

M2D2: Western Blot & System Conditions

03/10/16

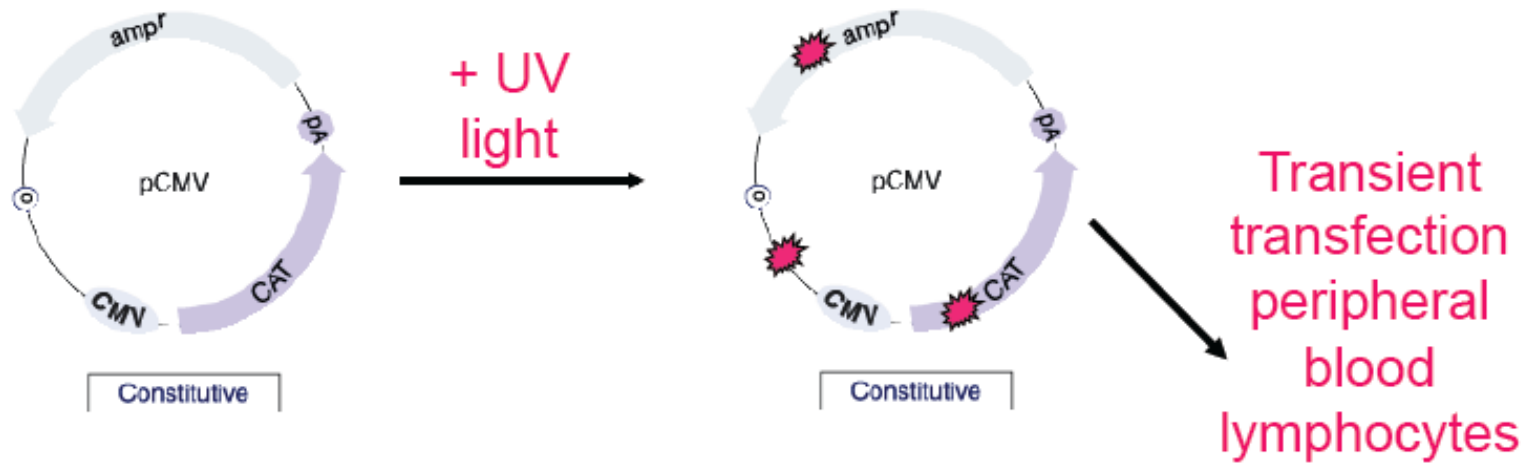
1. Pre-lab discussion
2. Lyse cells
3. Measure total protein concentration
4. Western Blot Analysis: SDS-PAGE & Transfer to nitrocellulose
5. Investigate DNA repair sensor — pick your damage conditions (add to discussion page)

MOD2 Major Assignments

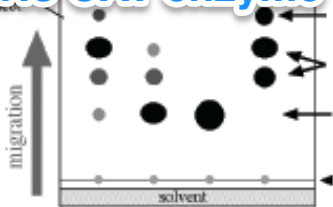
- Journal Club Presentation (10%)
in class March 17th or April 7th
- System Engineering Research Article (25%)
due at 5pm on Monday, April 18th
- M2D3 Homework:
 - 1) Schematic diagram (Figure, title and caption) of the NHEJ reporter plasmid with all features *relevant* to the NHEJ assay labeled.
 -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

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

Athas & GROSSMAN
Cancer Res. 1991



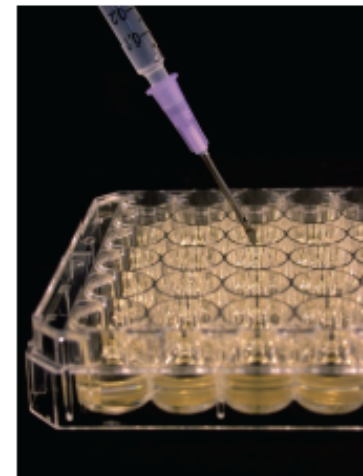
repair=CAT enzyme functional
no repair= No CAT enzyme



CAT Assay

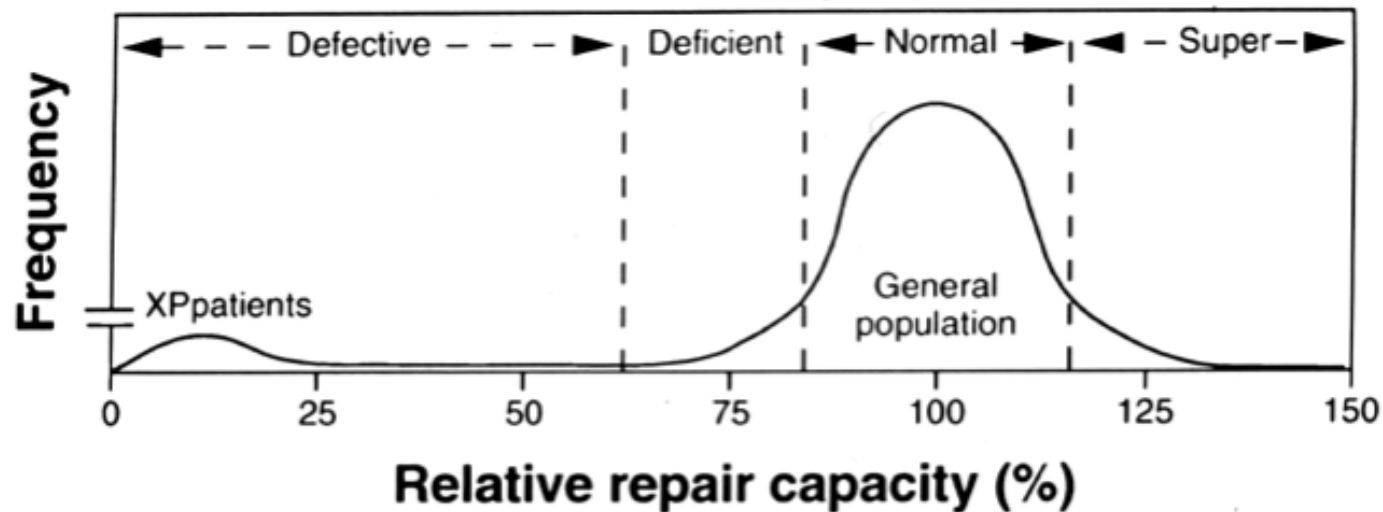


Time to repair



assay found variability in human population in regards to DNA repair

Interindividual Variation in DNA Repair Capacity

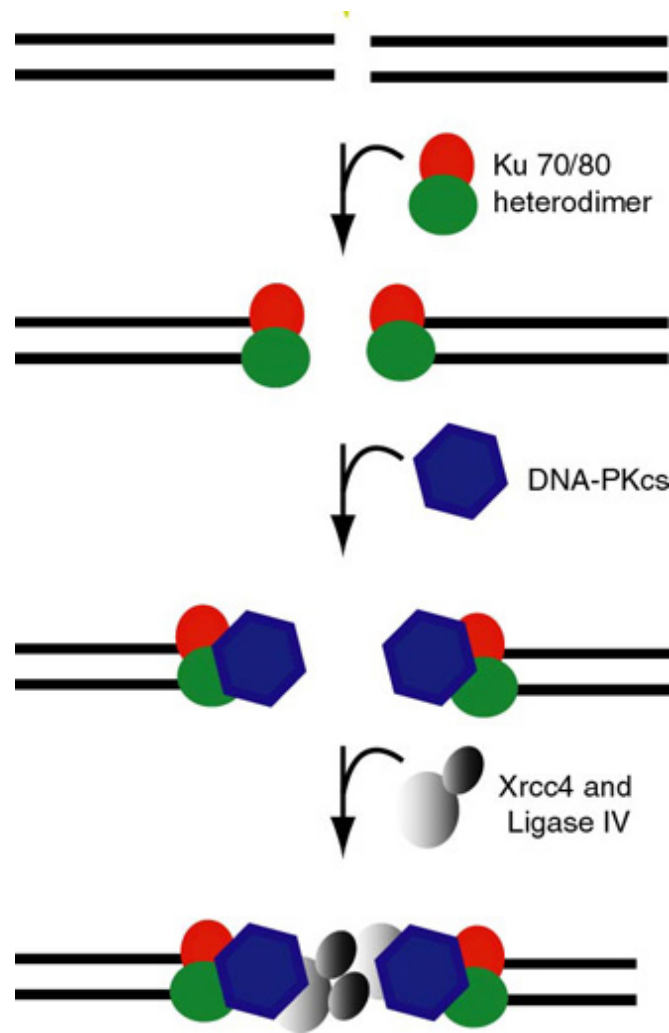


Adapted from **GROSSMAN and Wei (1995)** Clinical Chem 41: 1854-1863

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

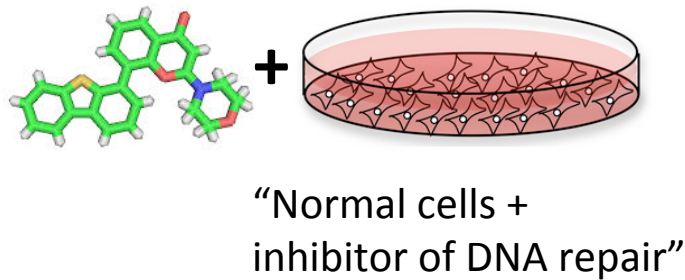
DNA-PKcs

Xrcc4

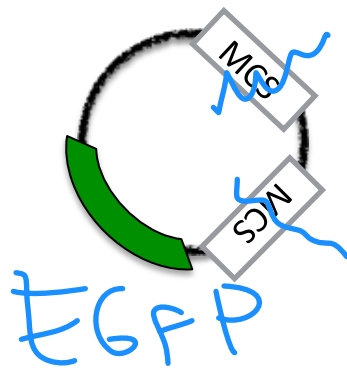
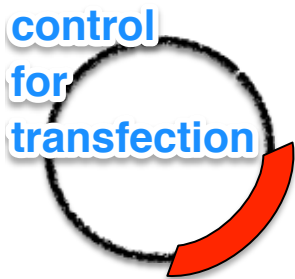
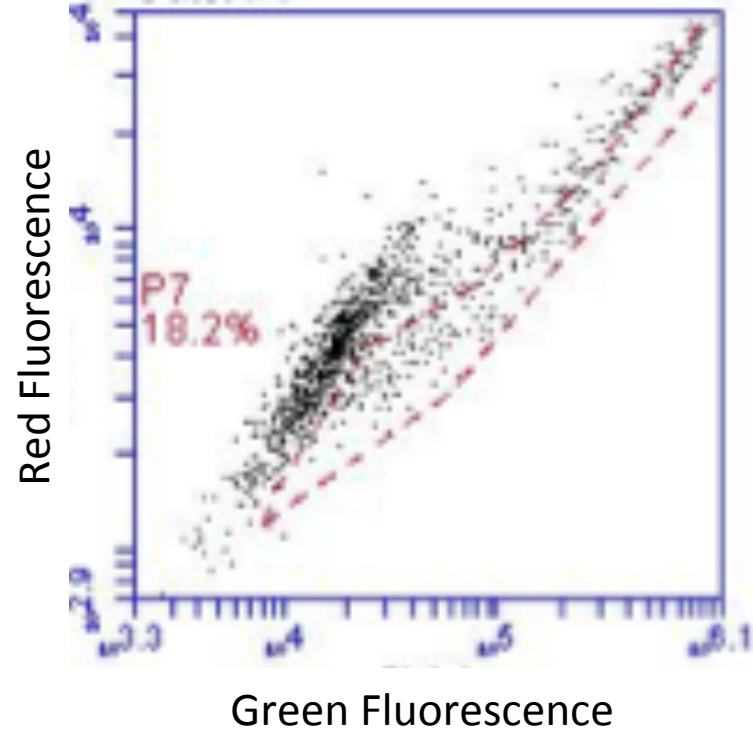
Ligase IV

Module 2 Experimental Goal:

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



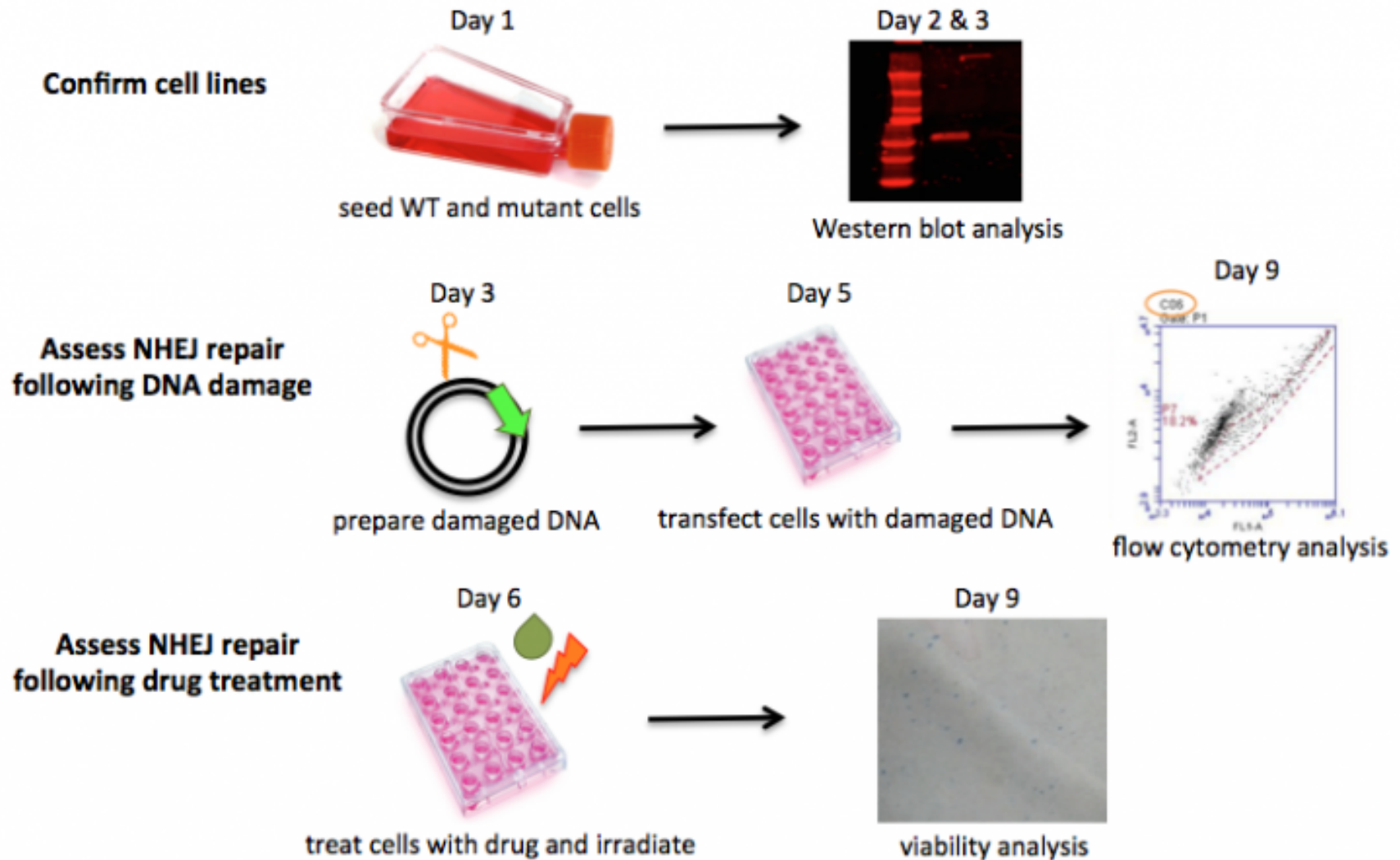
Repair by NHEJ



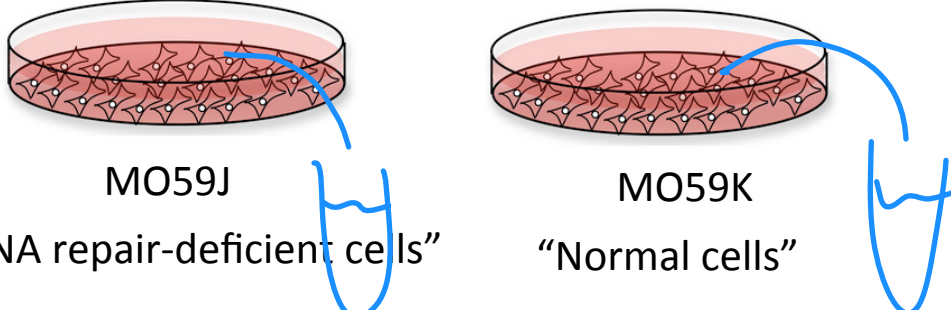
RE cut plasmid: to control the cut type
repair=green
no repair=no green

mCherry

Mod 2 experimental overview



Validation of the experimental system:



MO59J
"DNA repair-deficient cells"

MO59K
"Normal cells"

breaks all membranes

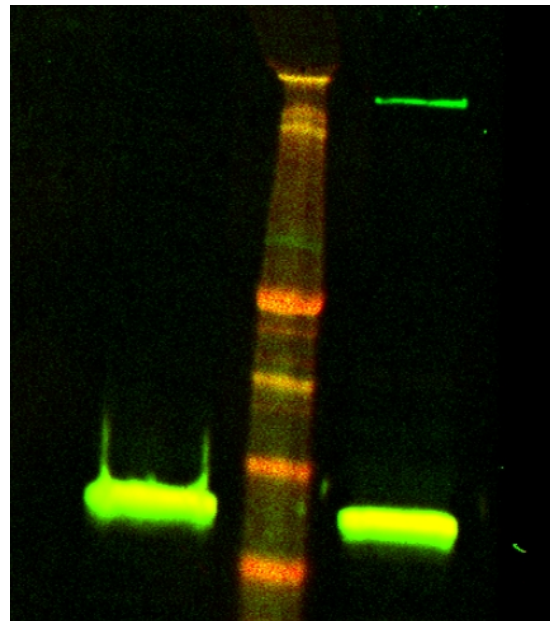
Mammalian Lysis Buffer, RIPA:

- 1% NP40 ; 0.1% SDS;
0.5% sodium deoxycholate
strong detergents
- protease inhibitors
stop protein degradation
- Tris-HCl pH 7.4: NaCl
physiological pH and salt concentration

Cell lysate protein concentration measured using Precision Red Protein Assay

LI-COR Western blot
secondary dye imaged in far red

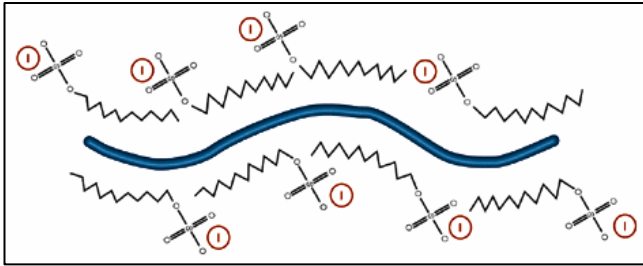
MO59J MO59K



-460kDa
-117kDa
-55kDa

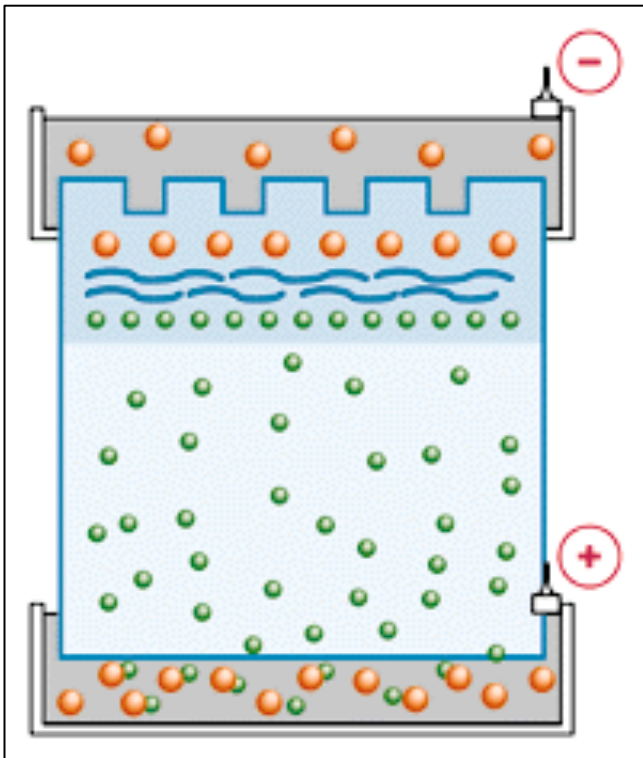
Western blot probed with:
anti-DNApk
anti-tubulin

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

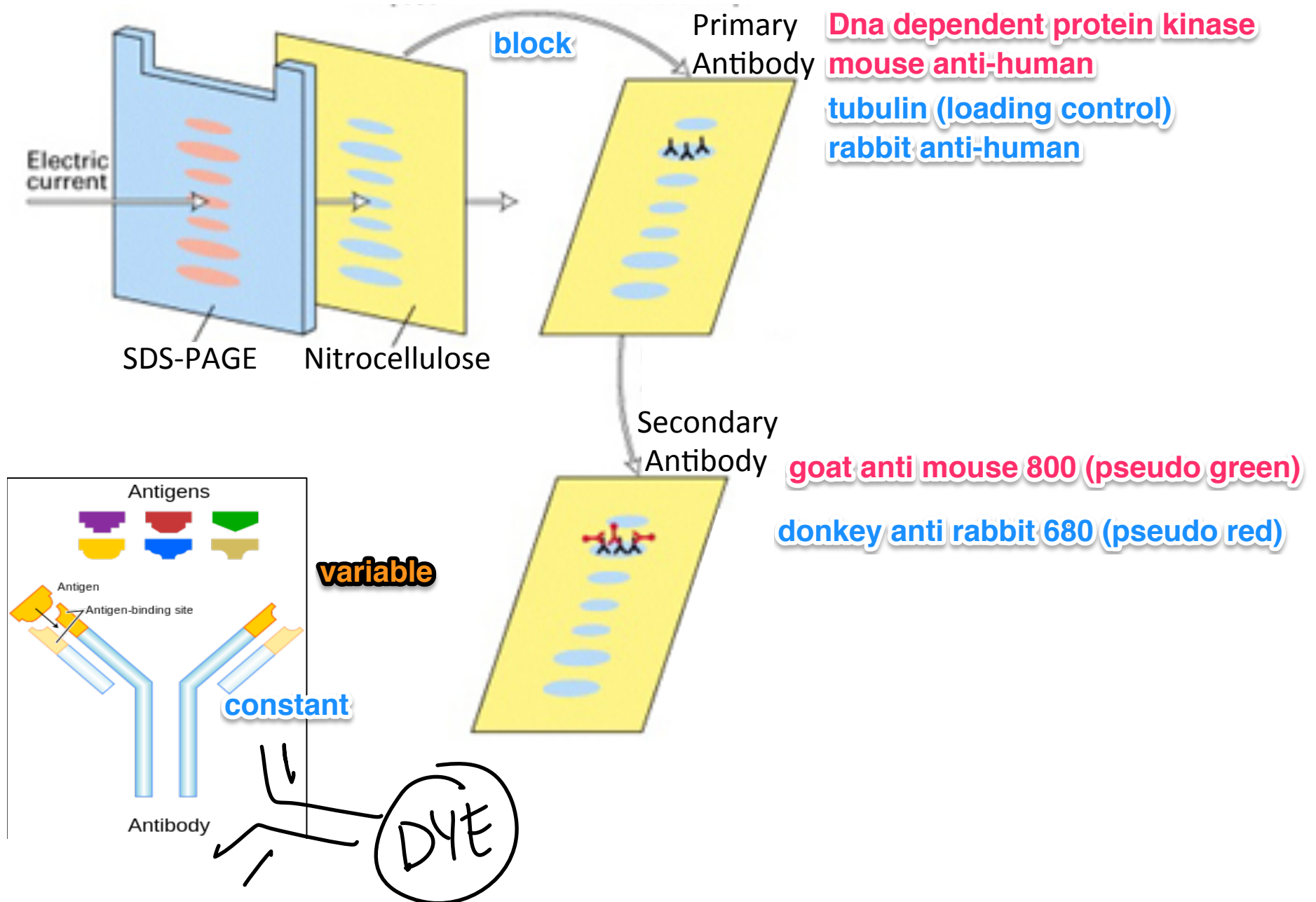


+ glycine

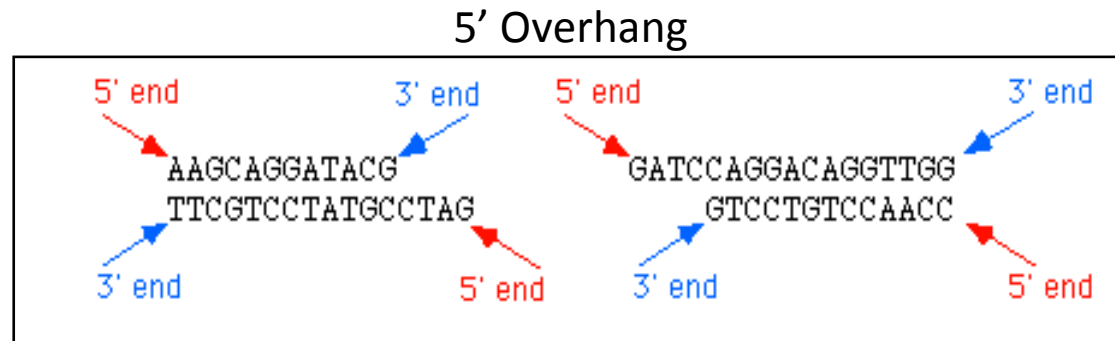


Hi Mark: prestained ladder for large MW proteins

Western blot Analysis (Step 2): Transfer and Immunoblotting



Restriction Enzymes digestion = DNA damage



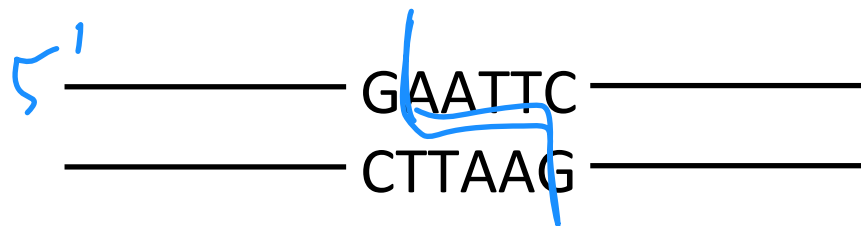
3' overhang

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

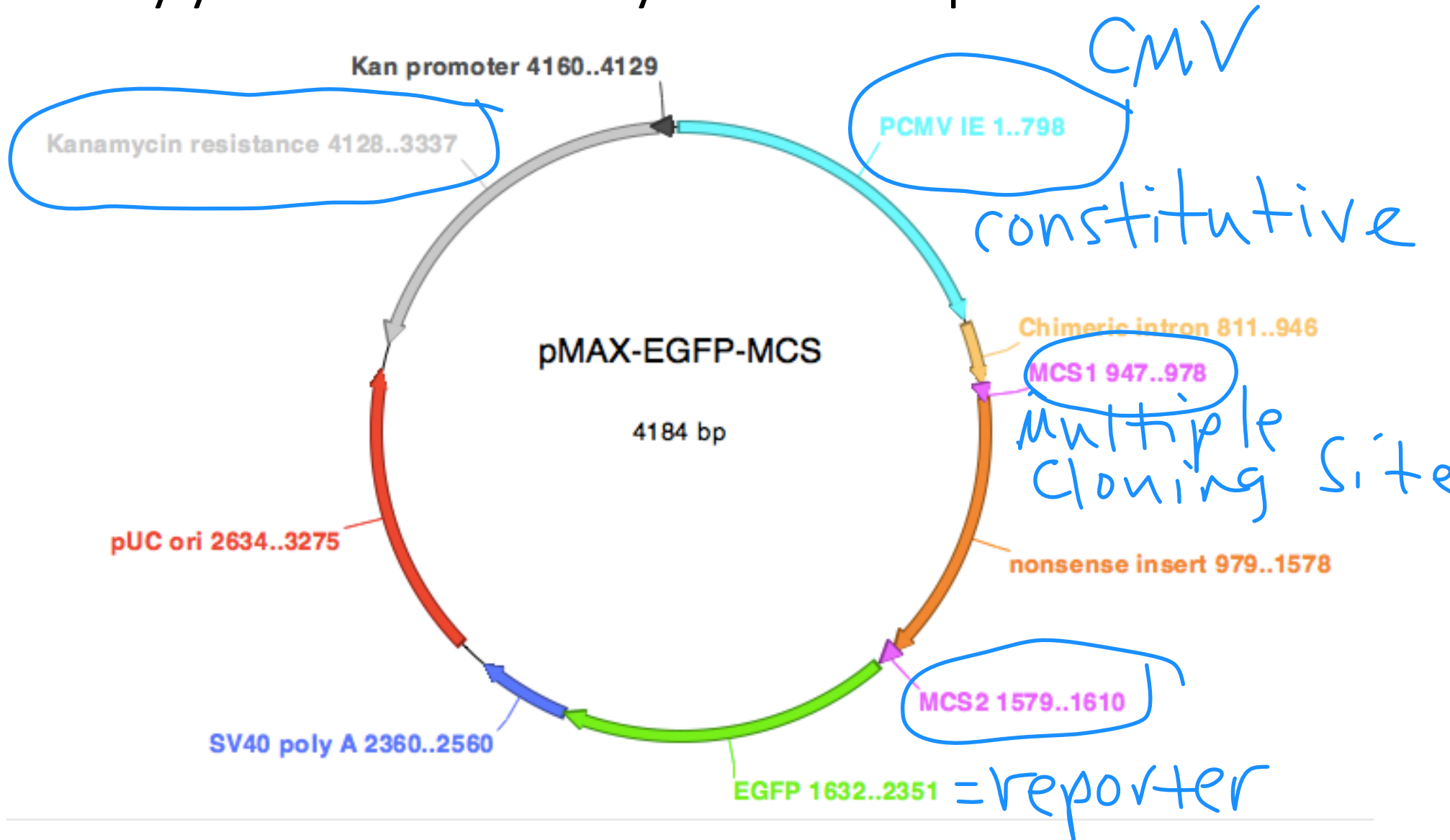
Hypothesis for NHEJ Repair capacity:

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