSHEET 4 - 8400, 2110

WORKSHEET 5 - 6000, 3,119

at 100 kg type

at 70 kg type

25.1

Blow 2.5 ac

Blow 1.0 ac

Blow 3.7 ac

5.2 ac

25.1

Blow 1.0 ac

Blow 5.7 ac

Blow 19.1

25.1

25.1

Blow 1.0 ac

Blow 12.5 ac

13.5 ac

11/11/04

11/11/04

11/11/04

11/11/04

11/11/04

11/11/04

11/11/04

11/11/04

Handwritten title: $\log_{10} \frac{1}{1 - \cos \theta} = \log_{10} \frac{1}{1 - \cos \theta}$

θ	(1)	(2)	(3)	(4)	(5)	(6)	(7)	(8)	(9)	(10)
0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
15	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
30	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
45	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
60	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
75	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
90	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0

θ	(1)	(2)	(3)	(4)	(5)	(6)	(7)	(8)	(9)	(10)
0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
15	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
30	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
45	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
60	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
75	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
90	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0

Handwritten note: $\log_{10} \frac{1}{1 - \cos \theta} = \log_{10} \frac{1}{1 - \cos \theta}$

	(-)	Exp	(-)	Exp	(-)	Exp	PCDD	PCDD
	6	7	7	7	8	8	8	8
	60g	100g	60g	100g	60g	100g	110g	110g
	2.5	4.2	2.5	4.2	2.5	4.2	inc	inc
Wt	2	2	2	2	2	2	2	2
Wt	2	2	2	2	2	2	2	2
Wt	-	0.1	-	0.1	-	0.1	-	0.1
Wt	13.5	11.2	13.5	11.2	13.5	11.2	15	13.9

(1.9)

Wt	Wt	Wt
1200	200	
Bottle	2400	2400
334 (100)	2700	2400
Wt	1.2	-
dH ₂ O	190.4	102
Wt	1958	173.6

10/10/20 chemistry exper 2ms3

with sph E - xtal 719, 1112

xtal I - xtal 636, 850

sph E - xtal I 529, 297

sph E - xtal

	(-) Exp 1	(-) Exp 2	(-) Exp 2	(-) Exp 2	PCOAN	xtal I	xtal I	xtal I
	1	1	2	2	(-)	xtal	sph E	xtal
plasma	100	100	100	100	100	100	100	100
Buffer #2	2	2	2	2	2	2	2	2
BS4	2	2	2	2	2	2	2	2
sph E	→	→	→	→	→	→	→	→
xtal	→	→	→	→	→	→	→	→
allho	15	15	15	15	15	15	15	15

1 = 2 hours
2 = 4 hours
stand @ 3:41pm

xtal I - xtal

sph E - xtal I

	(-) Exp 1	(-) Exp 2	(-) Exp 2	(-) Exp 2		(-) Exp 1	(-) Exp 2	(-) Exp 2	(-) Exp 2
	1	1	2	2		1	1	2	2
plasma	100	100	100	100	plasma	100	100	100	100
Buffer #2	2	2	2	2	Buffer #2	2	2	2	2
BS4	2	2	2	2	BS4	2	2	2	2
xtal	→	→	→	→	xtal	→	→	→	→
sph E	→	→	→	→	sph E	→	→	→	→
allho	15	15	15	15	allho	15	15	15	15

10/20/09

- seeds furess-aligned with MapA2E

DATA-shed align with PVAII

furess-herz align with spot of shed

because they were lost in the gel isolation etc

previous time ~~was~~ - error - the generation tubing was

broken did not have ethanol

Why some parents of 10/15/09

get info

furess aligned with MapA2E

#1 .1997 g

#2 .1348

#3 .3155 g

DATA-shed align with PVAII

#1 .2160

#2 .9902

#3 .1587

#4 .2798