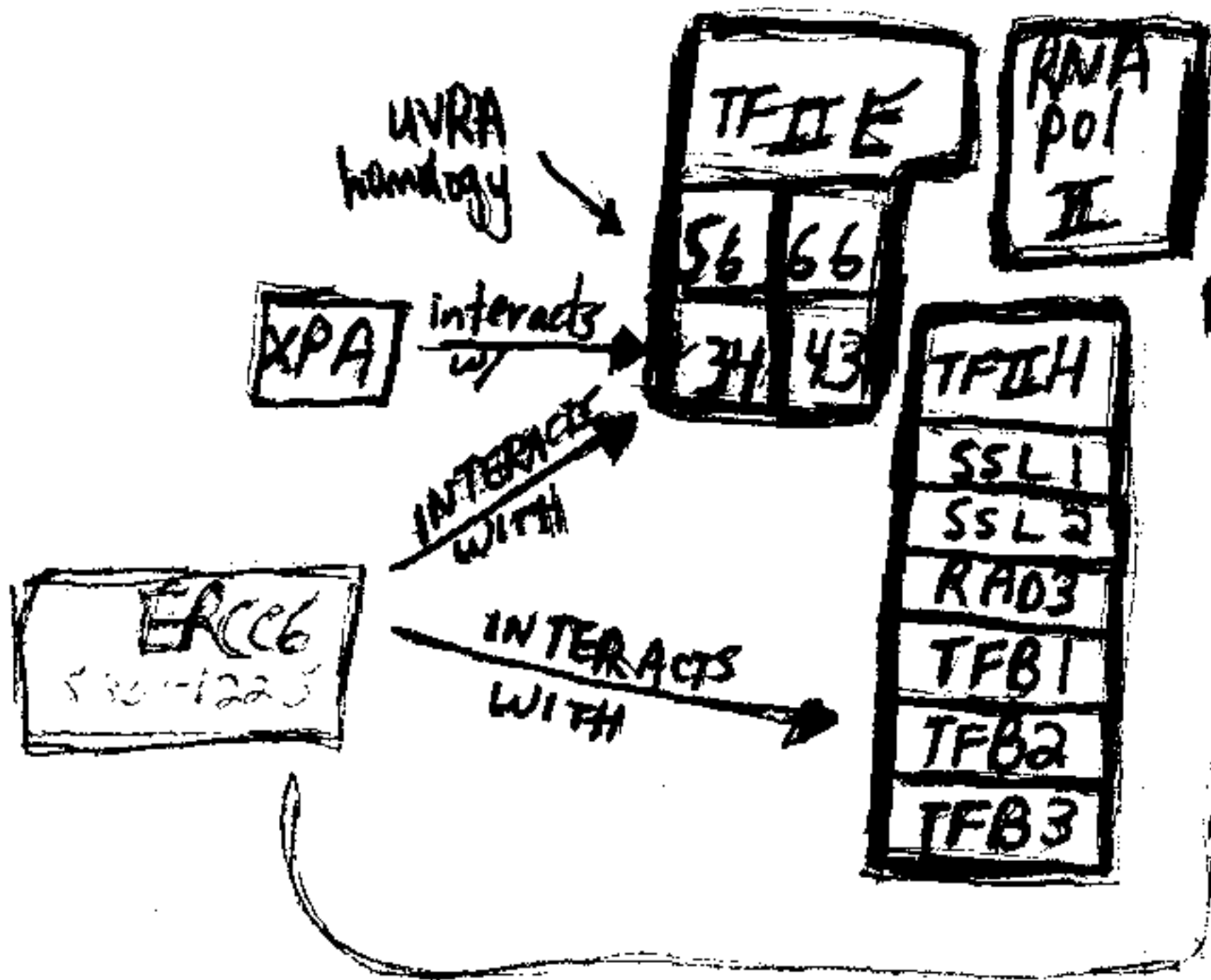
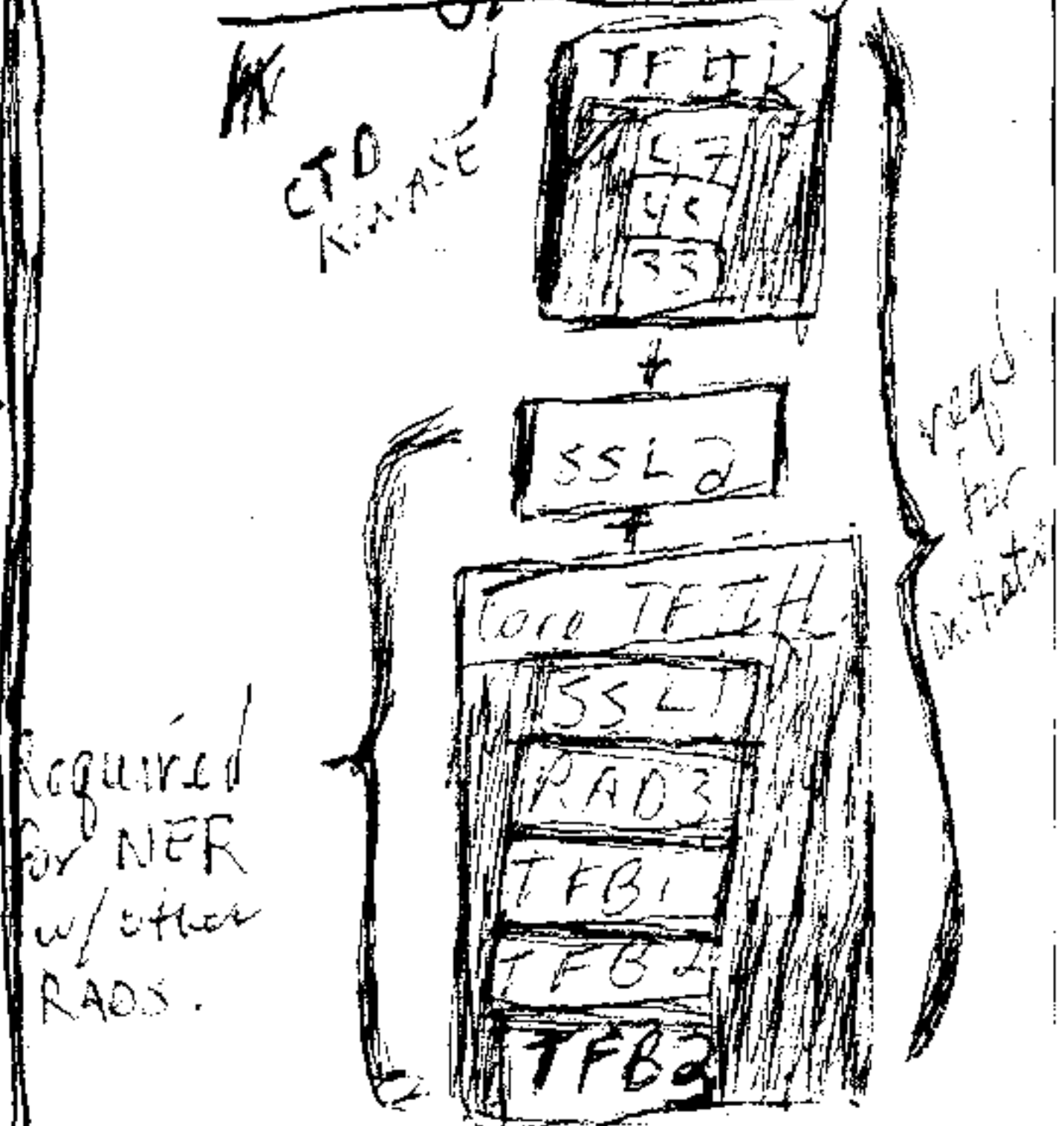


Phil - meeting report

Reinberg/Sancar



Kornberg/Friedberg



SSL2/XPB - active helicase needed for tx

RAD3/XPD - silent helicase needed for NER

3/15/95

Sumit wu

TFIIK

SSL0

RA03

p47

SSL2

TFB1

p45

SSL1

p33

p38

p55

TX

repair

TFIIK

NER PROTS

(RA01, 2, 4, 10, 14)

+

+

CORE-SSL2

~~Is SSL2 required for TX?~~

- no CTD kinase activity w/o TFIIK

- Is SSL2 required?

- ran TFIIH over column w/ SSL2 AB.

- reconst. assay - no tx w/o SSL2

Repair assay

2/10/90

Lab Meeting

~~Base~~

Meeting Updates



- Gordon conferences - Mammalian DNA Repair  
M. Smerdon - organizer

- A. Wolffe - chromatin structure

- acetylation of histones opens up regions  
for tx & replication

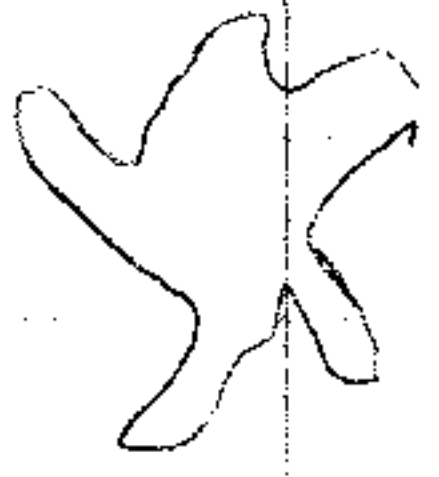
M. Bradbury

CPDs ... only ~50% accessible  
... some enhanced access

PUVA - Psoralen + UVA

- monoadduct is much more mutagenic  
per toxicity than crosslinks

MATS



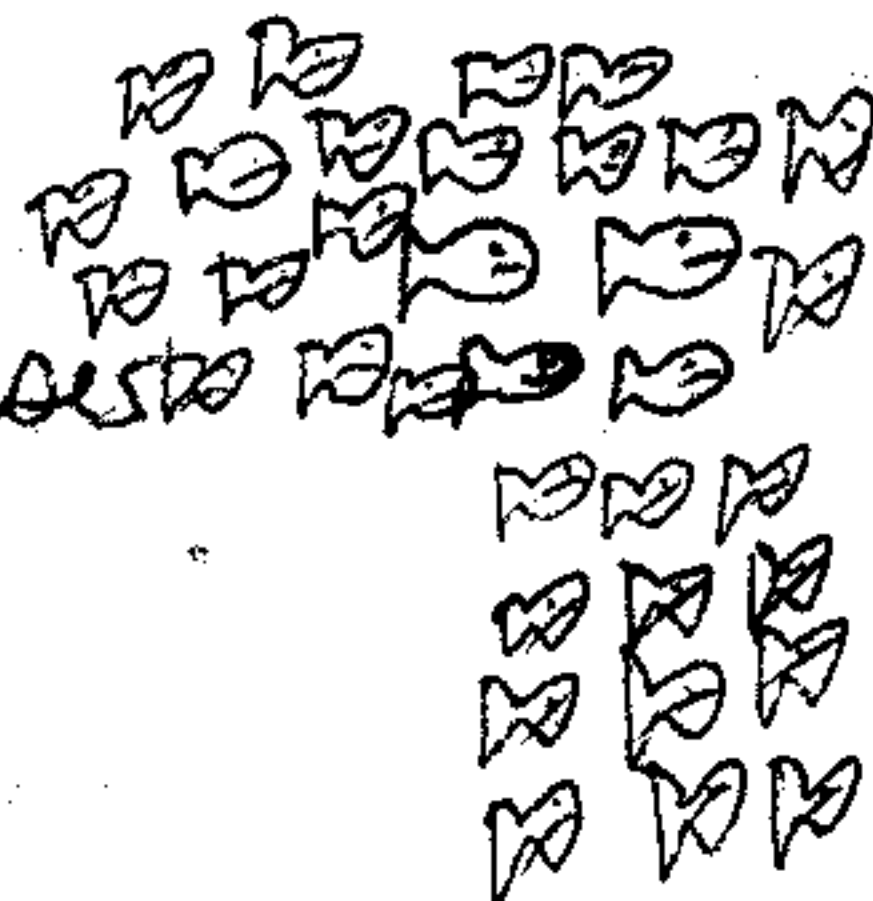


John Essigman

HUBF --- bind cis-plat adducts & binds cruciform  
--- tx. factor hijacking

M. Melend

BPDE -- run of G's binding increases

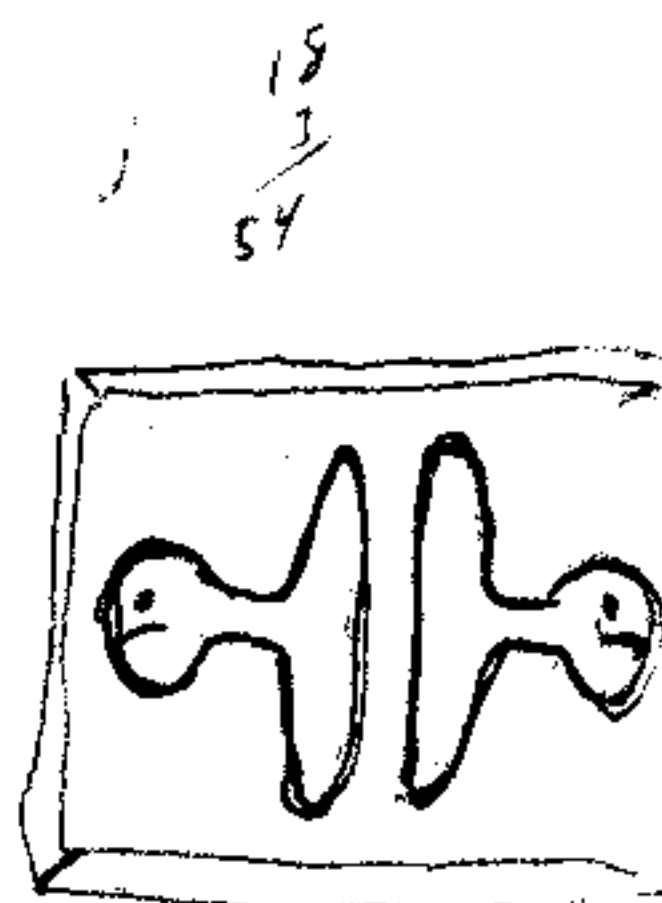
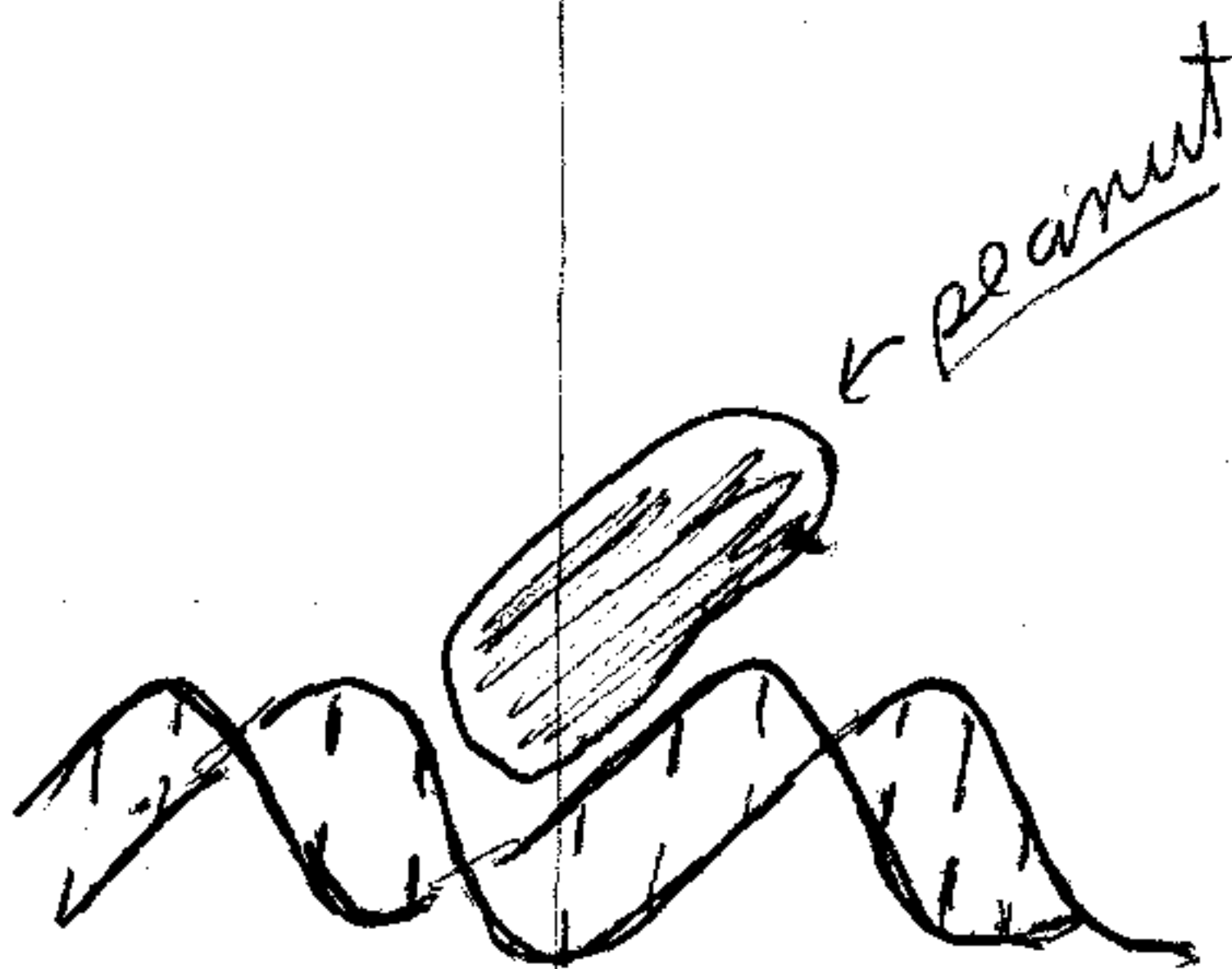


L. Fritz

- mammalian rRNA gene repair
- separate molecules from active **rDNA**
- no diff. in repair of diff. fractions

MOUSE

- bleomycin repair in human
- v. high in DHFR & rRNA



# TCR

~~PCR~~ / Melton

① In LoVo (MSH2<sup>-</sup>)

- defective in tcr in DHFR

② MLH1<sup>-</sup>

- same thing

③ MLH1 w/ chrom. complementation  
-TCR<sup>+</sup>

Pfeifer - fine structure

- many sites of hotspots are @ CpG

- 245 } methylated  
248 }

- some methylated sites not hotspots

- all methylated CpGs which are also DNase I resistant they are slow spots for repair

- methylation reduces G-Us

J. Hoeijmakers

ix problems

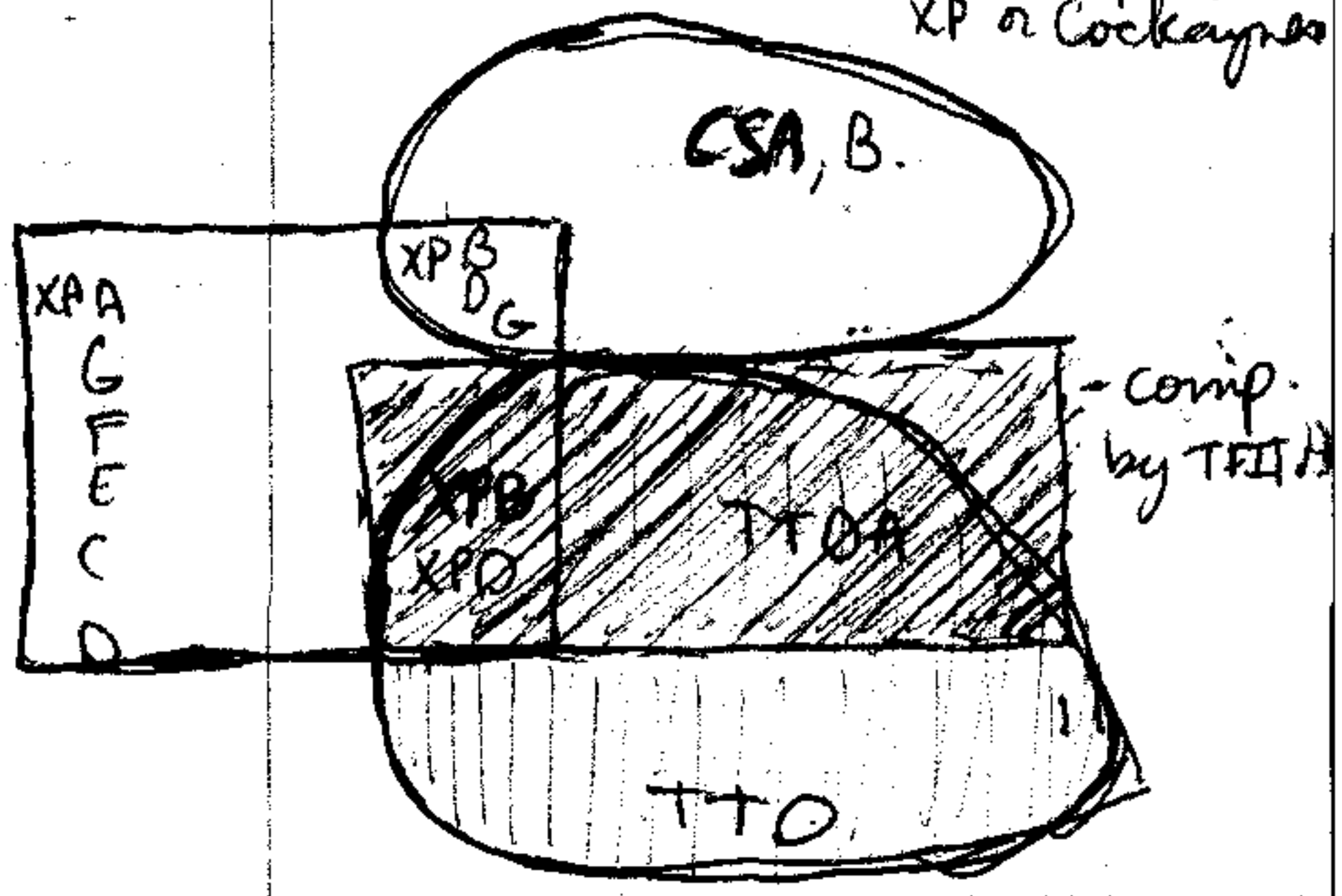
- PIB105
- haywire
- silver mouse

~~13. 13. 13. 13. 13.~~

TFIIH - components

-TTDA

- defective in NER
- complemented by TFIIH injection
- no overlap w/ XP or Cockayne



- ERCC1
  - XPA
  - XPC
  - XPG
  - XPF
  - CSB
- } DO NOT INTERACT w/ TFIIH

S. Wilson

- pol B ... mutants mice
- defective in BER
- sensitive to MMS

(double mutant) cell line

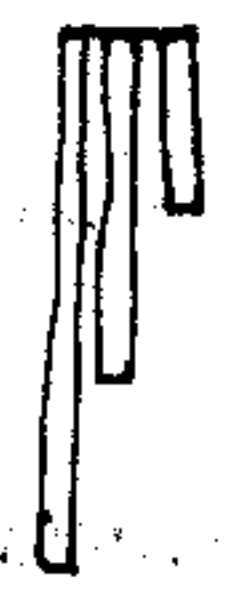
(double mutant) mice dead

Ayasui - N. Grassa

- ~~enzymes~~ enzymes cut at G-Uis, CPDs

- MUS18

- like S. pombe (Bowman...)

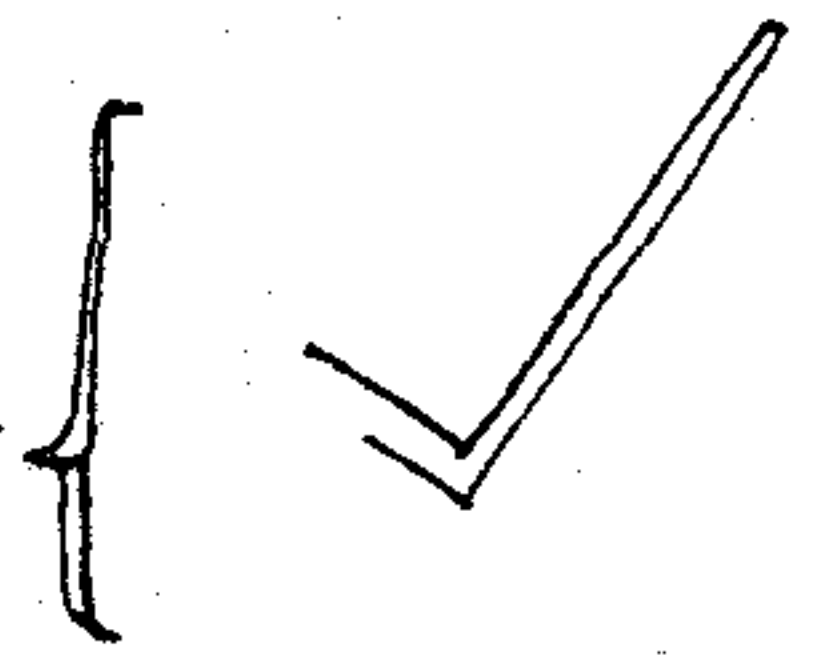
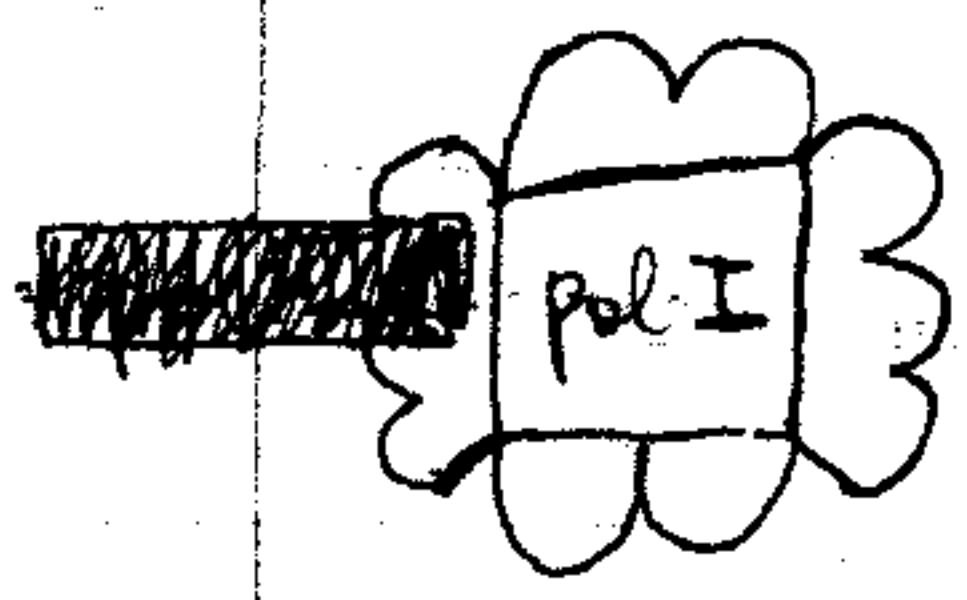


D. Bogenbragen

-> Xenopus extracts

-> 5s RNA gene

-> tx. blocks



P53 mutants

- survival lower



B. Stillman

- HDV

- PCNA not DNA binding ... just donut

- RFC ... opens up PCNA donut

Hoeymakers - mouse mutants

ERCC1 KO

- smaller
- can delete 100 nt from N-term
- can delete
- cannot delete C-term

- ERCC3 mutant

- homozygotes die as embryos

- ERCC6

- no TCR



- mutants normal phenotype but ~~not~~ not NER

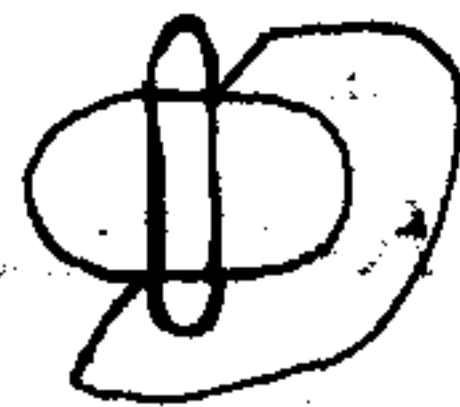
BAD - Cold Spring Harbor  
Vogelstein

HNPKC tumors - from 90% of the patients - microsat. instability

- HMSHZ  
- HPM51  
- HPM52

4 Causes of microsat. instability

- ① defect in mismatch repair
- ② repair/replication
- ③ clonal variation
- ④ art. facts



suggests some may be due to polo

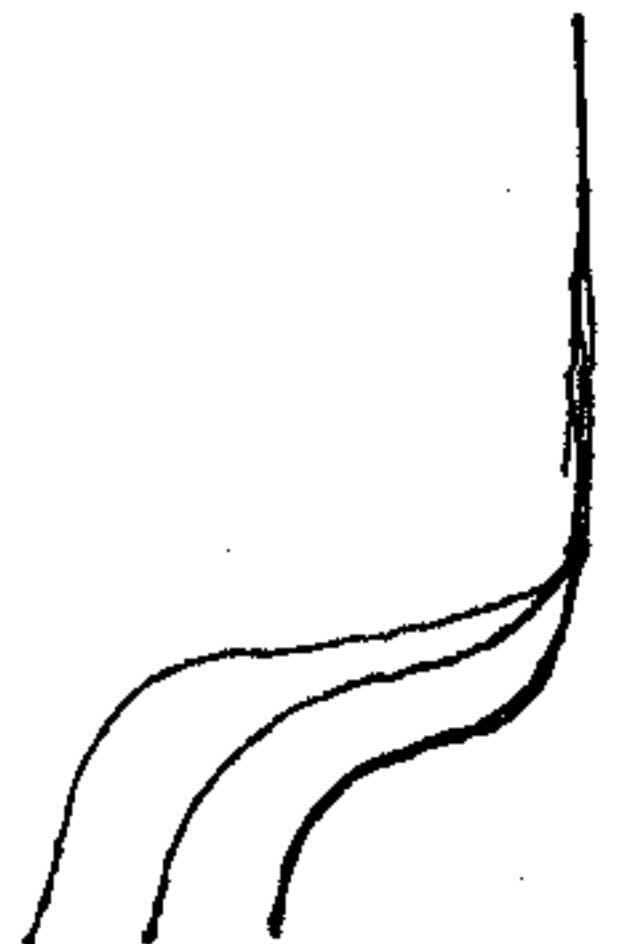
T. Petes

- 33 bp GT expands
  - 51 bp GT contracts
- } seems to think DNA slippage is more likely

one AT base pair in the middle stabilizes tract

Methyltransferase

- $mgt^-$  = sensitive to O-6-meG
- $mgt^- msh^-$  = resistant to O-6-meG





3/10/95

# Lab Meeting

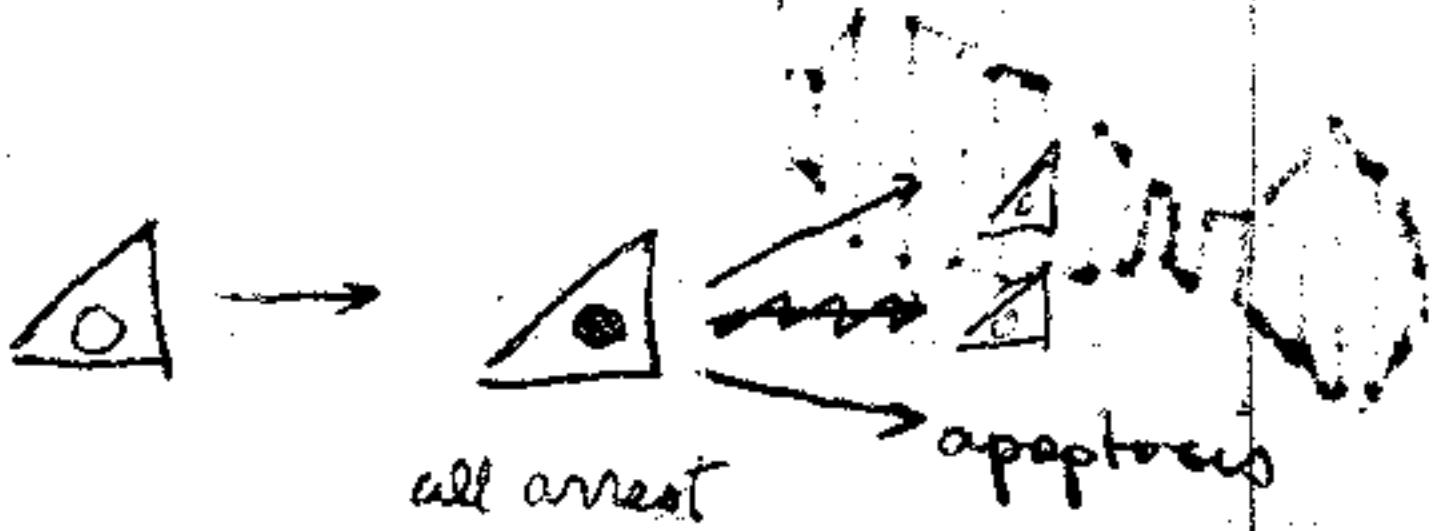


JIM FJORD

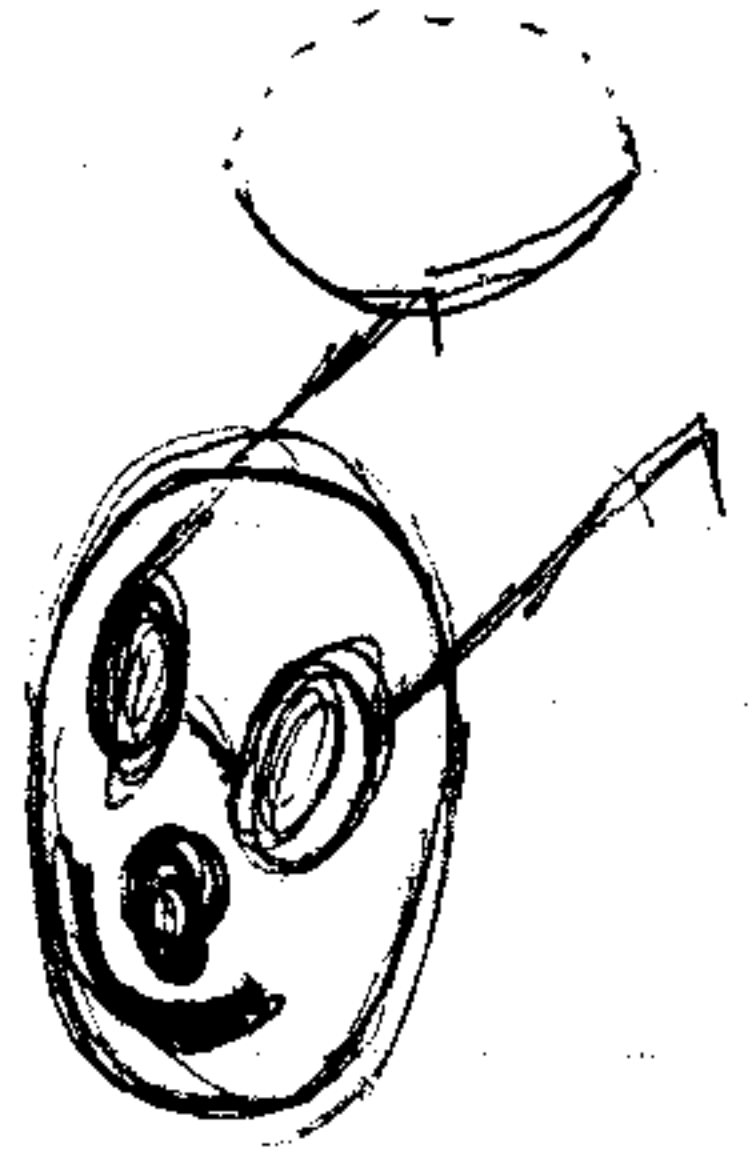
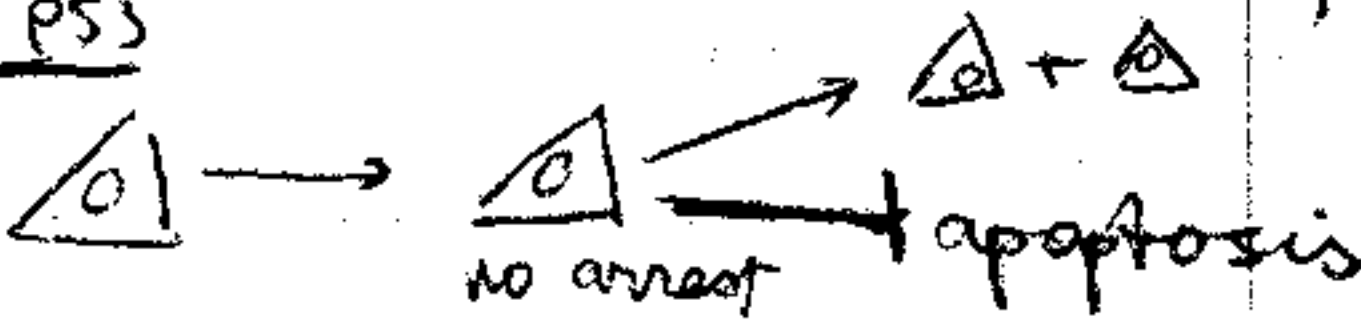


## REPAIR & UV SENSITIVITY IN P53 MUTANTS

WT P53



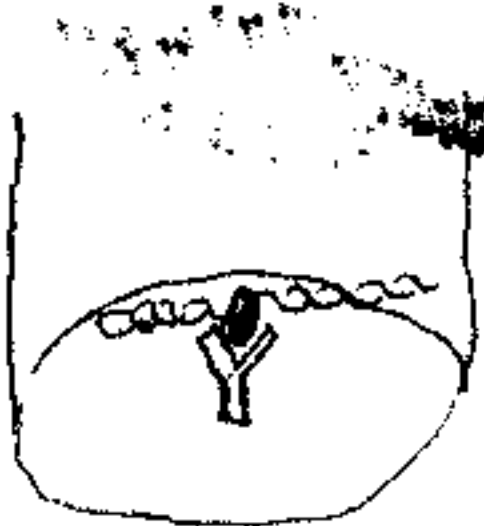
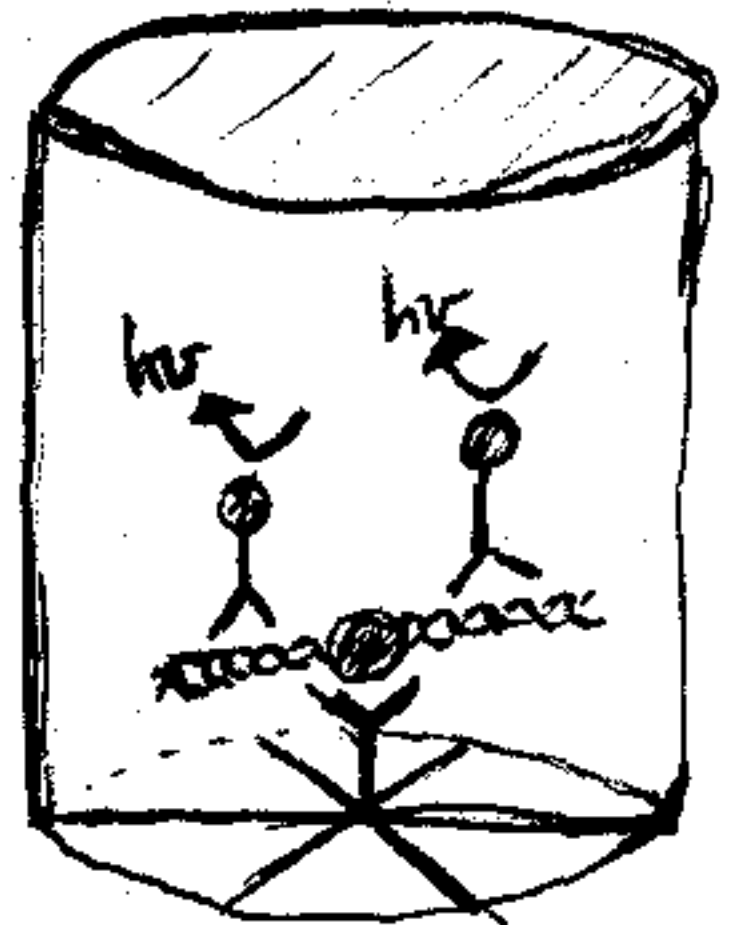
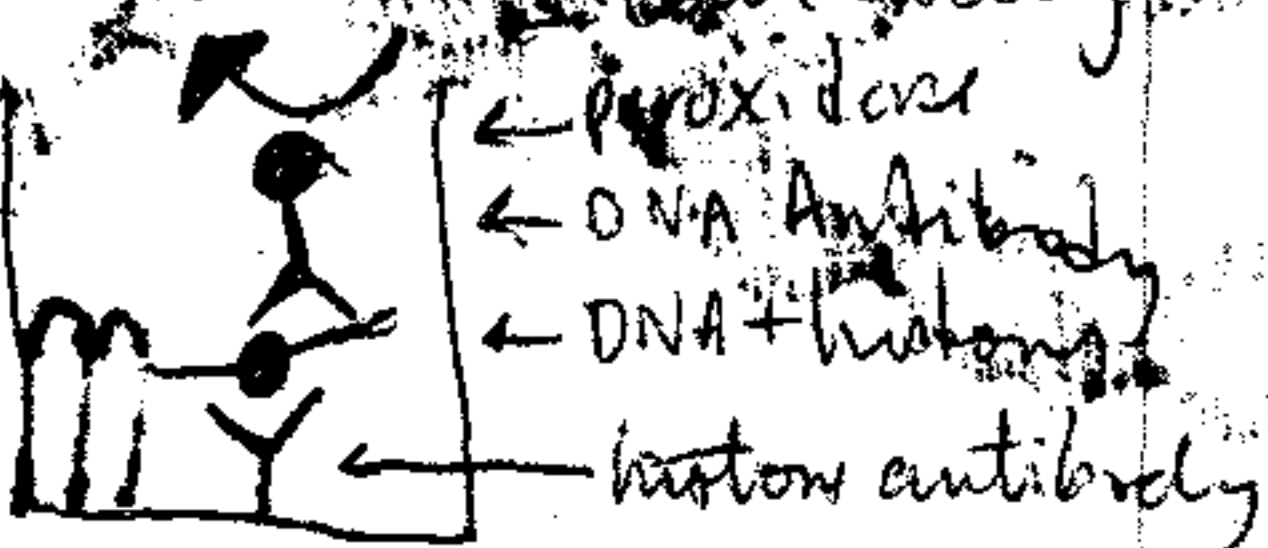
mutant P53



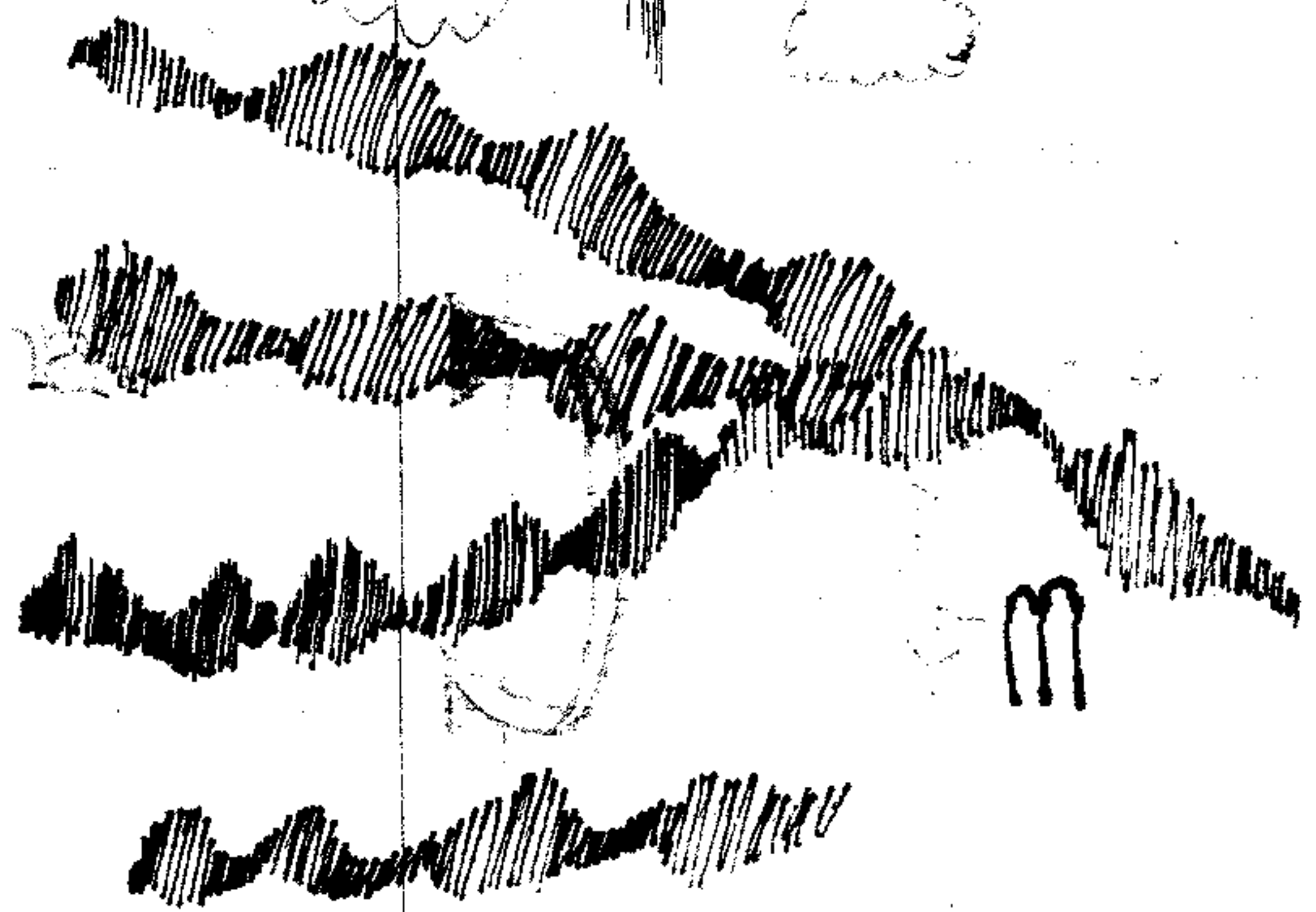
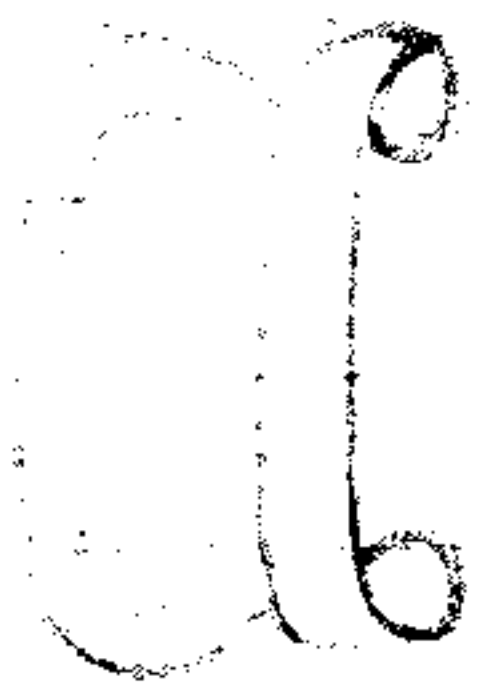
## LI-FRAUMENI SYNDROME

- mutants in P53 more resistant to UV
- mutants (-/-) lower repair of CPD's

## ELISA ASSAY



~~~~~



J

LONG

m