

# M2D2: Western Blot & System Conditions

03/10/16

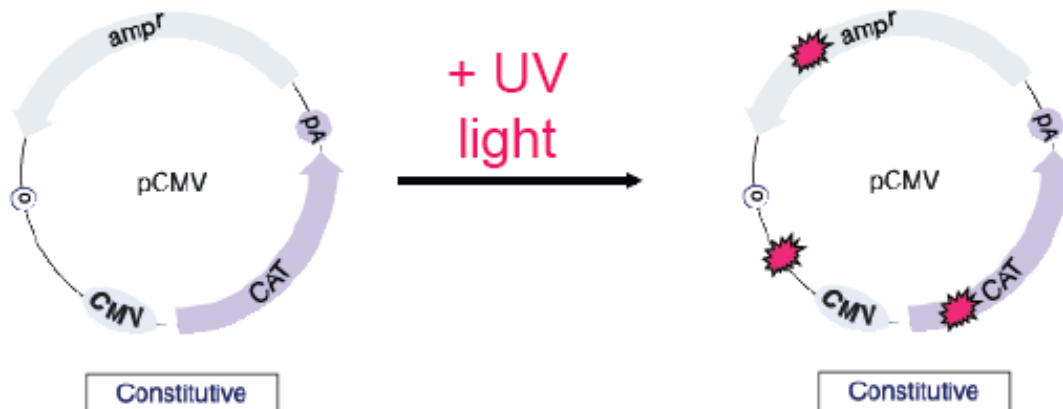
1. Pre-lab discussion
2. Lyse cells
3. Measure total protein concentration
4. Western Blot Analysis: SDS-PAGE & Transfer to nitrocellulose
5. Investigate DNA repair sensor — pick your damage conditions (add to discussion page)

# MOD2 Major Assignments

- Journal Club Presentation (10%)  
in class March 17<sup>th</sup> or April 7<sup>th</sup>
- System Engineering Research Article (25%)  
due at 5pm on Monday, April 18<sup>th</sup>
- M2D3 Homework:
  - 1) Schematic diagram (Figure, title and caption) of the NHEJ reporter plasmid with all features *relevant* to the NHEJ assay labeled.
  - 2) Diagnostic digest calculation

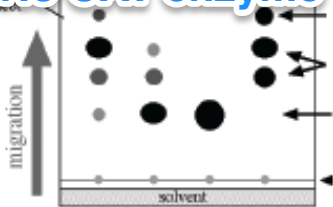
# Reactivation of UV damaged DNA by Host cell Reactivation (HCR)

Athas & GROSSMAN  
Cancer Res. 1991



Transient  
transfection  
peripheral  
blood  
lymphocytes

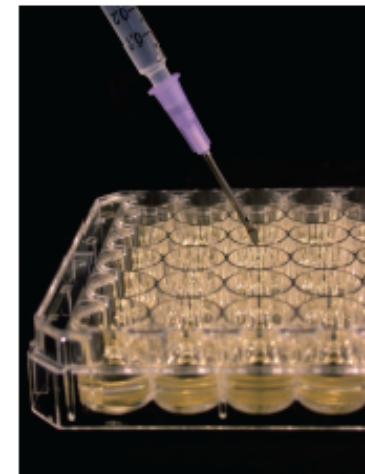
repair=CAT enzyme functional  
no repair= No CAT enzyme



CAT Assay

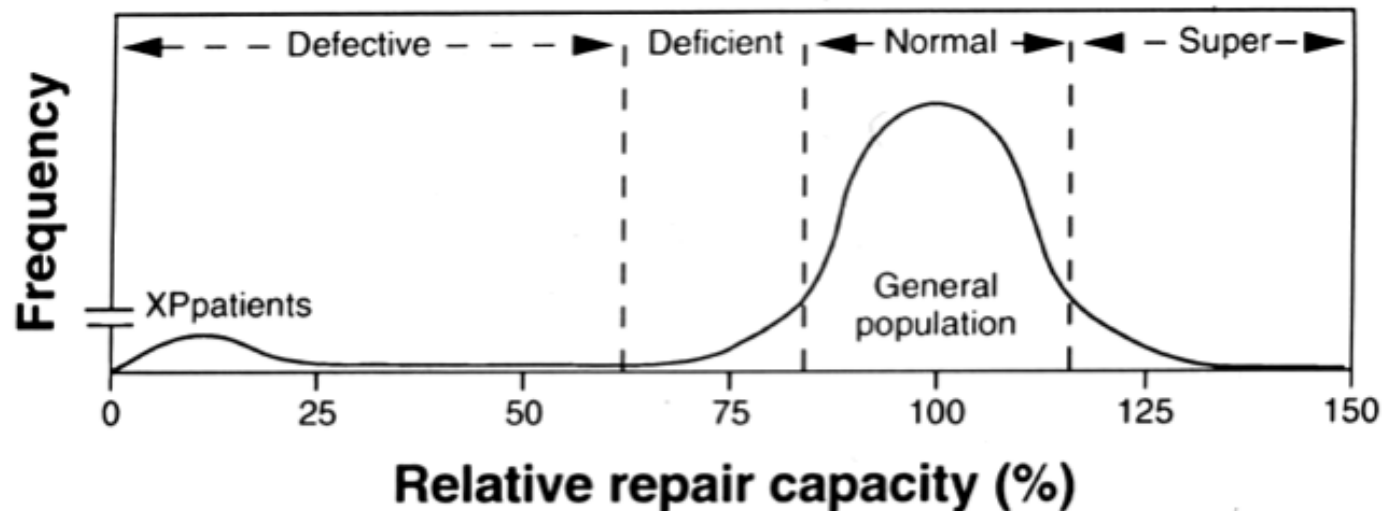


Time to repair



assay found variability in human population in regards to DNA repair

## Interindividual Variation in DNA Repair Capacity

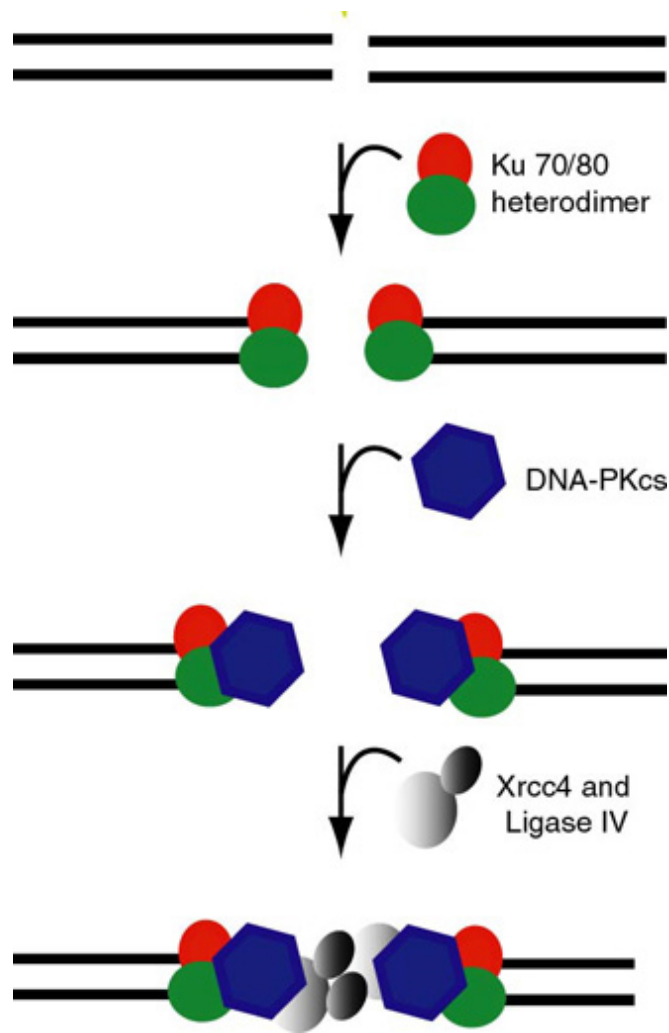


Adapted from **GROSSMAN and Wei (1995)** Clinical Chem 41: 1854-1863

XP frequency = ~1:250,000 giving a theoretical maximum of  
~28,000 cases worldwide with 2,000-fold increased risk

Even if just 1% of the population is relatively repair deficient,  
could have tens of millions with several-fold increased risk

# Non-Homologous End Joining (NHEJ)



Ku70

Ku80

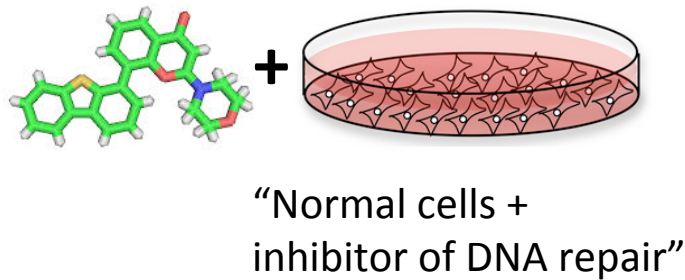
DNA-PKcs

Xrcc4

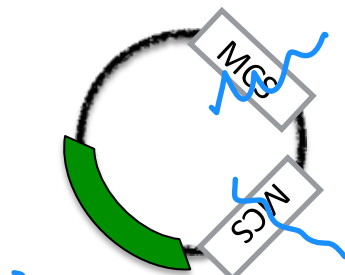
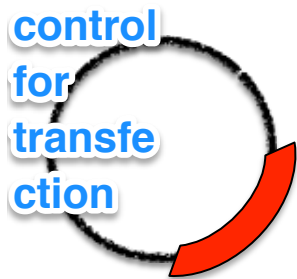
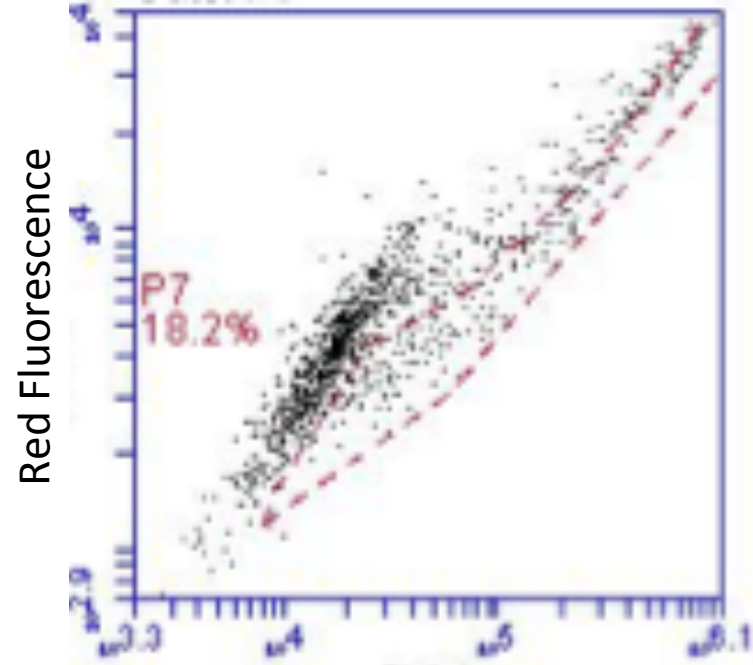
Ligase IV

# Module 2 Experimental Goal:

How efficiently does DNA repair via NHEJ act on DNA damage with different topologies?



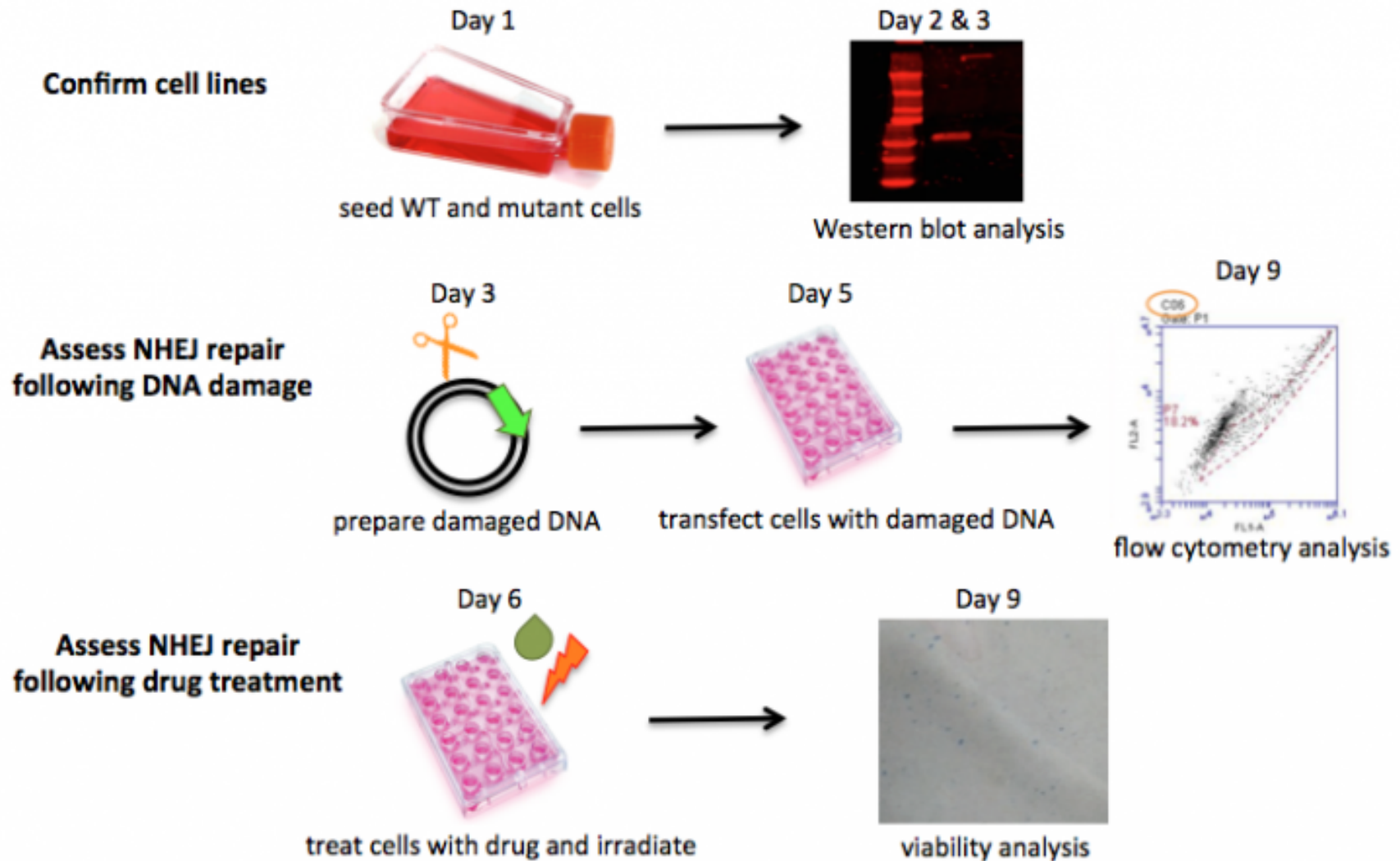
## Repair by NHEJ



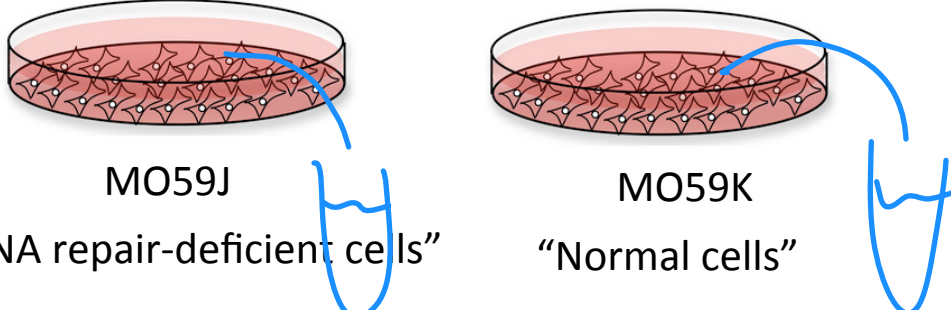
RE cut plasmid to control the cut type  
repair=green  
no repair=no green

mCherry

# Mod 2 experimental overview



# Validation of the experimental system:



MO59J  
"DNA repair-deficient cells"

MO59K  
"Normal cells"

**break all membranes**

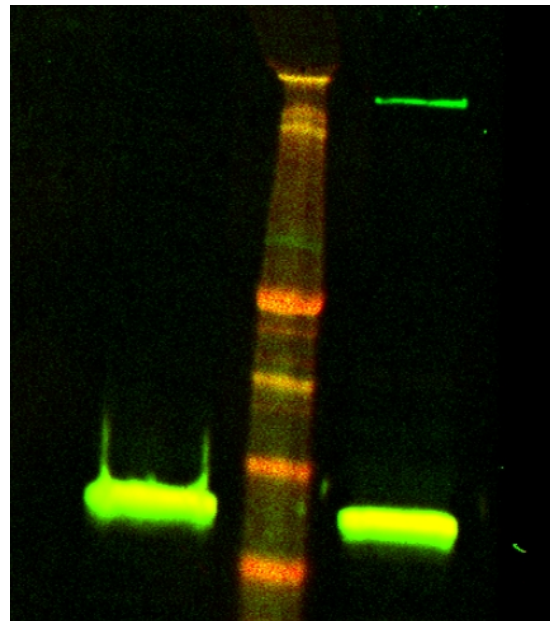
**Mammalian Lysis Buffer, RIPA:**

- 1% NP40 ; 0.1% SDS;  
0.5% sodium deoxycholate  
**strong detergents**
- protease inhibitors  
**stop protein degradation**
- Tris-HCl pH7.4: NaCl  
**physiological pH and salt concentration**

Cell lysate protein concentration measured using  
Precision Red Protein Assay

**LI-COR Western blot  
imaged in far red**

MO59J      MO59K

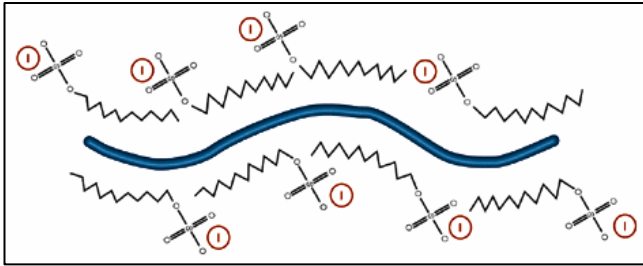


-460kDa  
-117kDa  
-55kDa

Western blot probed  
with: **anti-DNApk**  
**anti-tubulin**

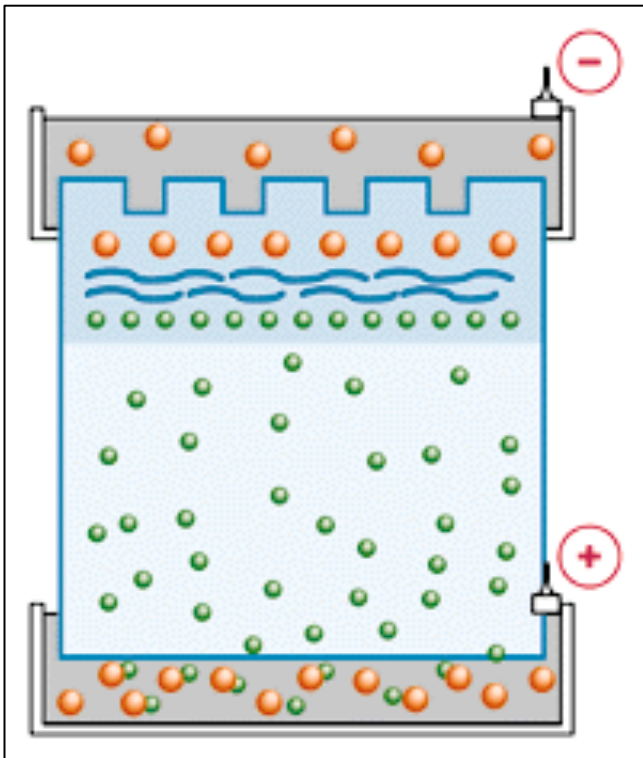


# Western Blot Analysis (Step 1): SDS-PAGE



- Laemmli sample buffer / loading dye:

**SDS, BME, bromophenol blue**



- boiling denatures higher-order structures

- TGS buffer

+ Tris-HCl



+ SDS/protein

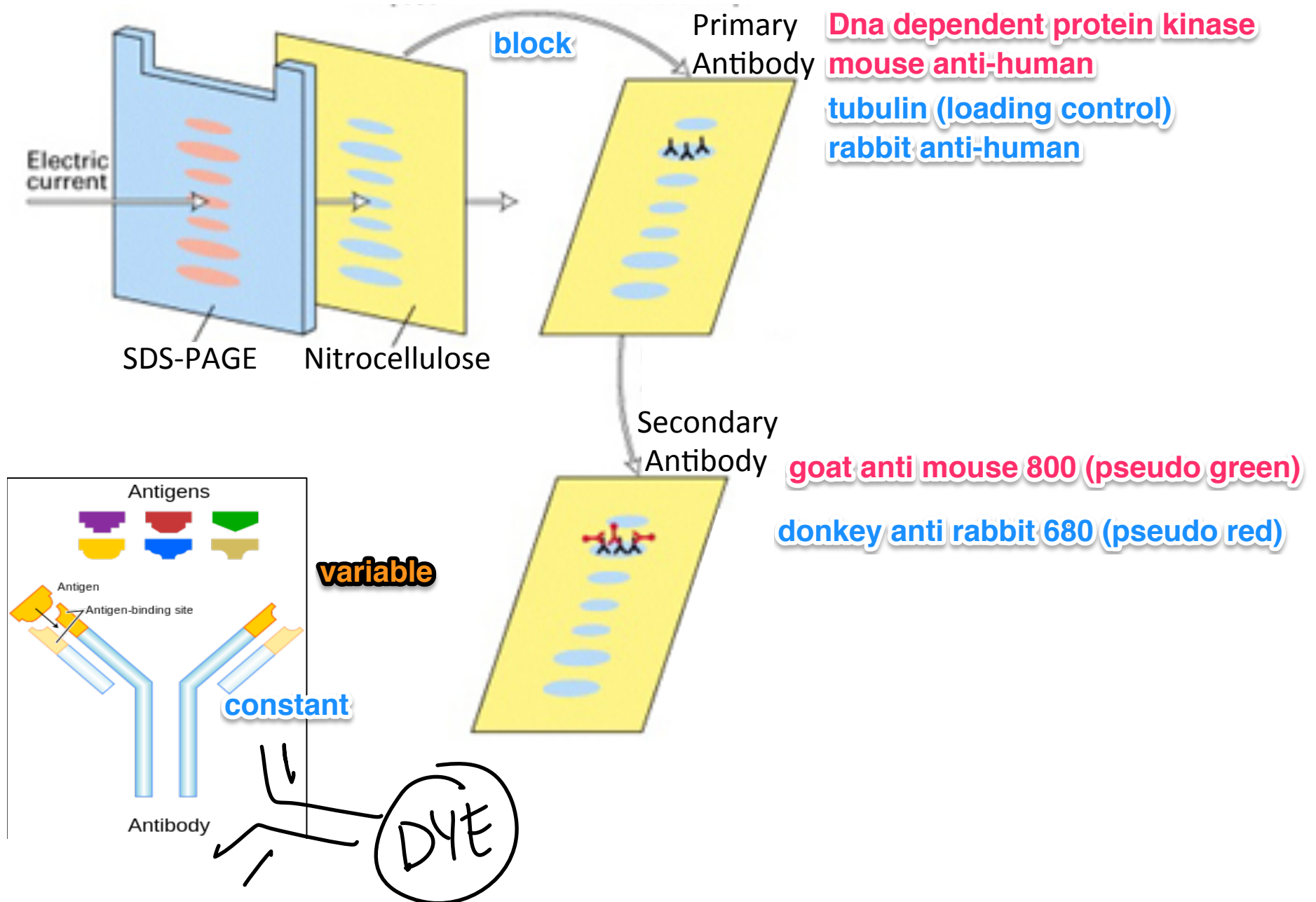


+ glycine



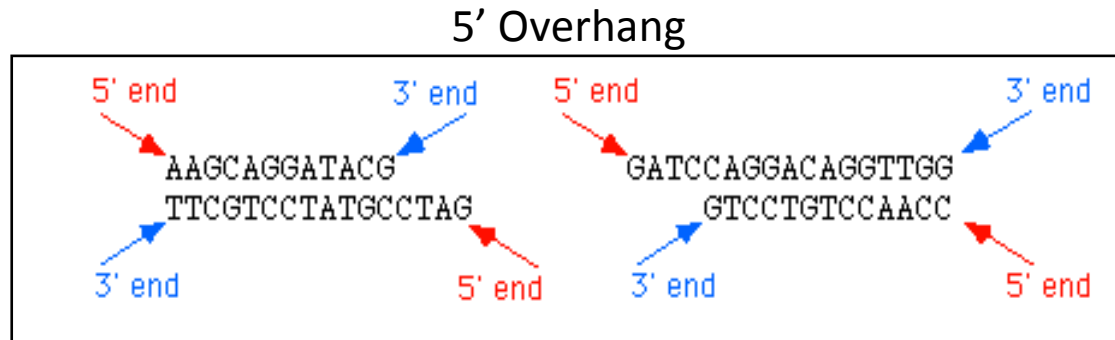
**Hi Mark: pertained large MW proteins**

# Western blot Analysis (Step 2): Transfer and Immunoblotting

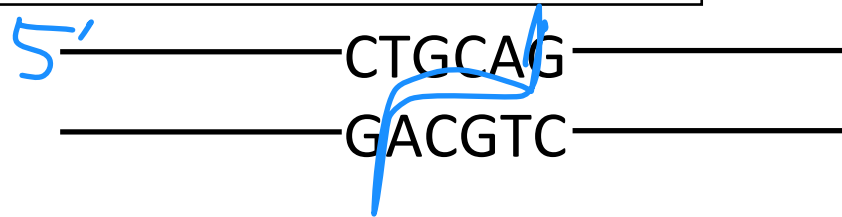


# Restriction Enzymes digestion = DNA damage

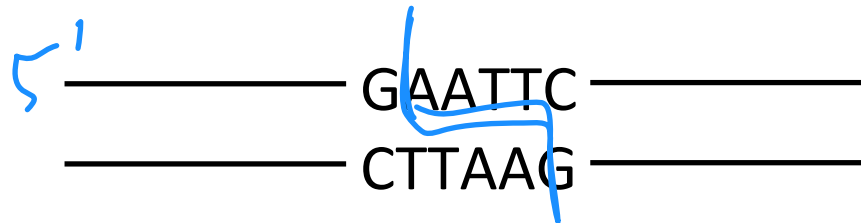
5



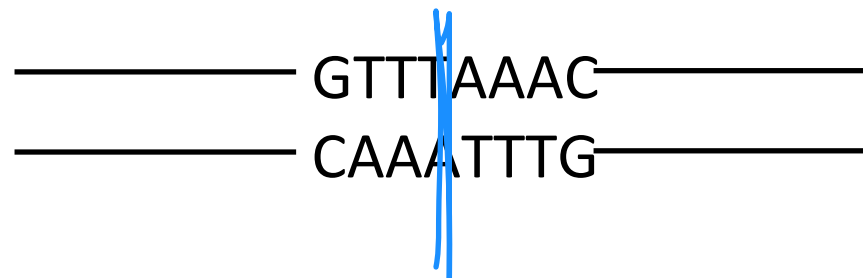
3' overhang  
ex. PstI



5' overhang  
ex. EcoRI



Blunt  
ex. PmeI



# M2 Model of DNA damage:

## Potential damage types:

blunt ends



compatible overhangs



incompatible overhangs



different color -  
diff sequence

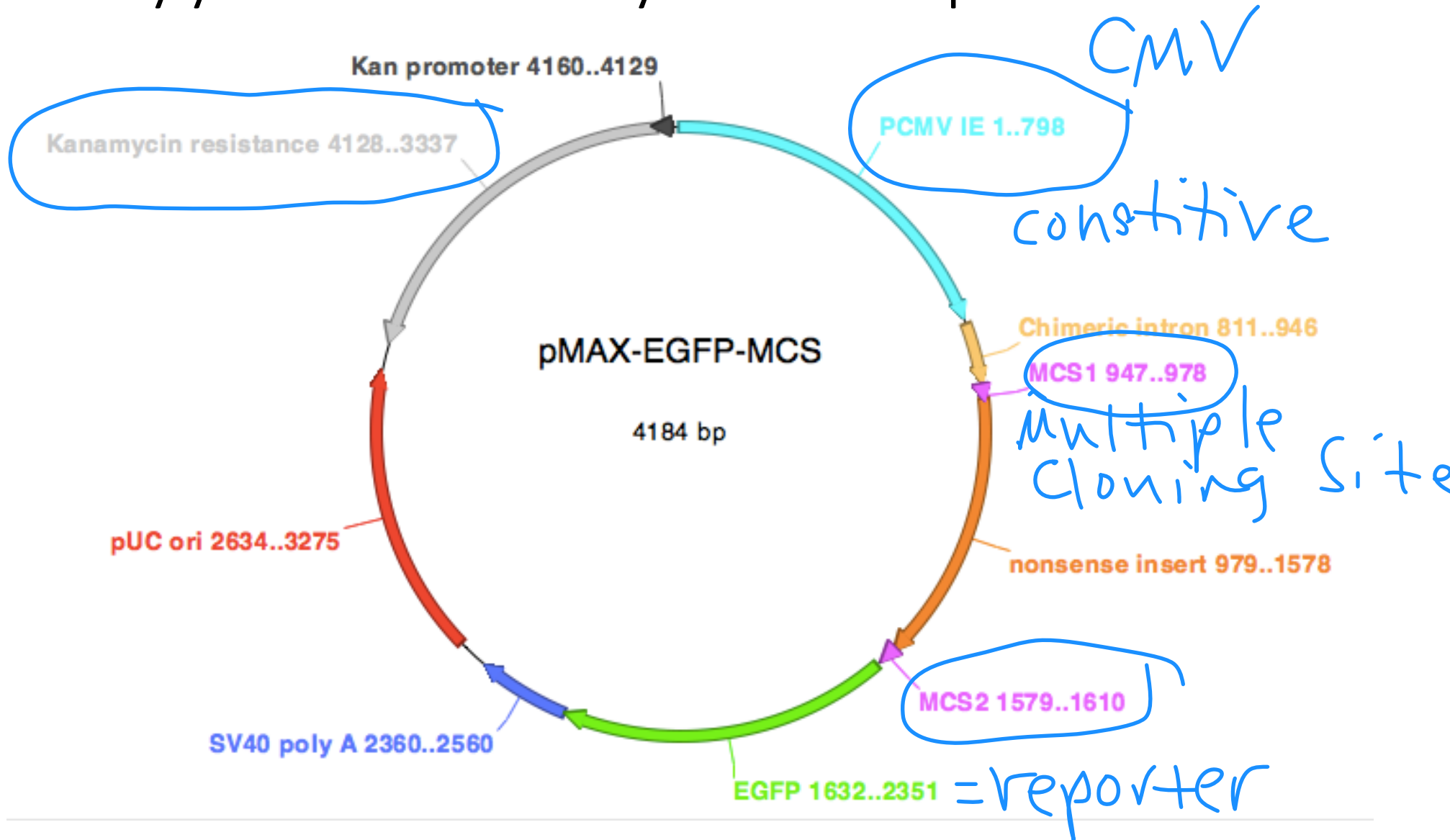
## Hypothesis for NHEJ Repair capacity:

① easiest

② easier

③ hardest

# Today you will familiarize yourself with pMAX-EGFP-MCS



## Today in lab:

- Lyse cells on ice, keep lysate cold!
- Measure total protein concentration with Precision Red
- Load samples on SDS-PAGE
- Transfer protein to nitrocellulose membrane
- Familiarize yourself with the NHEJ reporter
- Add you DNA damage choice to wiki discussion page