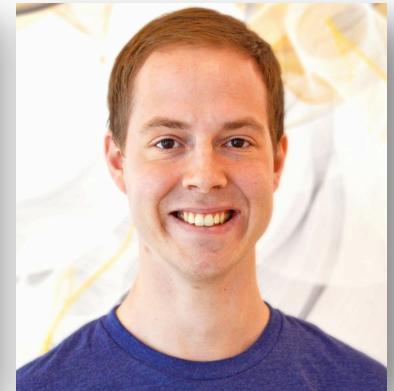


Module 2 – Lecture 2 & 3 Gene Expression Engineering (March 14 & 16, 2017)



Noreen Lyell
Leslie McLain
Maxine Jonas
Rob Wilson
Leona Samson (Lectures)

What experimental question will you ask in Module 2?

How does DNA repair affect the ability of cancer chemotherapy drugs to kill cancer cells?

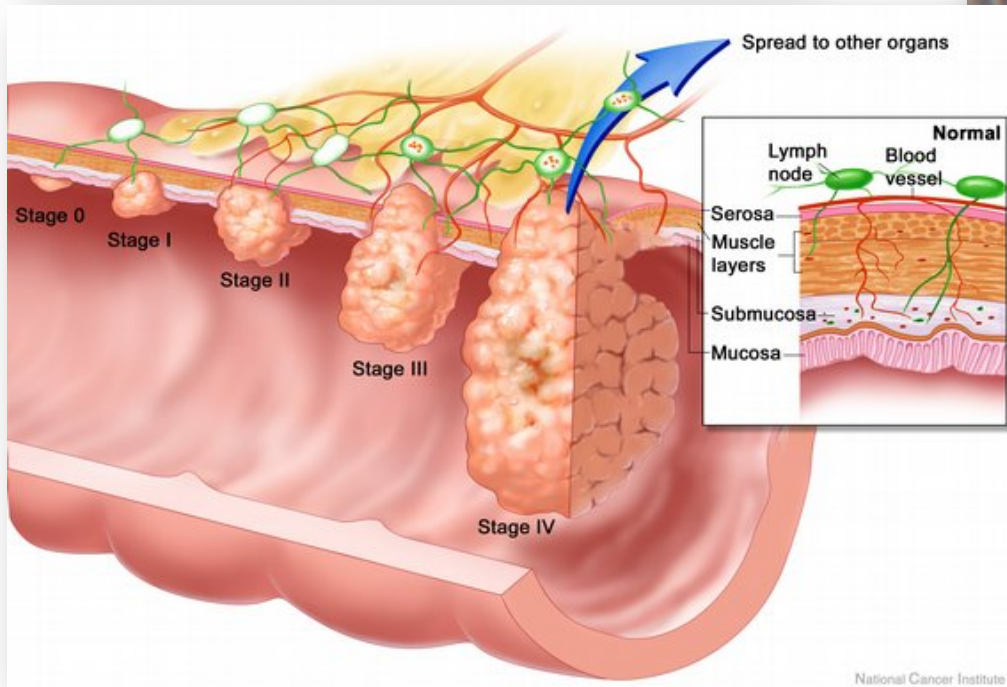
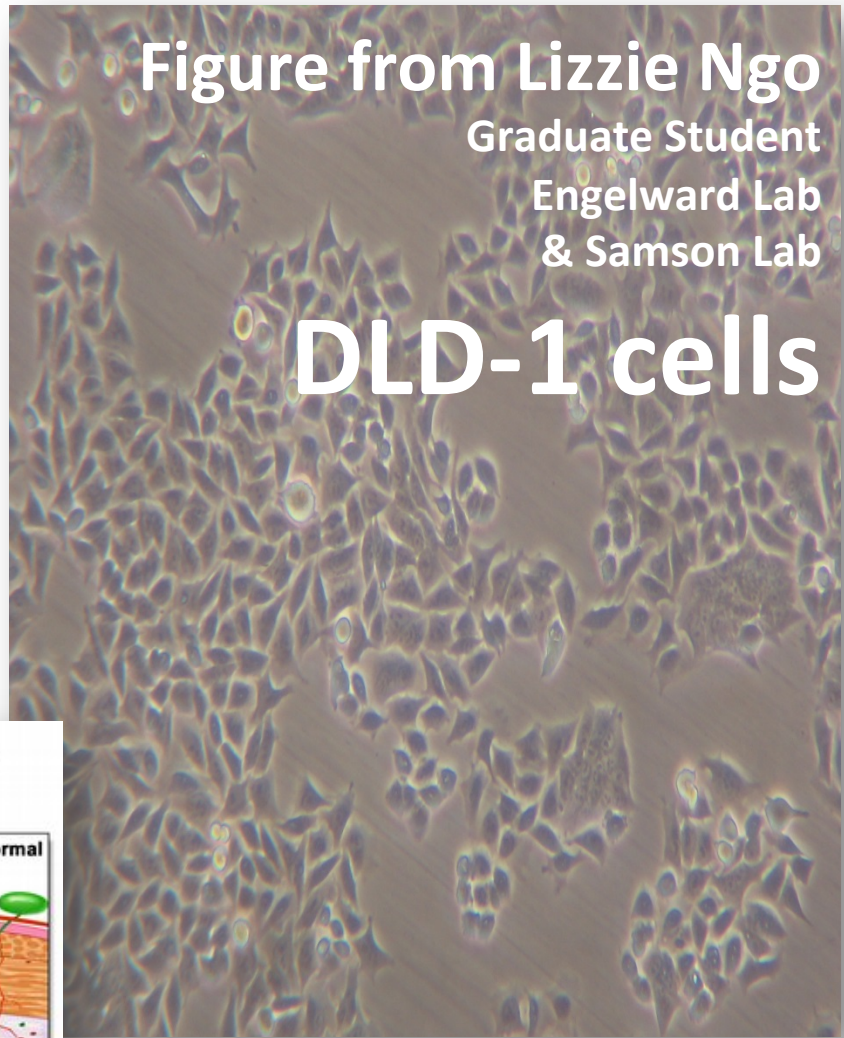
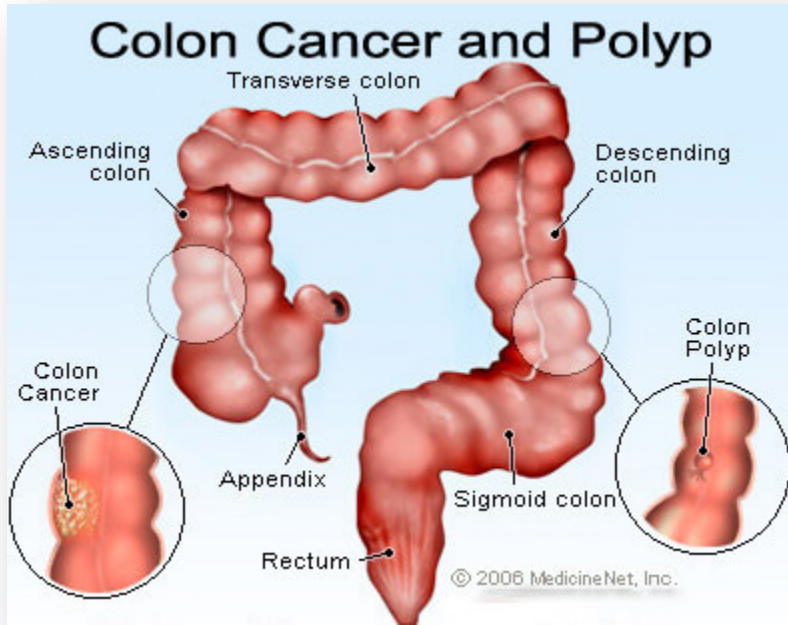
How does cancer chemotherapy affect gene expression?

This raises the following questions

- How does DNA get damaged?
- What is DNA repair?
- Why does DNA repair exist?

Key Experimental Methods for Module 2

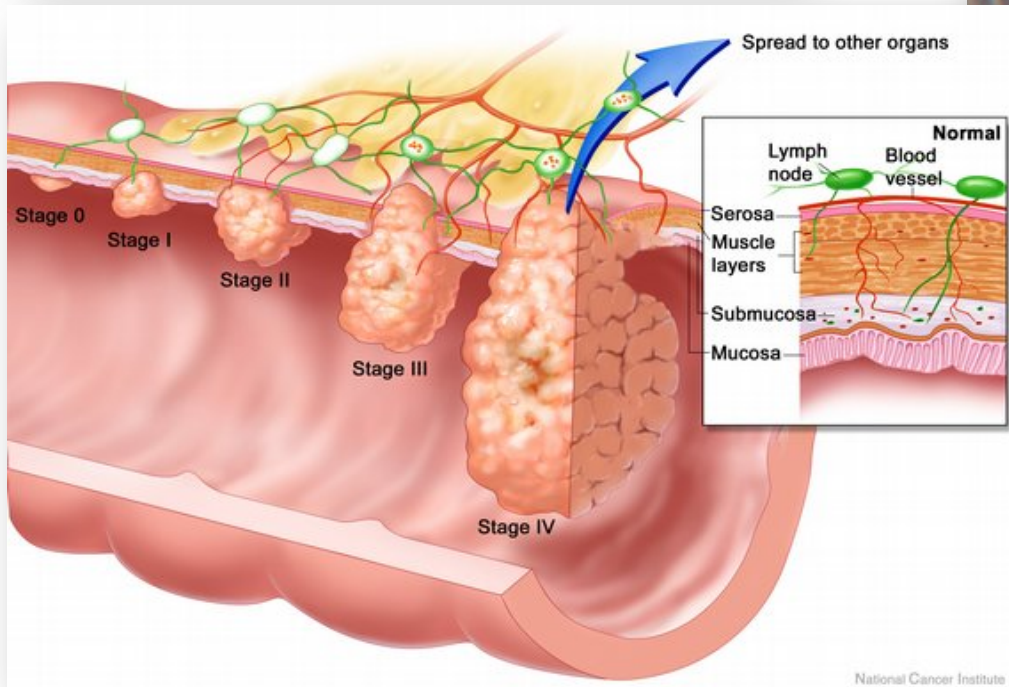
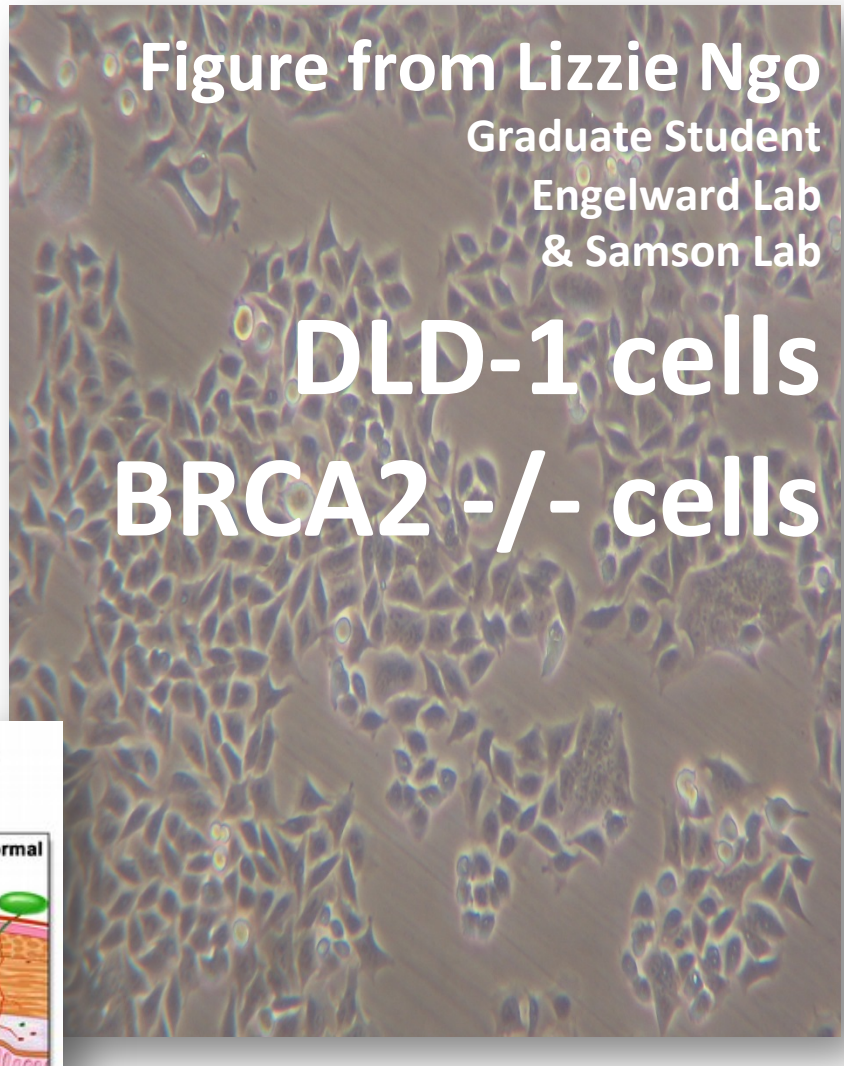
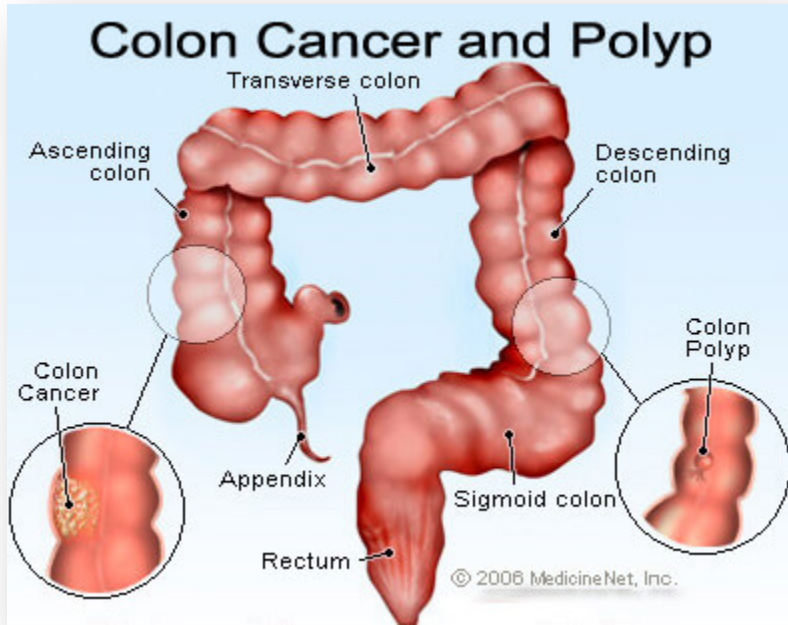
- Grow human cancer cells in tissue cell culture
- Monitor specific protein levels by Western blot
- Kill cancer cells with chemotherapy drugs
- Engineer the inhibition of DNA Repair pathways
- Monitor changes in a gene's expression (qPCR)
- Analyze RNAseq dataset measuring expression of ~ 20,000 genes (BIG DATA!)
- Statistical analysis of all biological data



[CANCER RESEARCH 39, 1020-1025, March 1979]
0008-5472/79/0039-0000\$02.00

https://www.google.com/search?q=colon+cancer&rlz=1T4GGHP_enUS635US636&source=Inms&tbm=isch&sa=X&ved=0ahUKEwj1hMLwrsfSAhWX0YMKHSFIB-EQ_AUICgB&biw=1451&bih=669&dpr=1.75#imgrc=3X3SToZHyQtBFM

https://www.google.com/search?q=colon+cancer&rlz=1T4GGHP_enUS635US636&source=Inms&tbm=isch&sa=X&ved=0ahUKEwj1hMLwrsfSAhWX0YMKHSFIB-EQ_AUICgB&biw=1451&bih=669&dpr=1.75#imgrc=SVmB_b-jLkVYQM



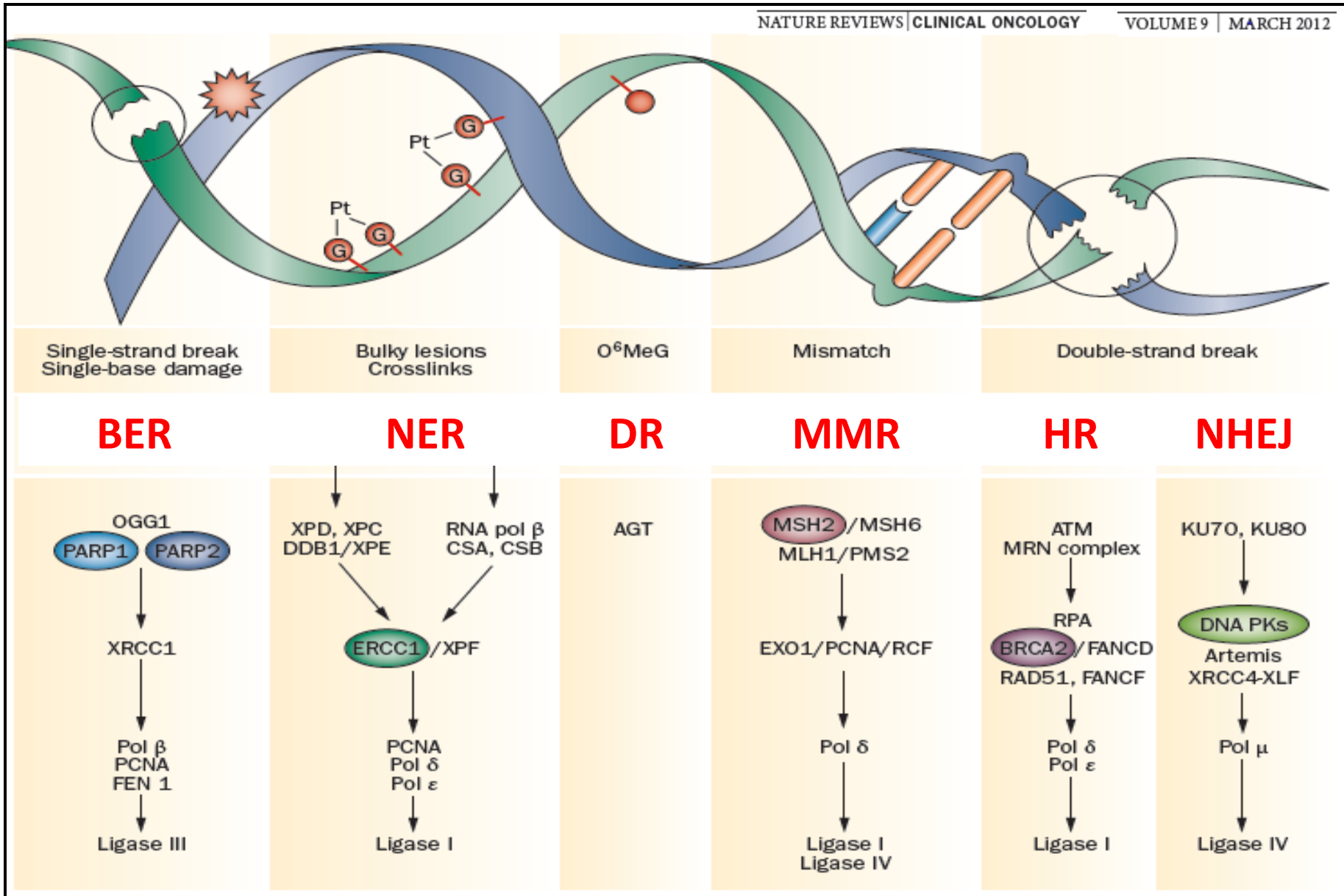
[CANCER RESEARCH 39, 1020-1025, March 1979]
0008-5472/79/0039-0000\$02.00

https://www.google.com/search?q=colon+cancer&rlz=1T4GGHP_enUS635US636&source=Inms&tbn=isch&sa=X&ved=0ahUKEwj1hMLwrsfSAhWXY0MKHSFIB-EQ_AUICgB&biw=1451&bih=669&dpr=1.75#imgcr=3X35ToZHyQtBFM

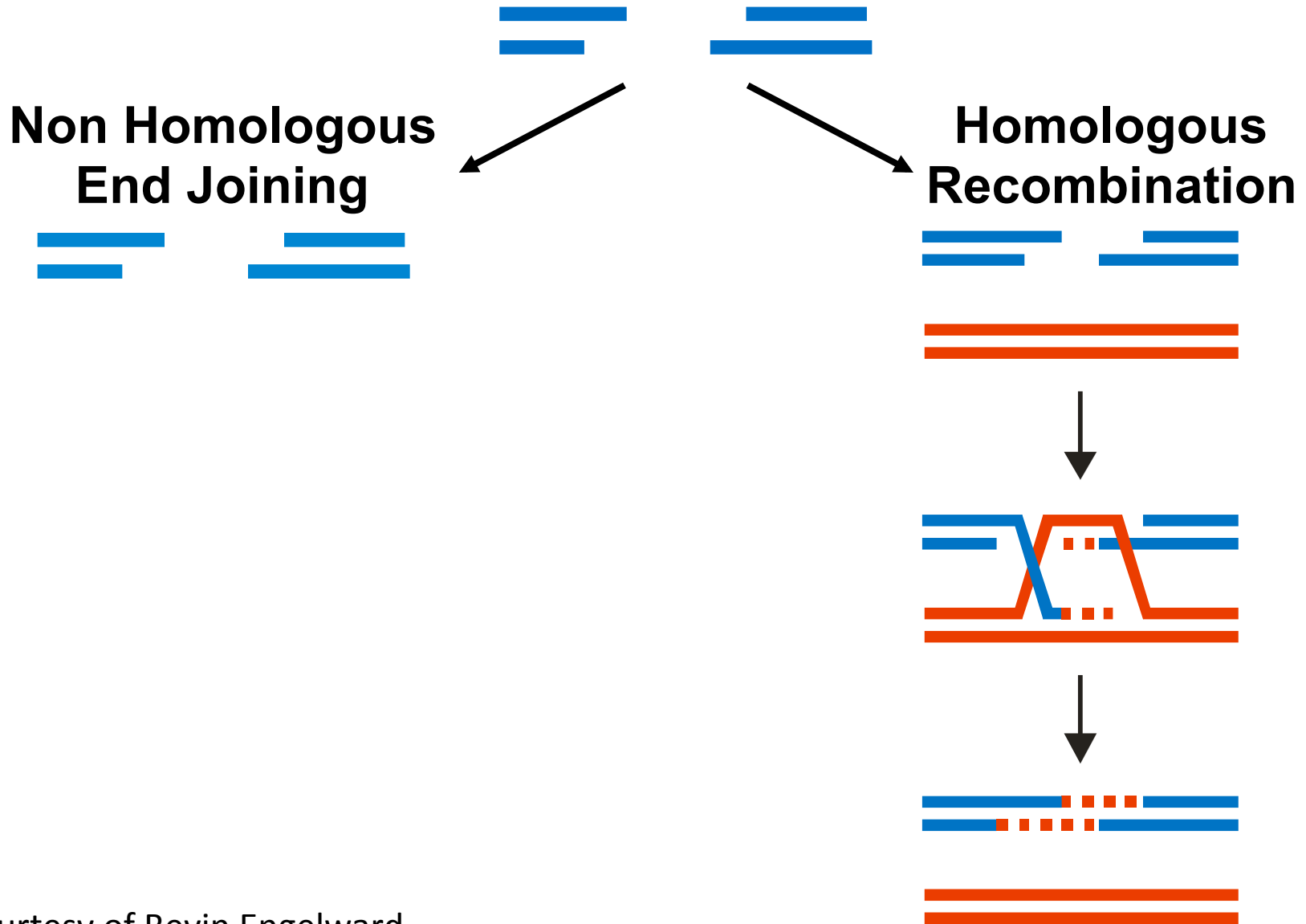
https://www.google.com/search?q=colon+cancer&rlz=1T4GGHP_enUS635US636&source=Inms&tbn=isch&sa=X&ved=0ahUKEwj1hMLwrsfSAhWXY0MKHSFIB-EQ_AUICgB&biw=1451&bih=669&dpr=1.75#imgcr=SVmB_b-jLkVYQM

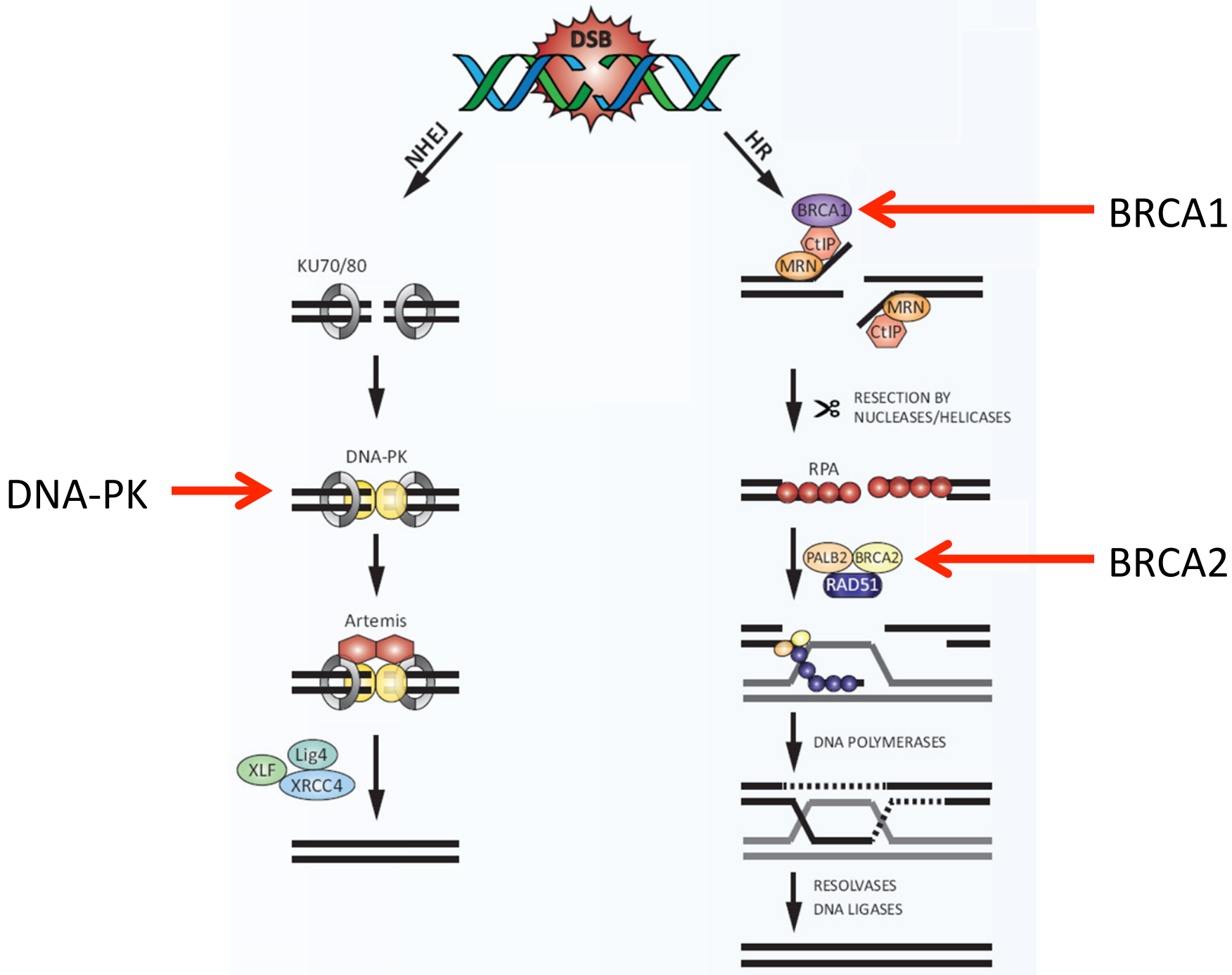
Six Major DNA Repair Pathways

NATURE REVIEWS | CLINICAL ONCOLOGY | VOLUME 9 | MARCH 2012



DNA double-strand break repair





Non-Homologous End Joining

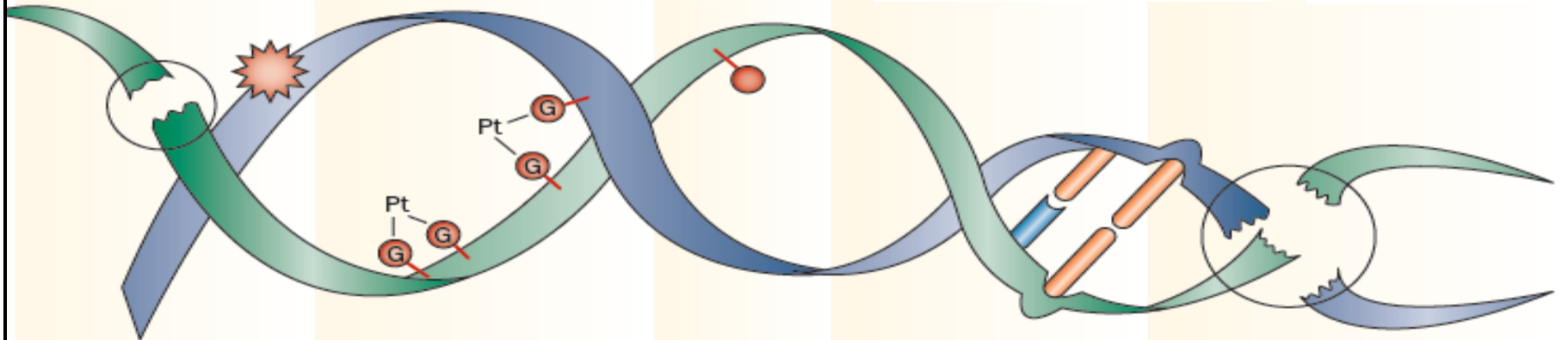
<http://web.mit.edu/engelward-lab/animations/NHEJ.html>

Synthesis-Dependent Strand Annealing (Homologous Recombination)

<http://web.mit.edu/engelward-lab/animations/SDSA.html>

Six Major DNA Repair Pathways

NATURE REVIEWS | CLINICAL ONCOLOGY | VOLUME 9 | MARCH 2012



Single-strand break
Single-base damage

Bulky lesions
Crosslinks

O⁶MeG

Mismatch

Double-strand break

BER

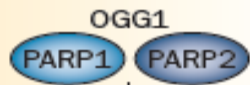
NER

DR

MMR

HR

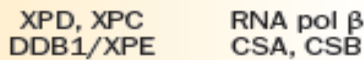
NHEJ



XRCC1

Pol β
PCNA
FEN 1

Ligase III

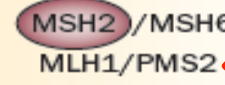


ERCC1/XPF

PCNA
Pol δ
Pol ε

Ligase I

AGT



EXO1/P

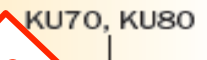
Pol δ

Ligase I
Ligase IV



Pol δ
Pol ε

Ligase I



Pol μ

Ligase IV

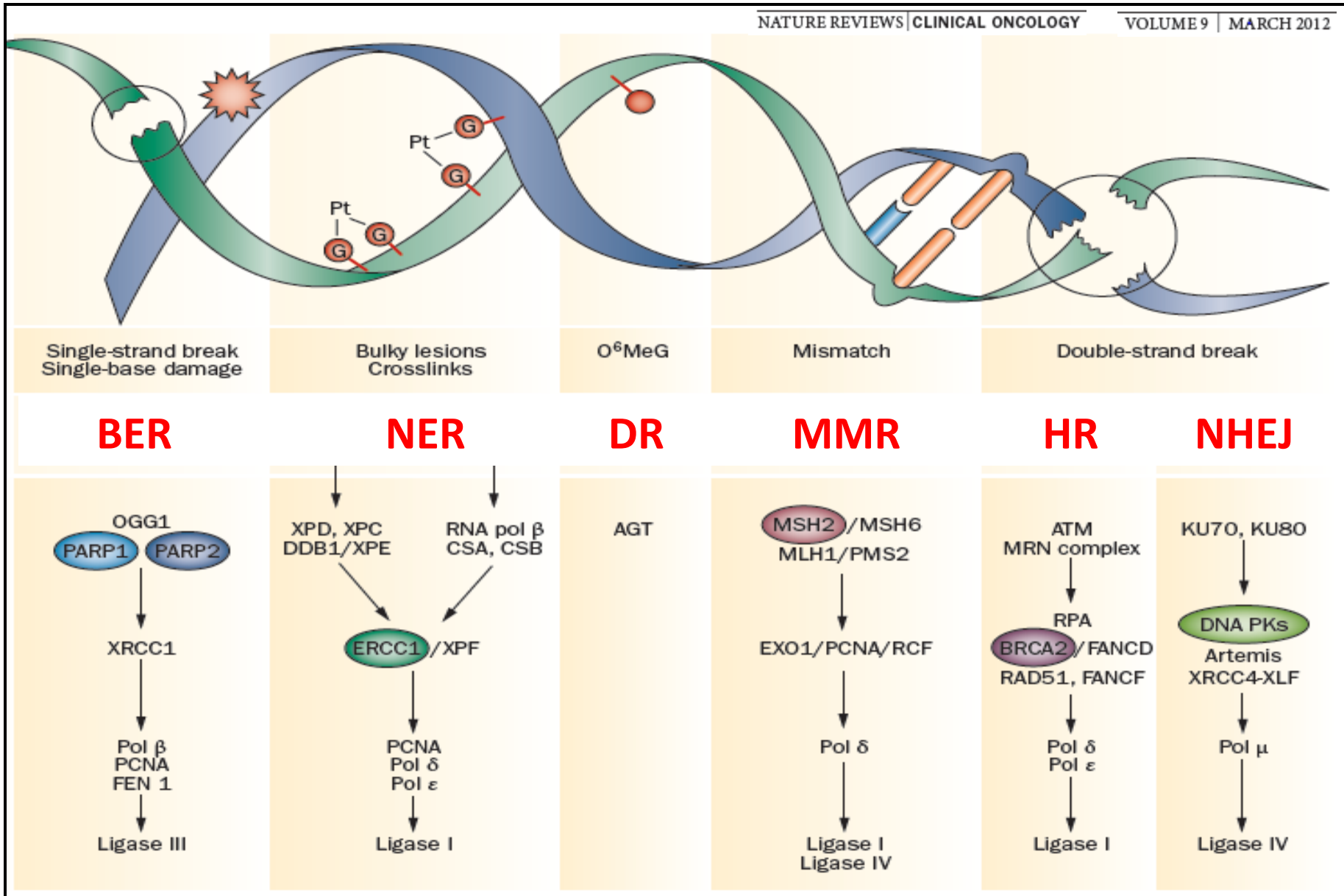
PANCREATIC CANCER

OVARIAN CANCER

BREAST CANCER

Six Major DNA Repair Pathways

NATURE REVIEWS | CLINICAL ONCOLOGY | VOLUME 9 | MARCH 2012



Mary-Claire King – Established genetic susceptibility to Breast Cancer



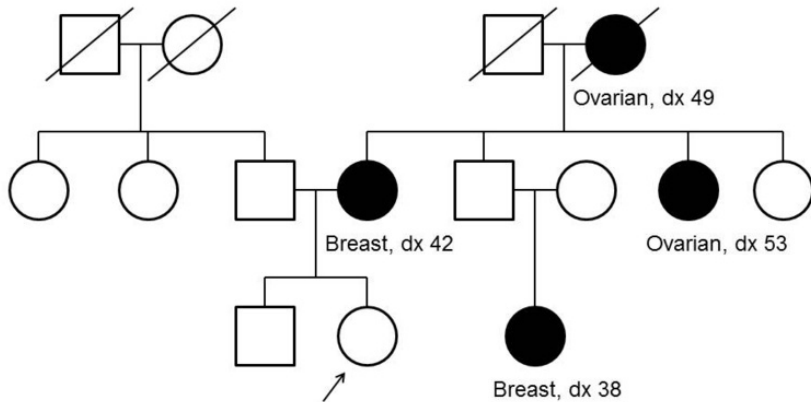
1974 – 1990 mapping gene to Chromosome 17 in Breast Cancer Families (17 years!!)

1994 BRCA1 gene was pinpointed on Chr 17

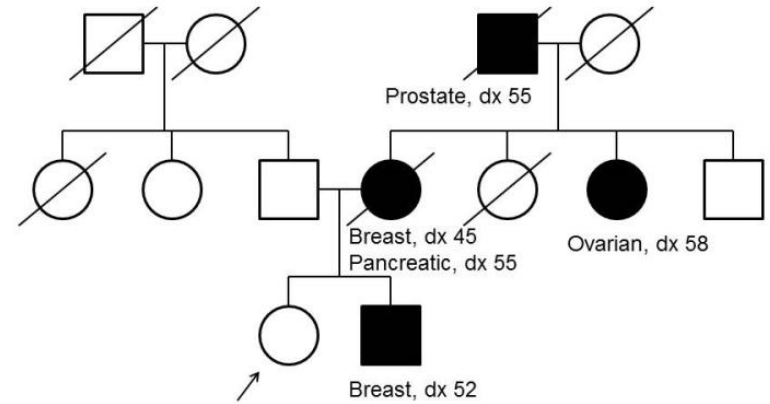
1994 BRCA2 gene was pinpointed on Chr 13

Mary-Claire King – Established genetic susceptibility to Breast Cancer

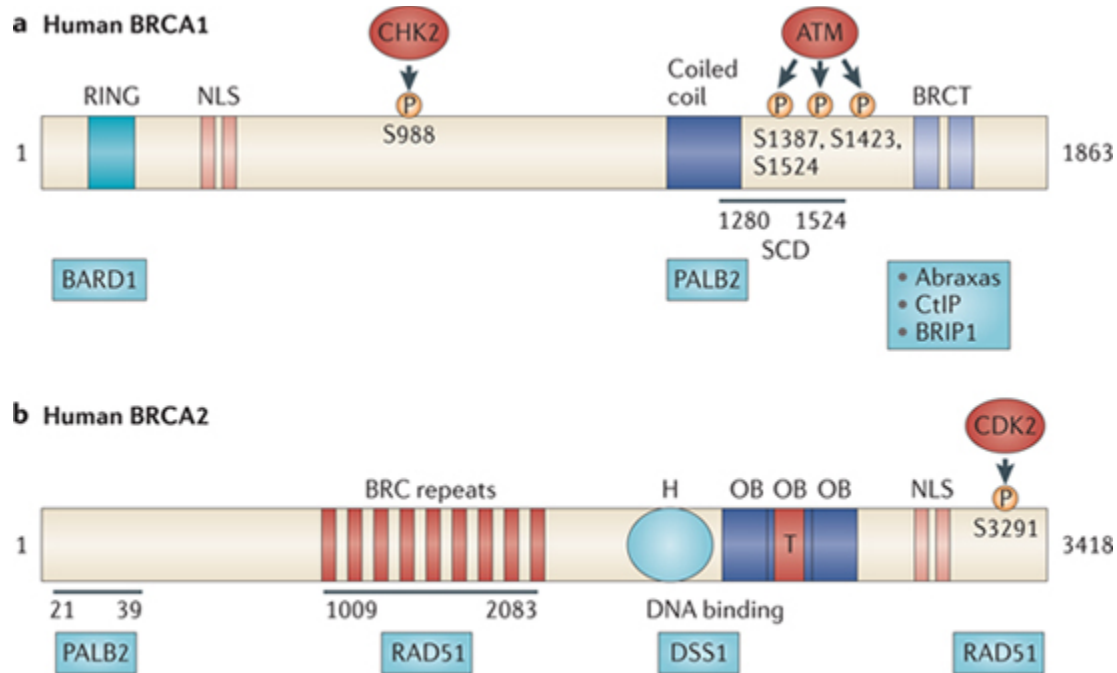
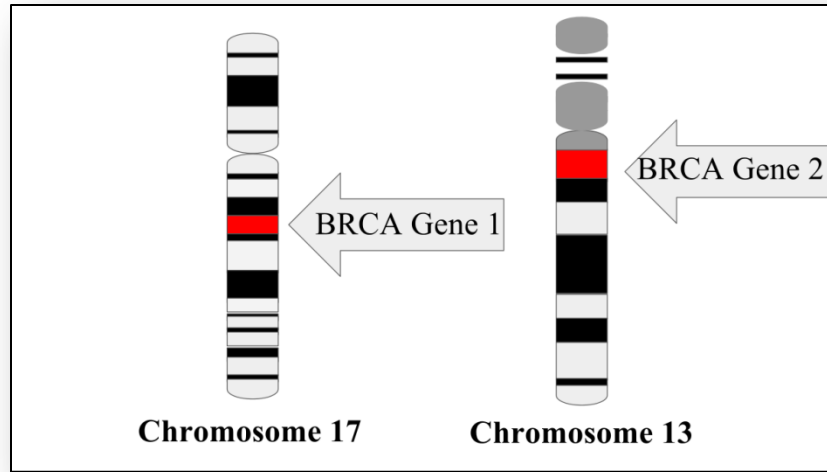
Classic *BRCA1* Pedigree

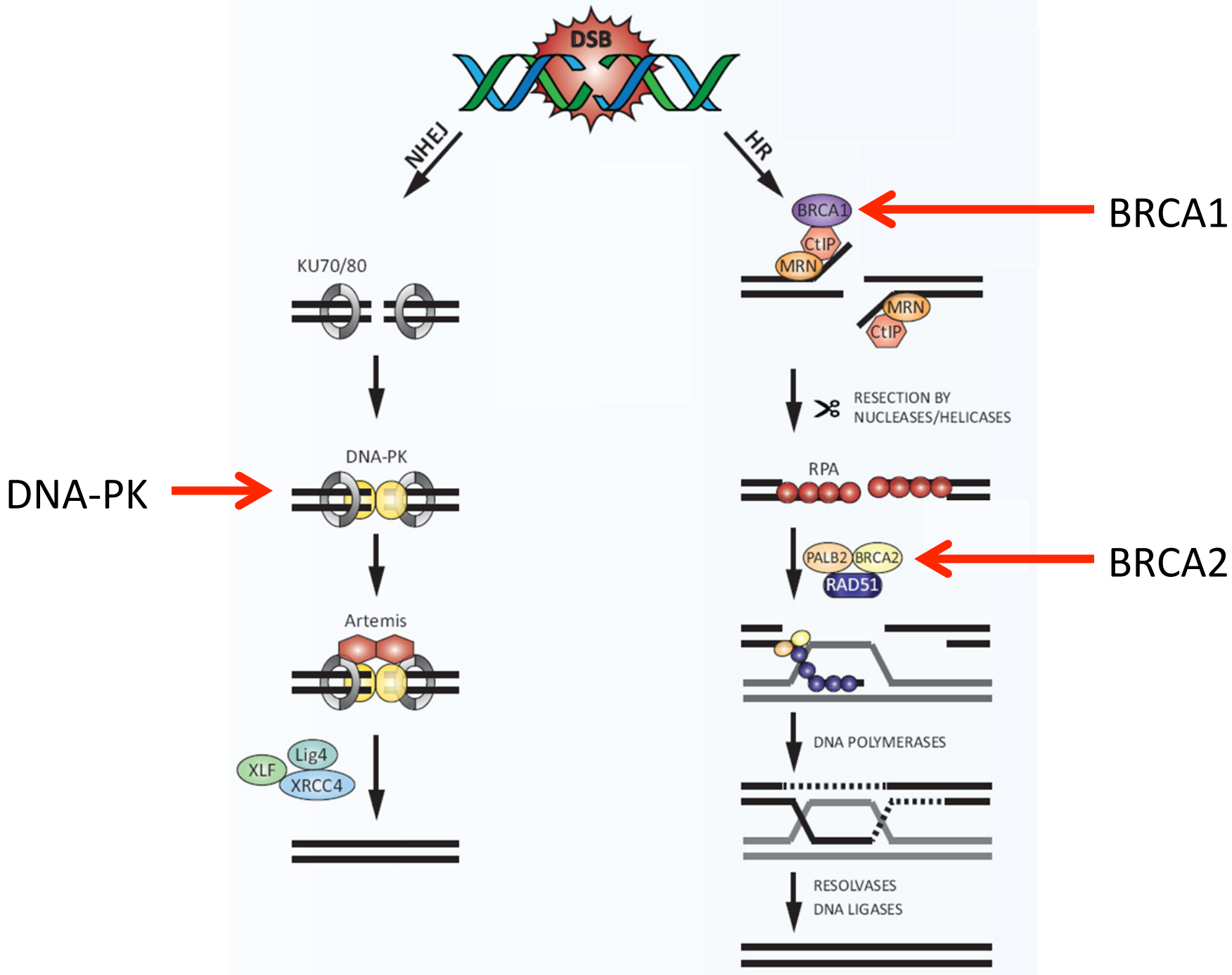


Classic *BRCA2* Pedigree

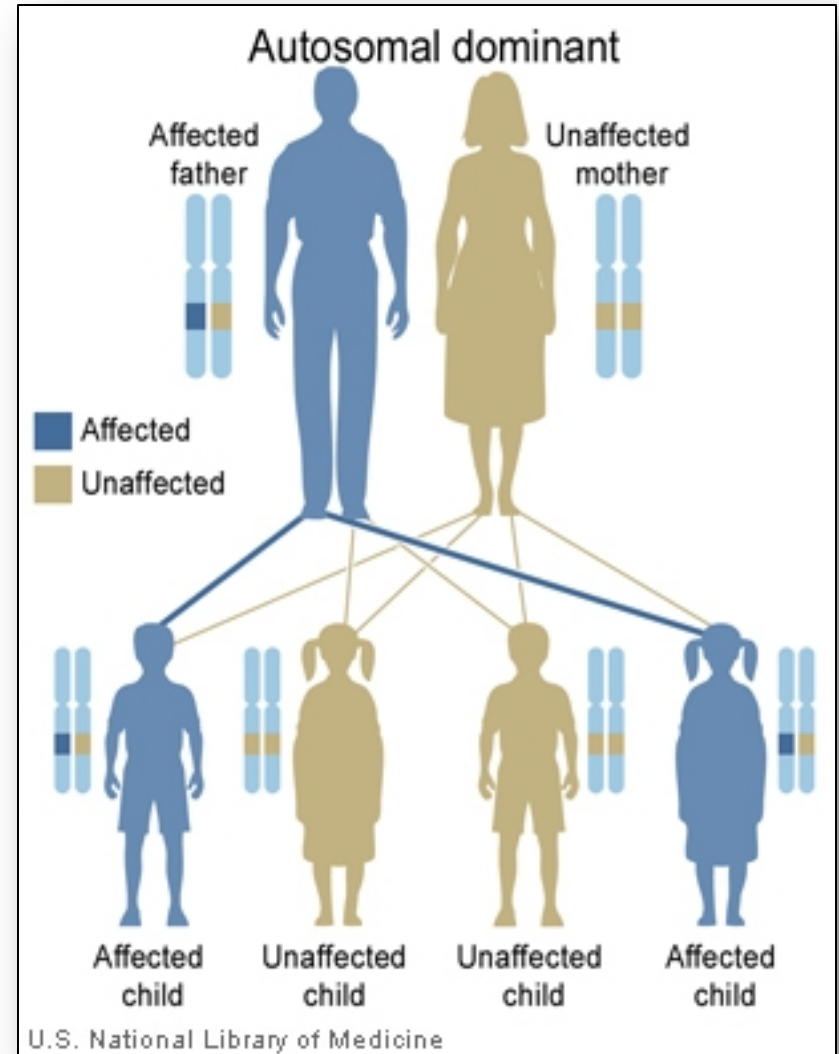
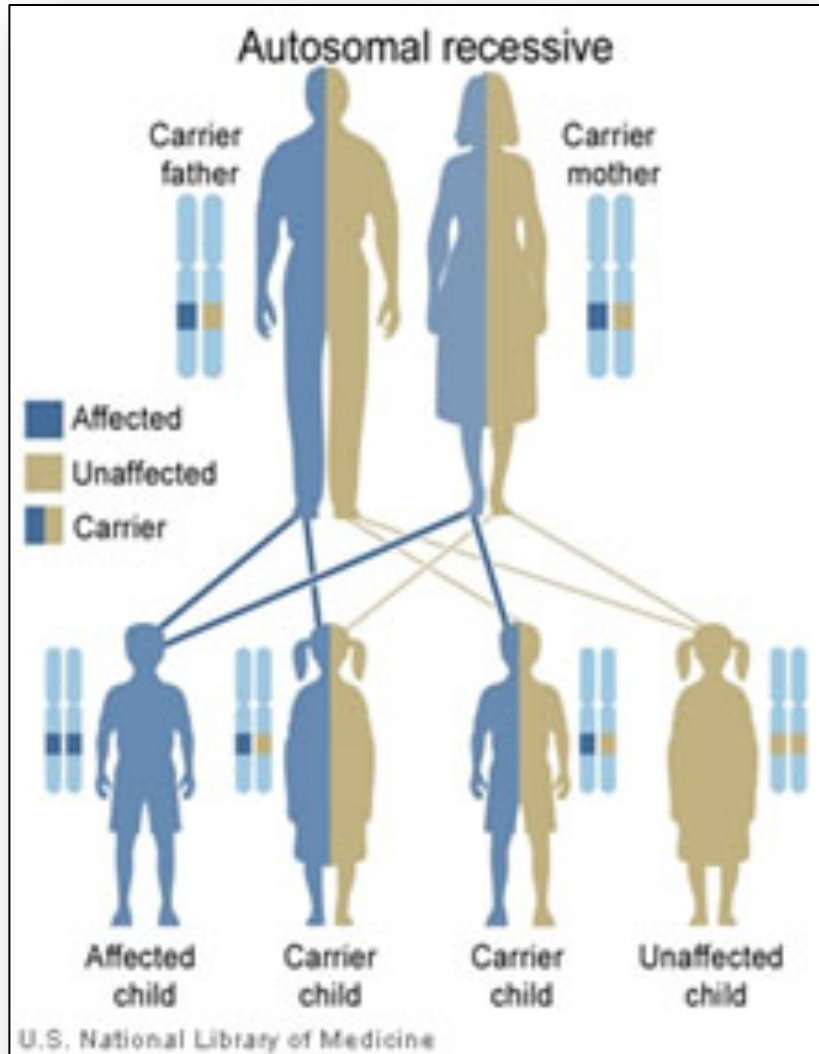


<https://www.dnalc.org/view/15126-Using-pedigress-in-the-hunt-for-BRCA1-Mary-Claire-King.html>

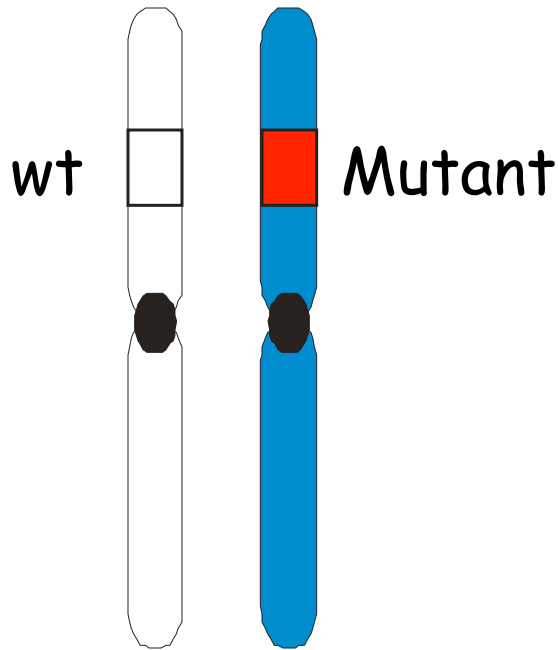




BRCA1 and *BRCA2* mutations behave as autosomal dominant



How is the second BRCA1/2 allele rendered non-functional?



Heterozygous for BRCA mutation

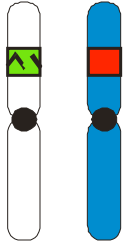


Loss of Heterozygosity

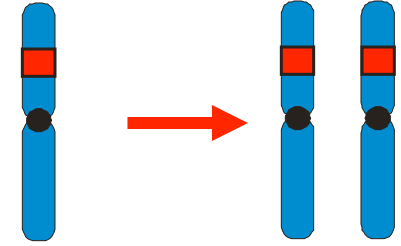
LOH

This can happen in several ways

Point Mutation



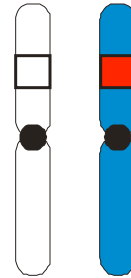
Non-Disjunction



Chromosome loss

Chromosome loss & duplication

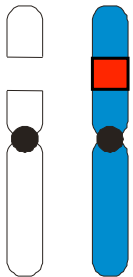
wt



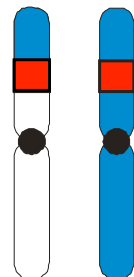
Mutant



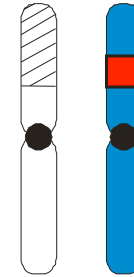
Recombination



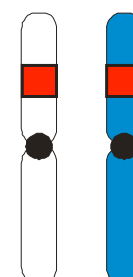
Deletion



Interchromosomal Recombination



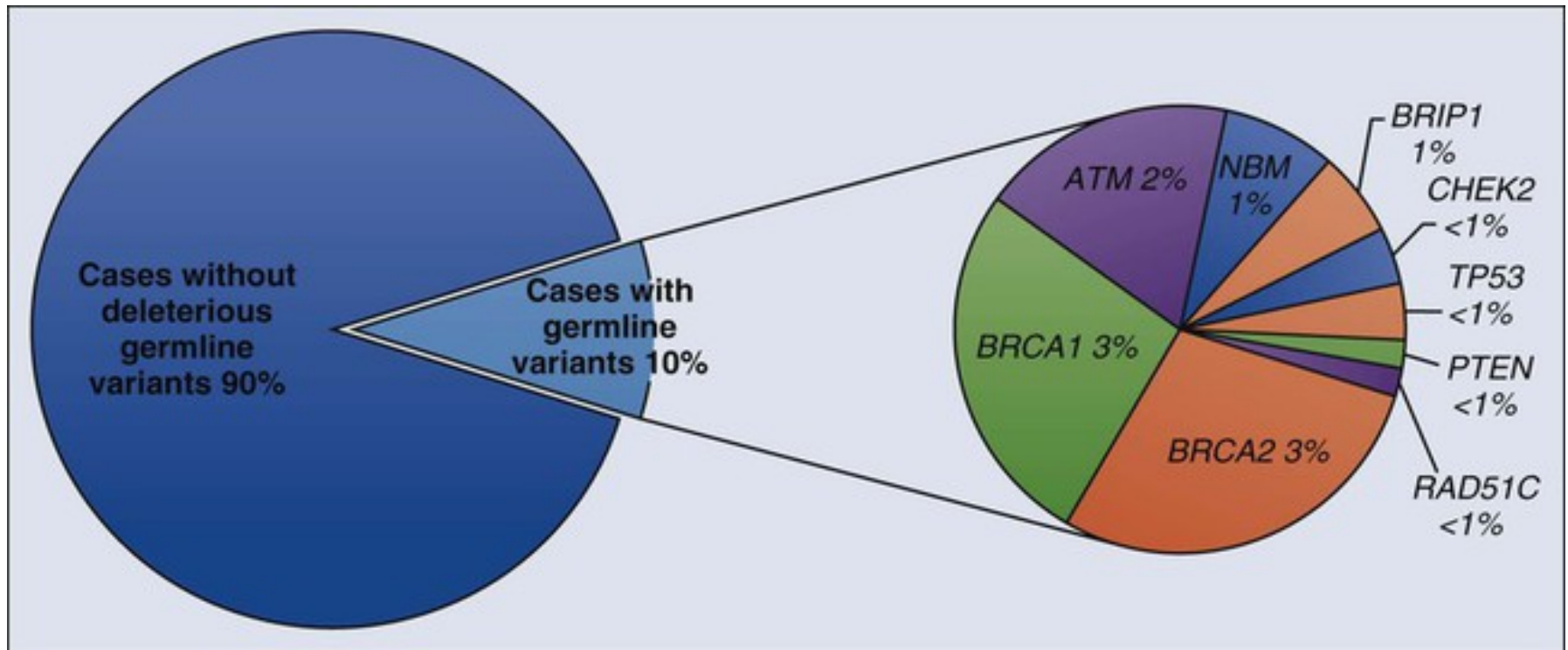
Translocation



Gene Conversion

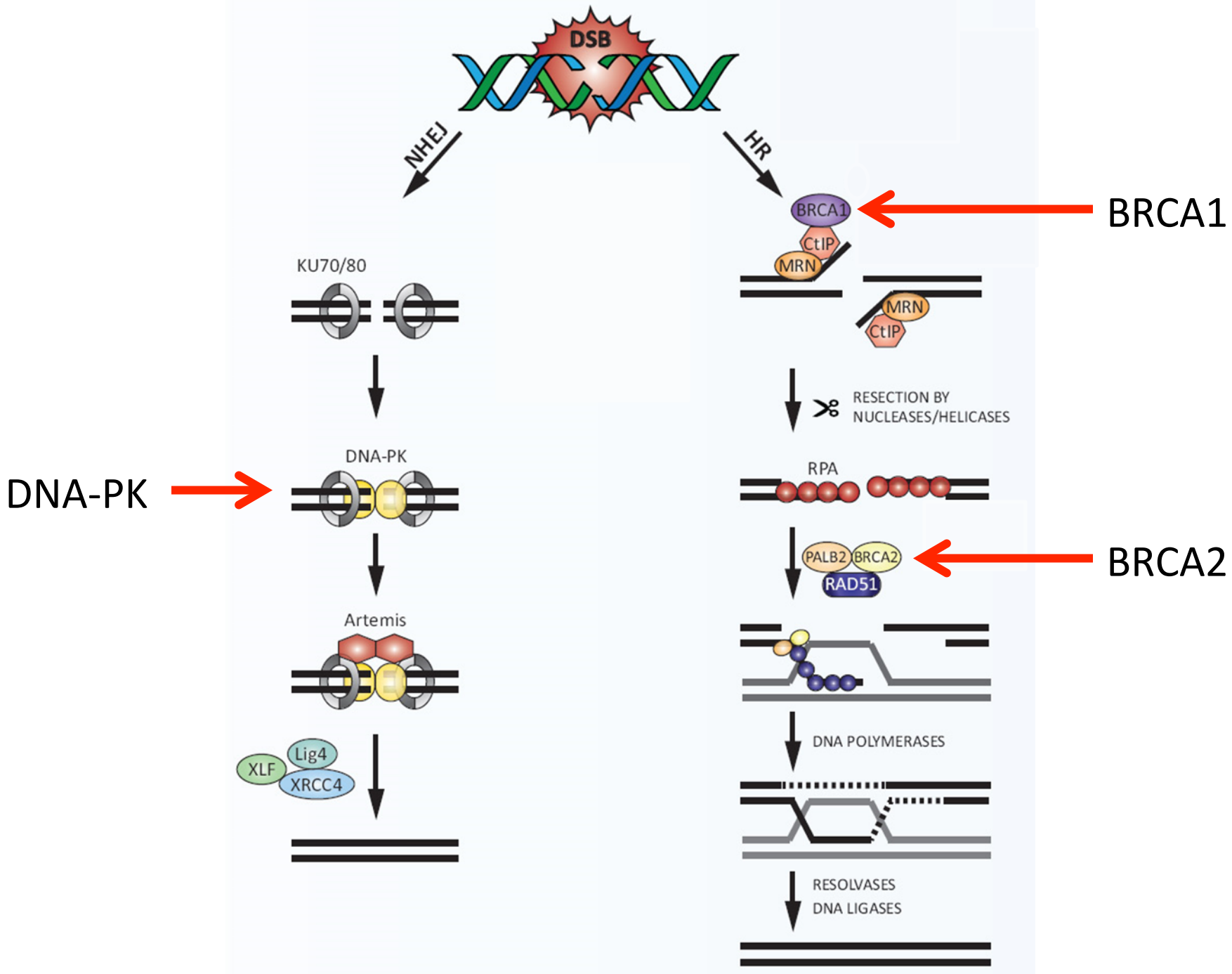
Familial Breast Cancers ~ 10%

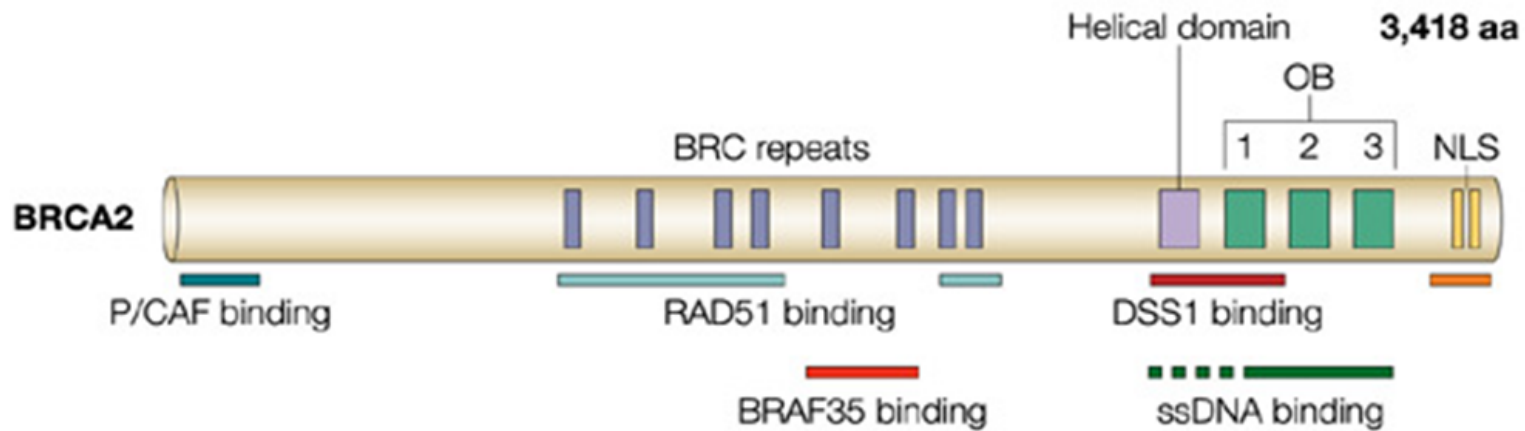
Sporadic Breast Cancers ~ 90%



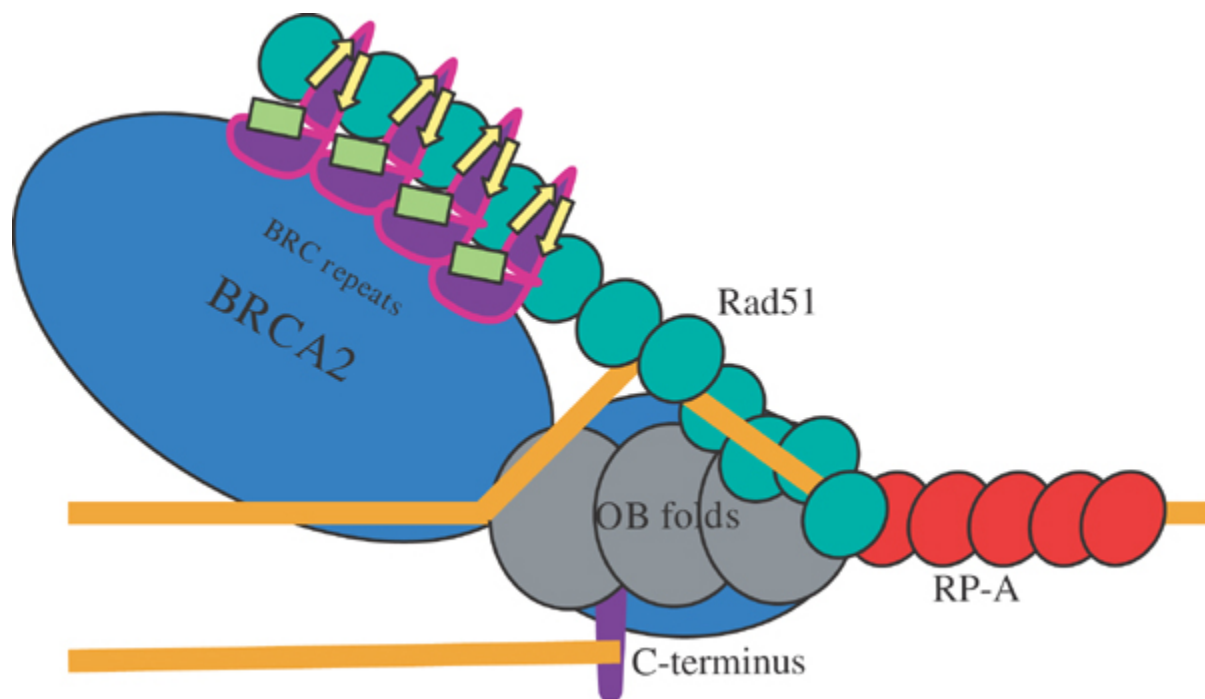
Total breast cancer cases (n = 507) analyzed in The Cancer Genome Atlas with deleterious germline variants. (Cancer Genome Atlas Network. Comprehensive molecular portraits of human breast tumours. Nature 2012;490:61–70.) Kasmintan A. Schrader, Ravi Sharaf, Shaheen Alanee and Kenneth Offit

<http://clinicalgate.com/genetic-factors-hereditary-cancer-predisposition-syndromes-2/>





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GeneArt™ Engineered Cell Models from Thermo Fisher Scientific



The largest collection of ready-to-go CRISPR engineered cell lines

The GeneArt Engineered Cell Models have been engineered using the latest genome editing tools including CRISPRs and rAAV editing to create both knockout and knock-in models. Each cell line ships with the unedited, isogenic parental cell lines providing the perfect control.

GeneArt Engineered Cell Models provide researchers with instant access to a collection of thousands of affordable, pre-engineered cell lines that will accelerate their research programs and eliminate the risk, cost, and time associated with developing their own cell lines or commissioning custom cell line development program.

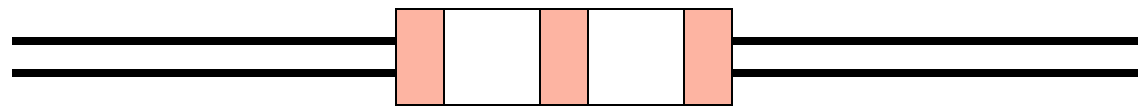
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Thermo Fisher Scientific for
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Models*

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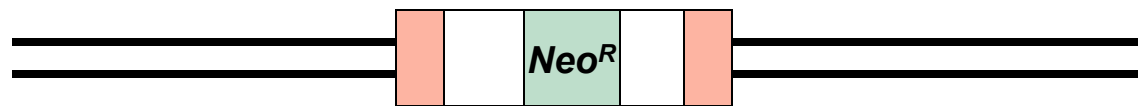
[Supplier Page](#)

Gene X e.g., *BRCA2* gene – in the test tube

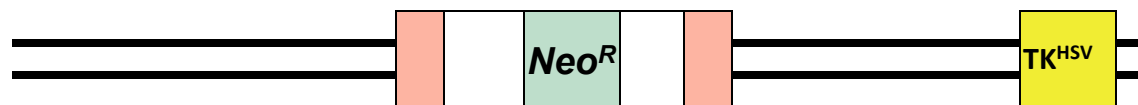


exons

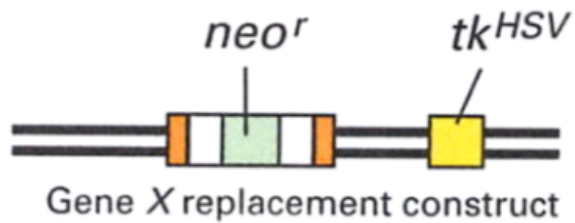
Replace an exon with a **positive** selectable marker, e.g., Neo^R



Mutated or “Knocked out” gene construct



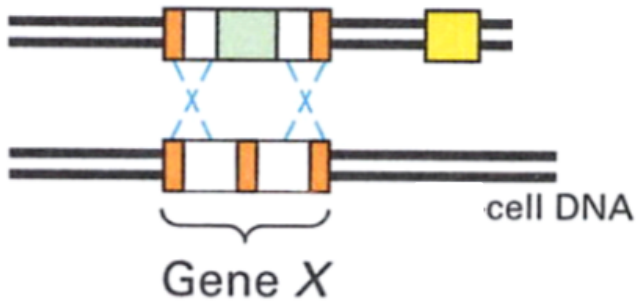
Add some distance away, a **negative** selectable marker, e.g., TK^{HSV}



0.01% of the time

Homologous recombination

Cells



Gene-targeted insertion

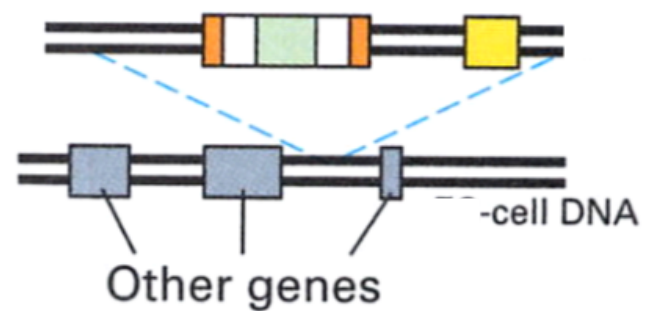


Cells are resistant to G-418 and ganciclovir

Cells

Nonhomologous recombination

99.9% of the time

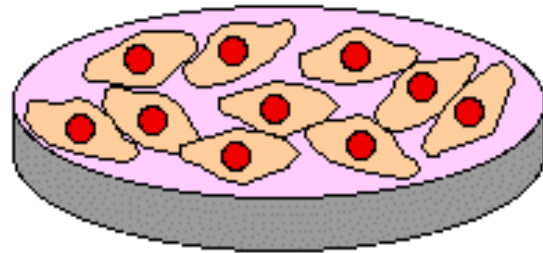
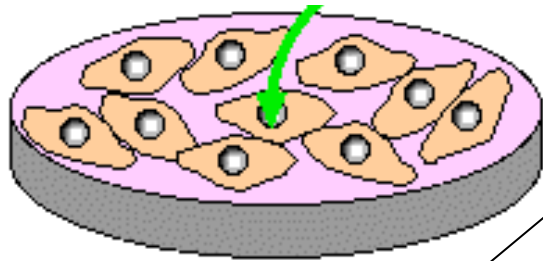
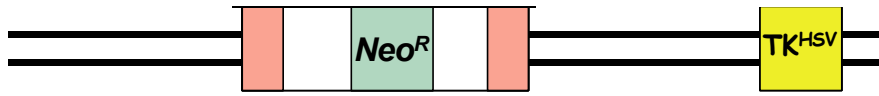


Random insertion



Cells are resistant to G-418 but sensitive to ganciclovir

Targeting Construct

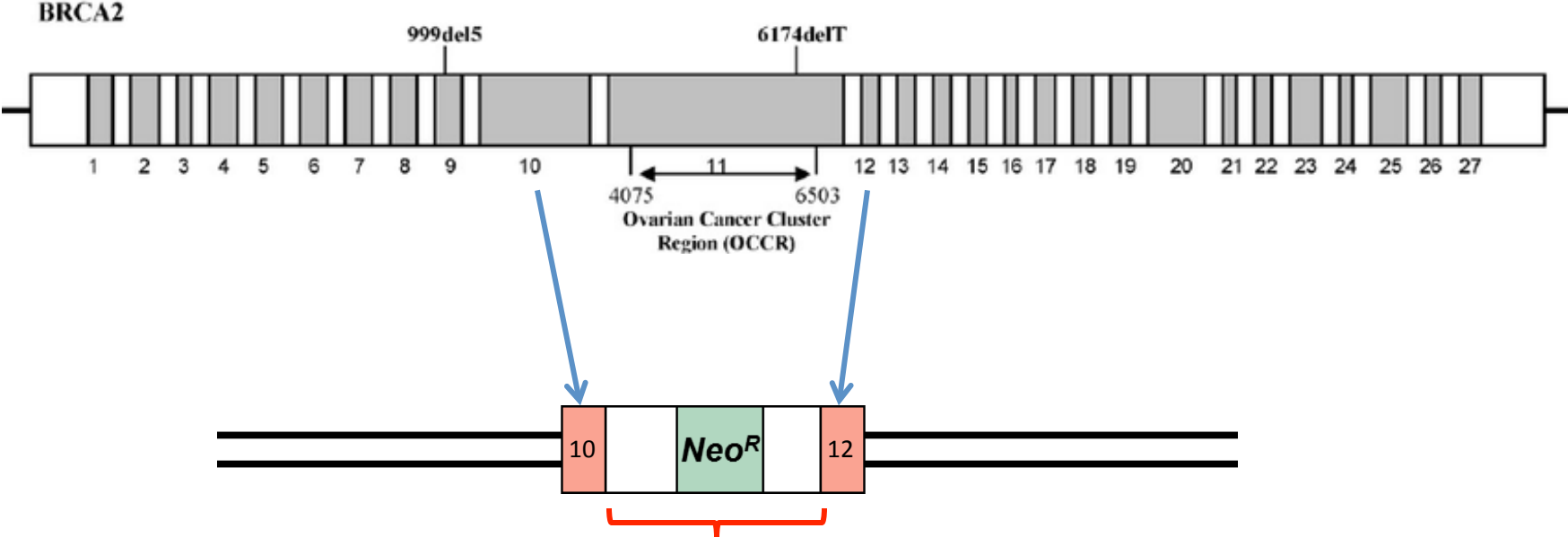


Select for the genetically altered cells you want

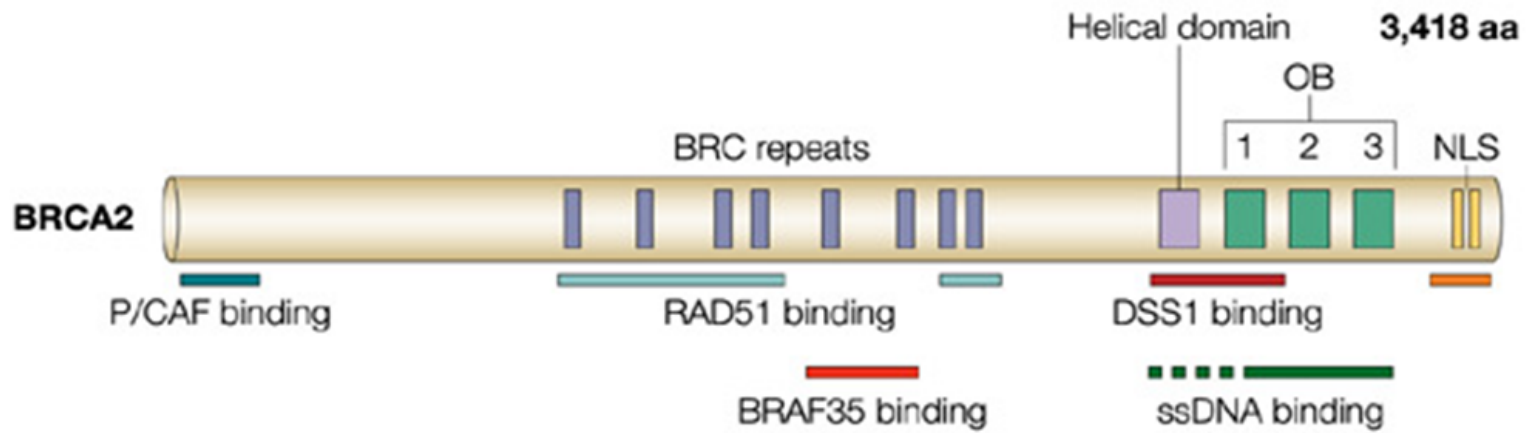
Select **for** the Neo^R gene and **against** the TK^{HSV} gene using **Neomycin** and **Gancyclovir**

The only cells to survive have undergone a targeted homologous recombination event at the gene of interest

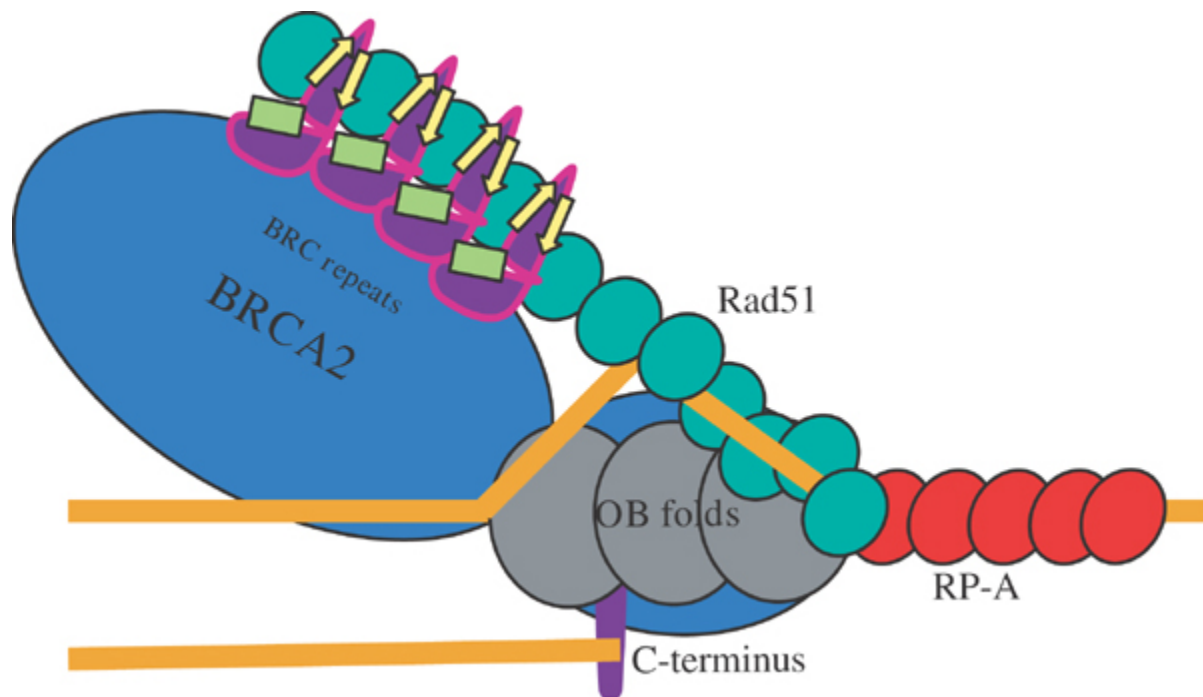
BOTH *BRCA2* alleles have exon 11 replaced with a selectable marker gene (for Neomycin resistance)



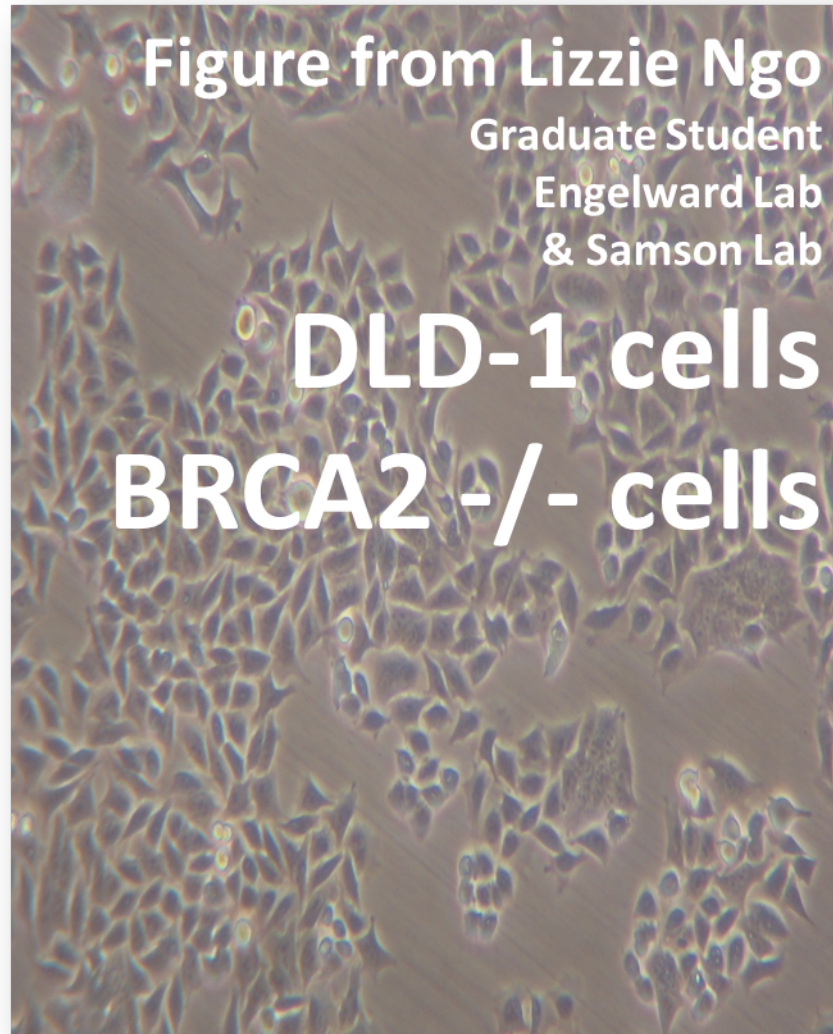
Exon 11 replaced with selectable marker



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BOTH *BRCA2* alleles have exon 11 replaced with a selectable marker gene (for Neomycin resistance)



GeneArt™ Engineered Cell Models from Thermo Fisher Scientific



The largest collection of ready-to-go CRISPR engineered cell lines

The GeneArt Engineered Cell Models have been engineered using the latest genome editing tools including CRISPRs and rAAV editing to create both knockout and knock-in models. Each cell line ships with the unedited, isogenic parental cell lines providing the perfect control.

GeneArt Engineered Cell Models provide researchers with instant access to a collection of thousands of affordable, pre-engineered cell lines that will accelerate their research programs and eliminate the risk, cost, and time associated with developing their own cell lines or commissioning custom cell line development program.

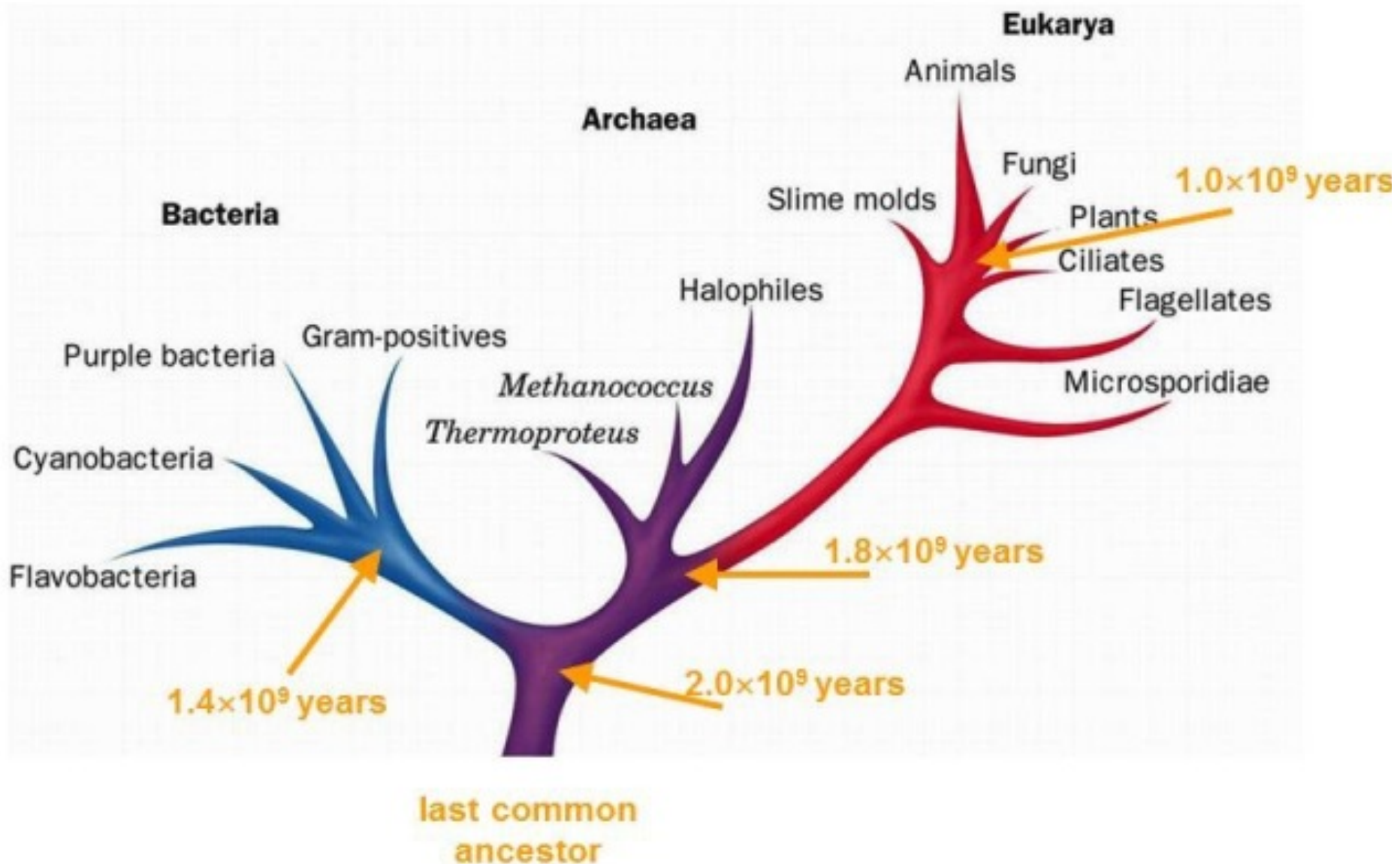
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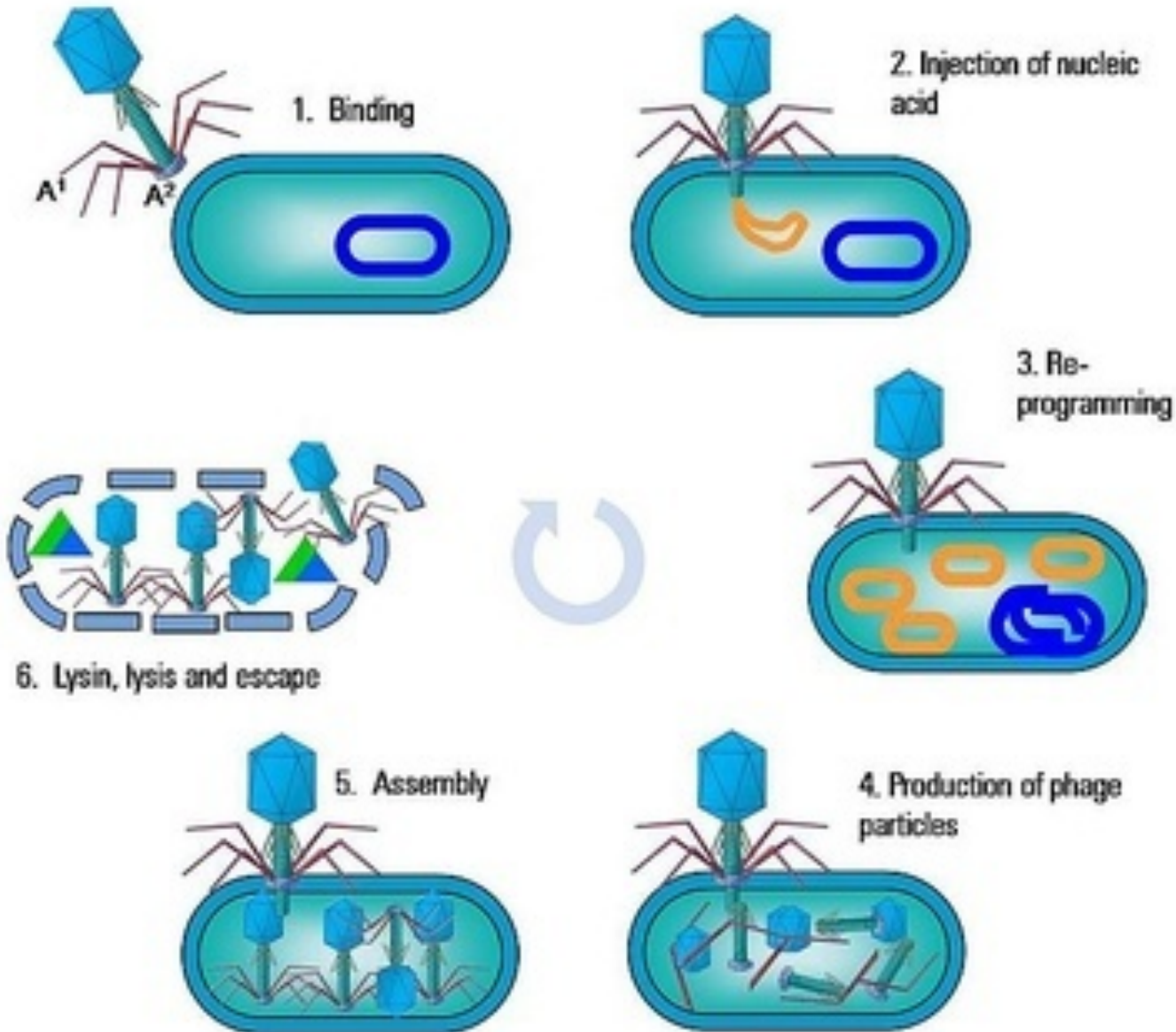
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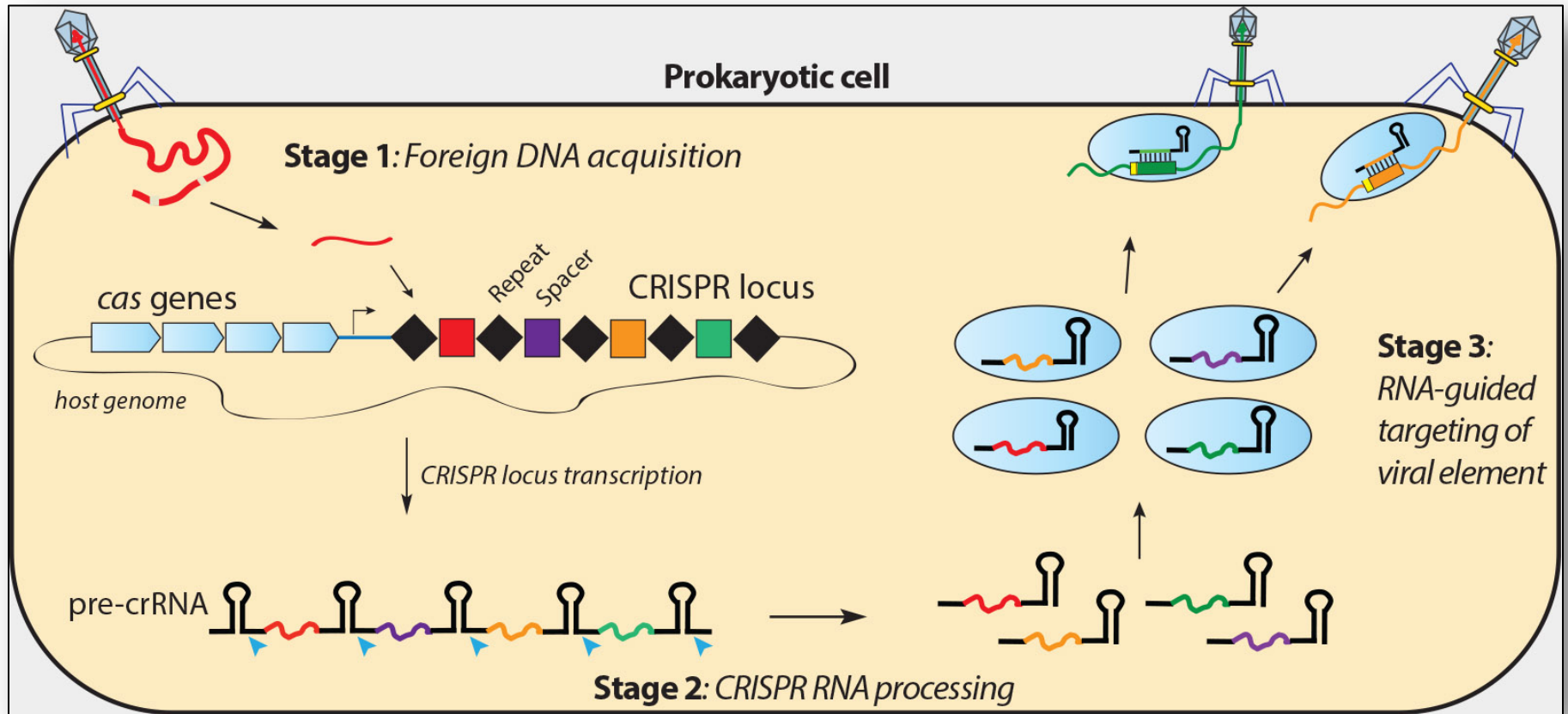
All known life forms are based on DNA



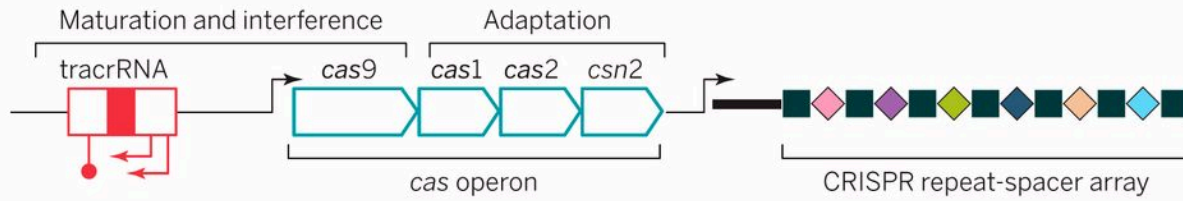
Bacteriophage (bacterial virus) infecting *E. coli*



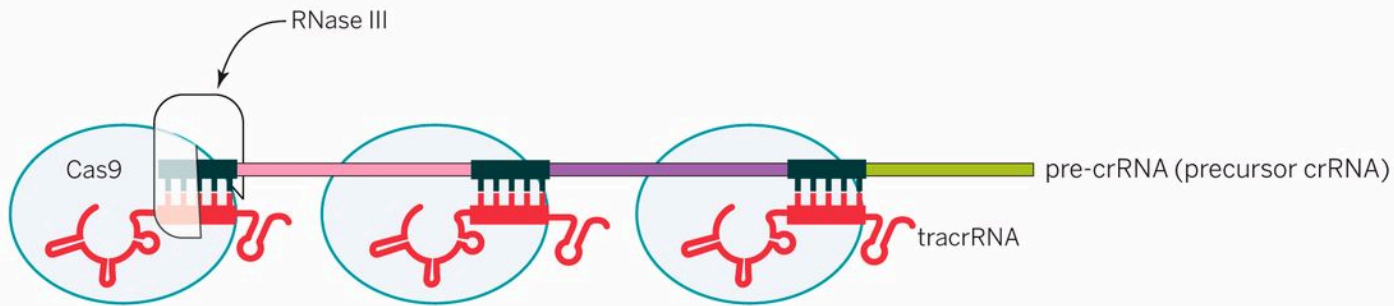
CRISPR - **C**lustered **R**egularly **I**nterspaced **S**hort **P**alindromic **R**epeats
CAS genes – **C**RISPR **A**sociated genes



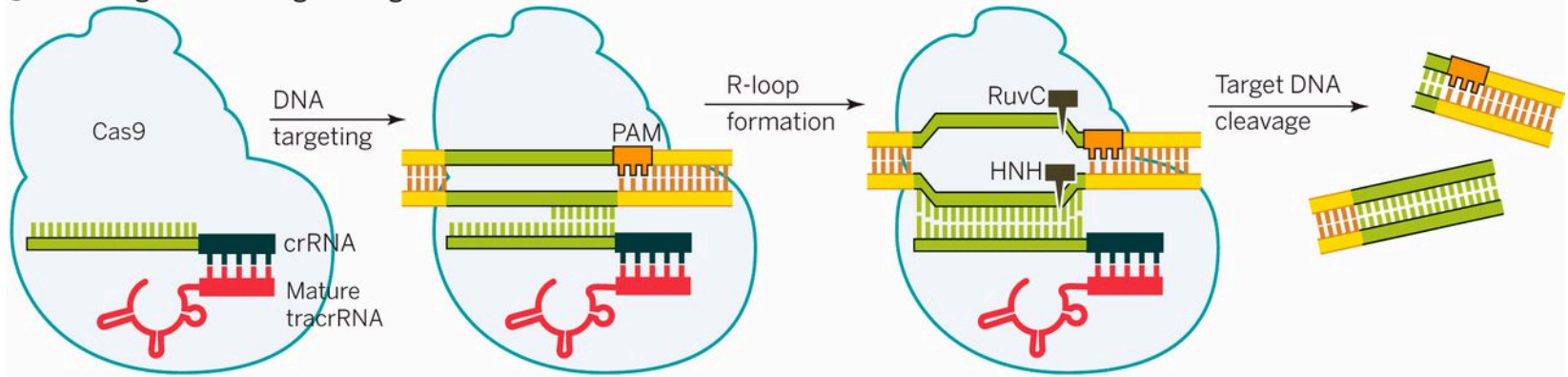
A Genomic CRISPR locus



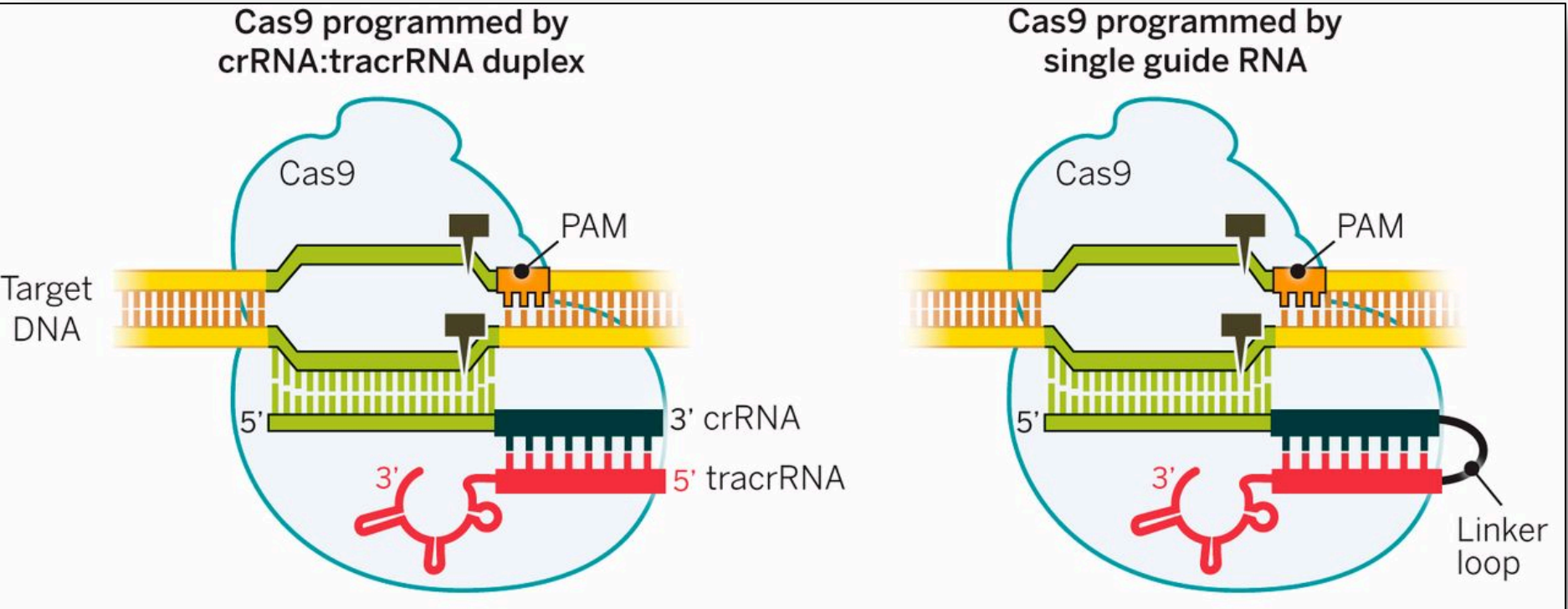
B *tracrRNA*:*crRNA* co-maturation and Cas9 co-complex formation



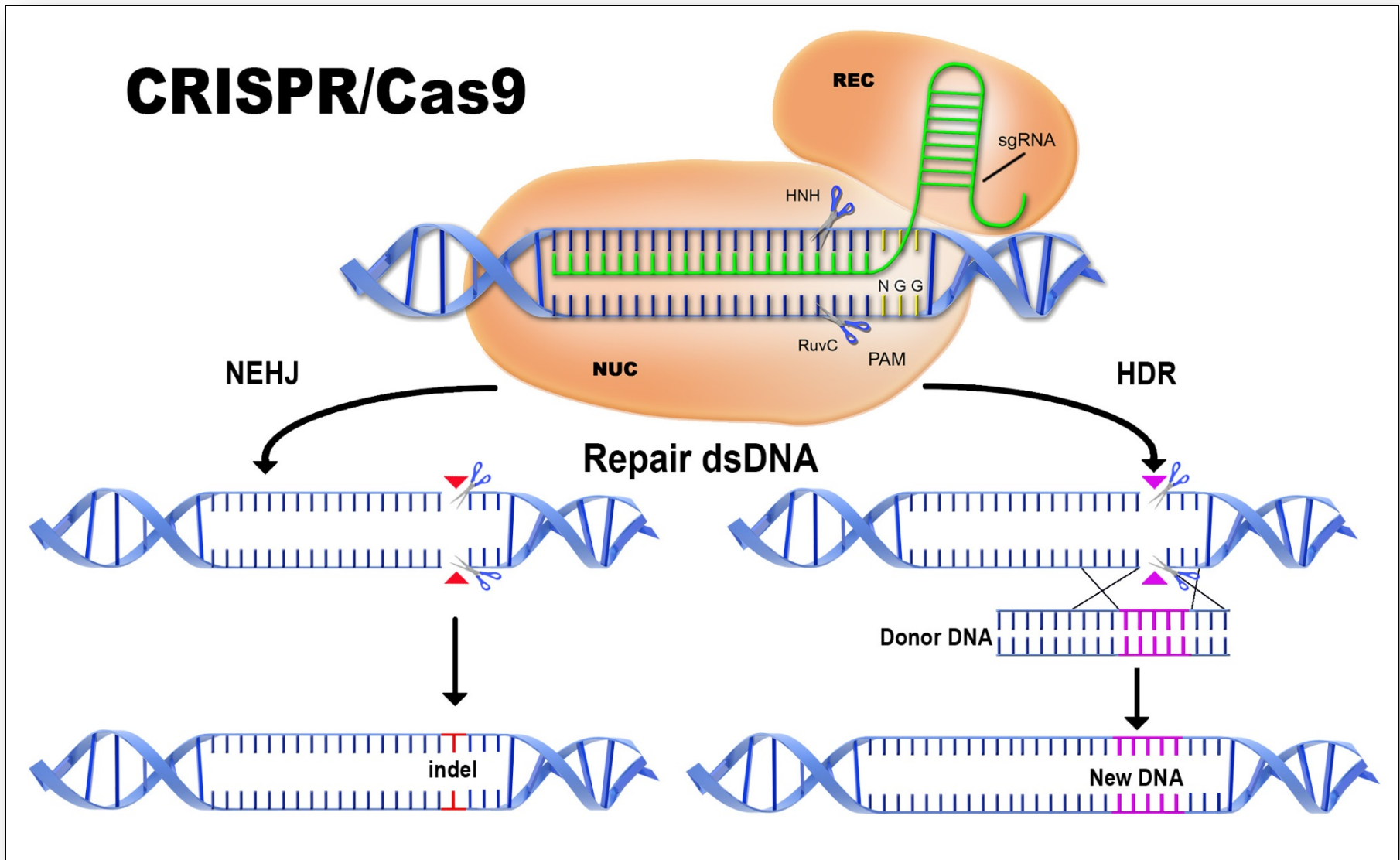
C RNA-guided cleavage of target DNA



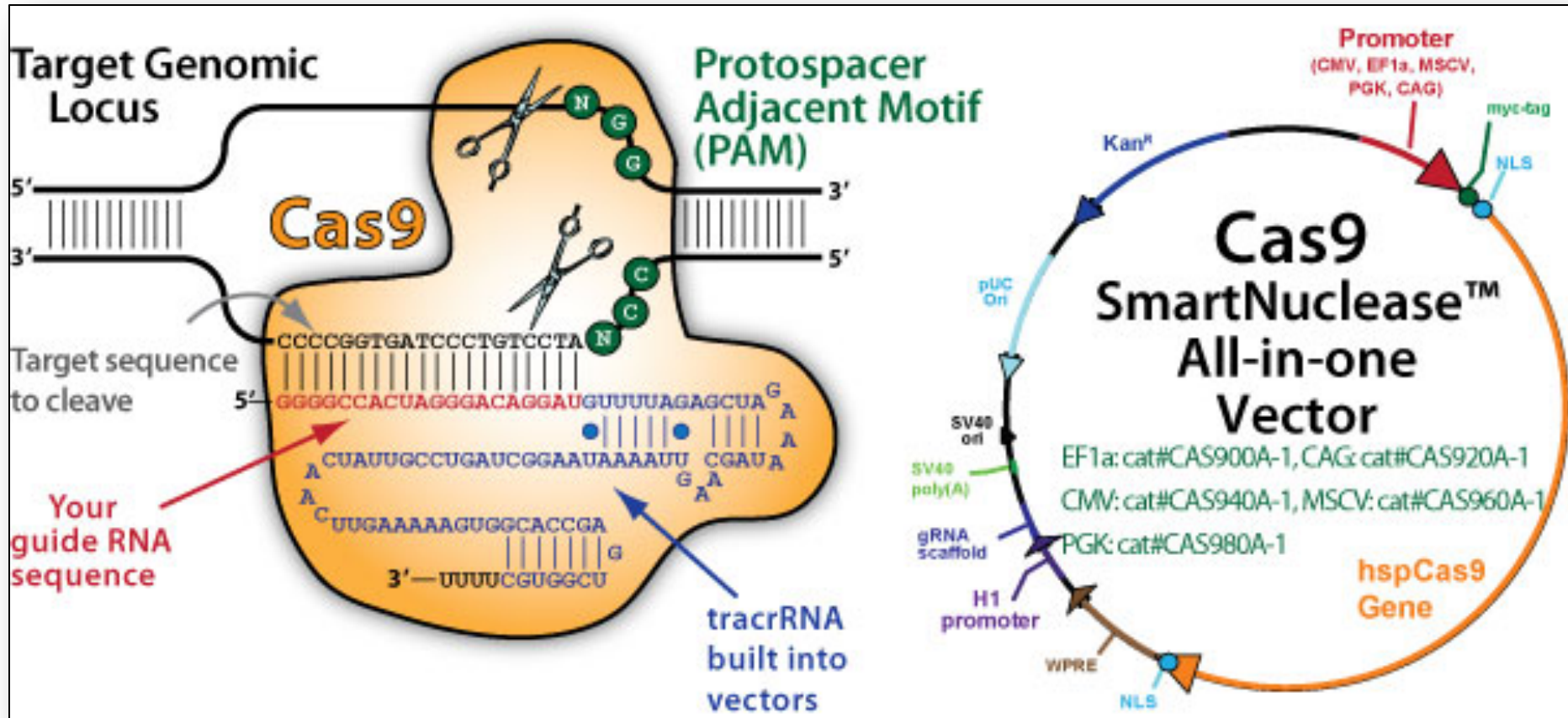
BREAKTHROUGH – fuse the crRNA and tracrRNA to make a single “Guide RNA”



CRISPR/Cas9 can help **make** a mutation or **fix** a mutation NHEJ vs Homologous Recombination **IN HUMAN CELLS**



Genome Editing Kits Galore



<https://www.youtube.com/watch?v=2pp17E4E-O8>

Awards Galore



Emmanuelle Charpentier and Jennifer Doudna

Awards Galore

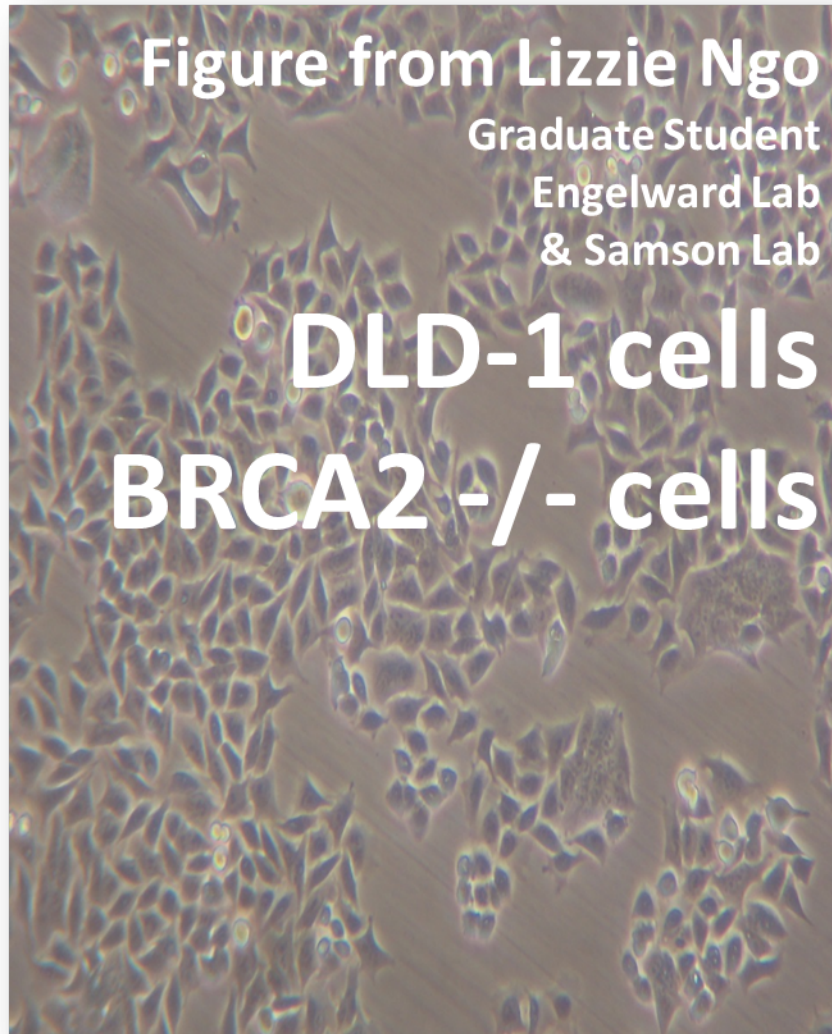
2016 Canada Gairdner Awards



LES PRIX CANADA GAIRDNER AWARDS

Feng Zhang - MIT

BOTH *BRCA2* alleles have exon 11 replaced with a selectable marker gene (for Neomycin resistance)



It is **essential** to confirm the identity of the cells you are working with

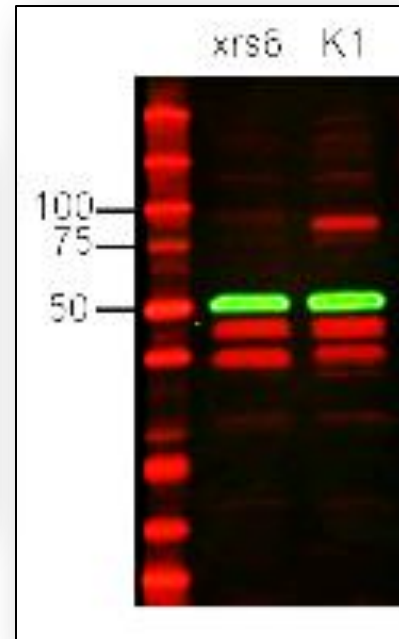
Western Blot Analysis



α - tubulin

Ku80

(Or in your case, BRCA2)



Key Experimental Methods for Module 2

- Grow human cancer cells in tissue cell culture
- Monitor specific protein levels by Western blot
- Kill cancer cells with chemotherapy drugs
- Engineer the inhibition of DNA Repair pathways
- Monitor changes in a gene's expression (qPCR)
- Analyze RNAseq dataset measuring expression of ~ 20,000 genes (BIG DATA!)
- Statistical analysis of all biological data

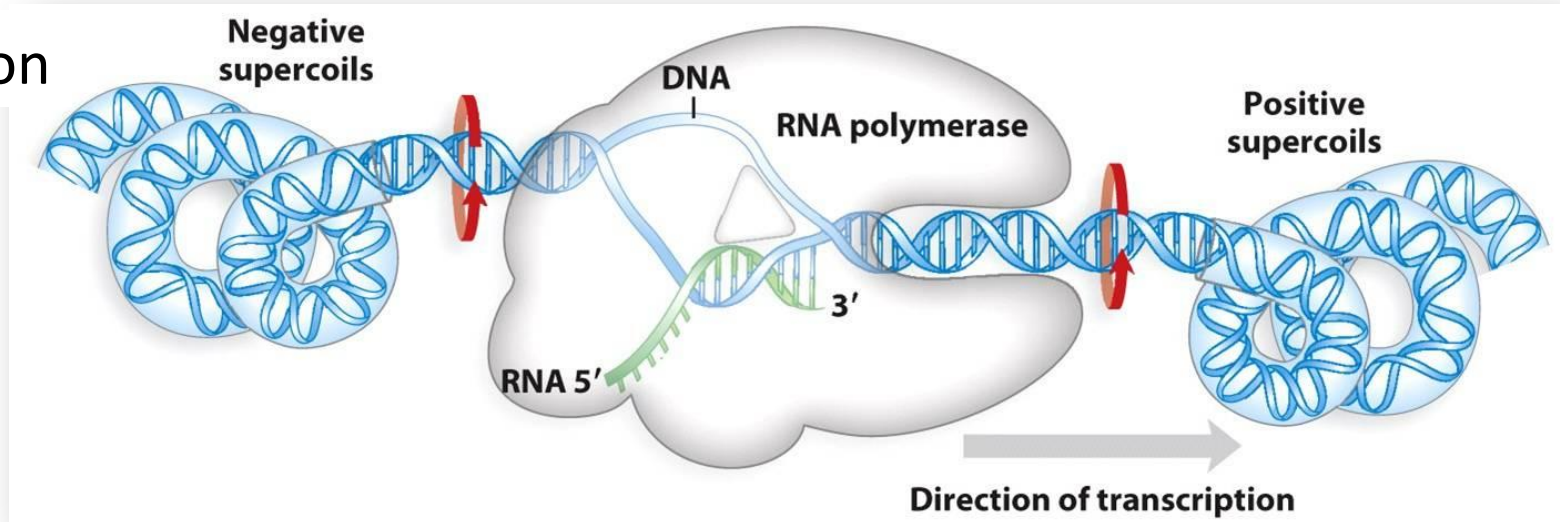
Some cancer Chemotherapy agents and all Radiotherapies
CAUSE DNA DAMAGE



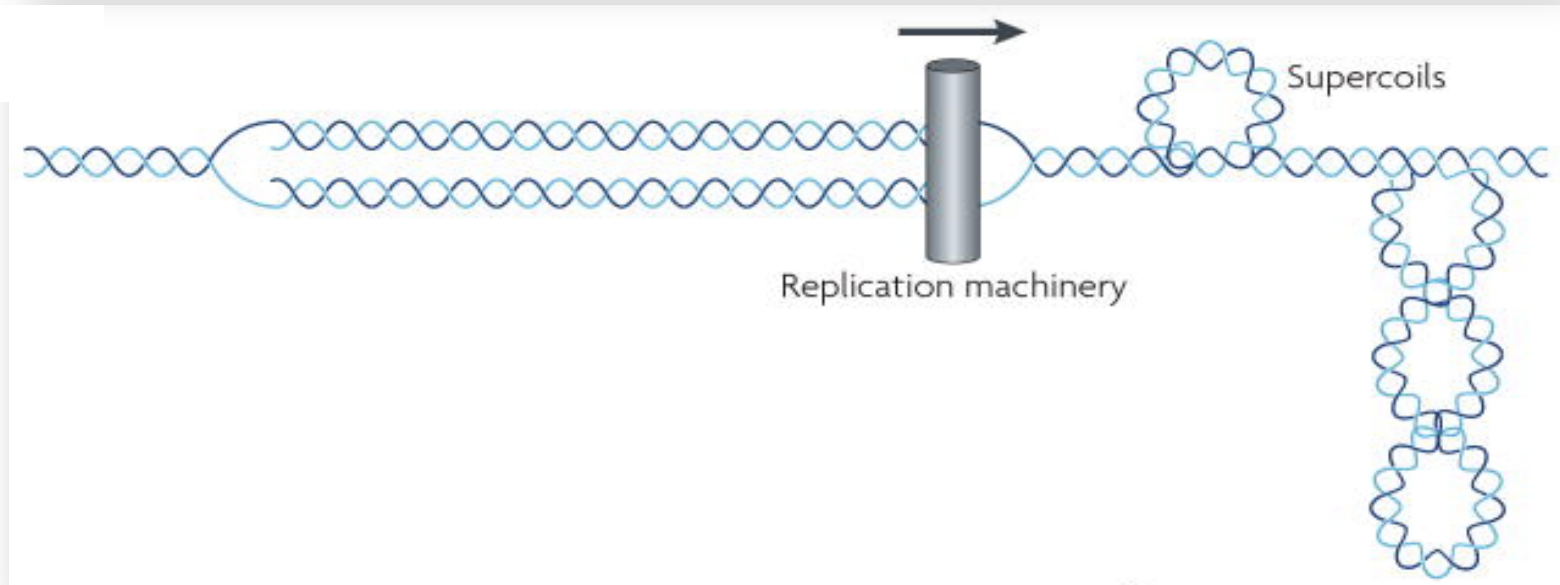
Etoposide induces **DNA double strand breaks** (DSBs) by inhibiting an enzyme that undoes the supercoiling of DNA

DNA Replication and RNA Transcription cause DNA SUPERCOILING

Transcription

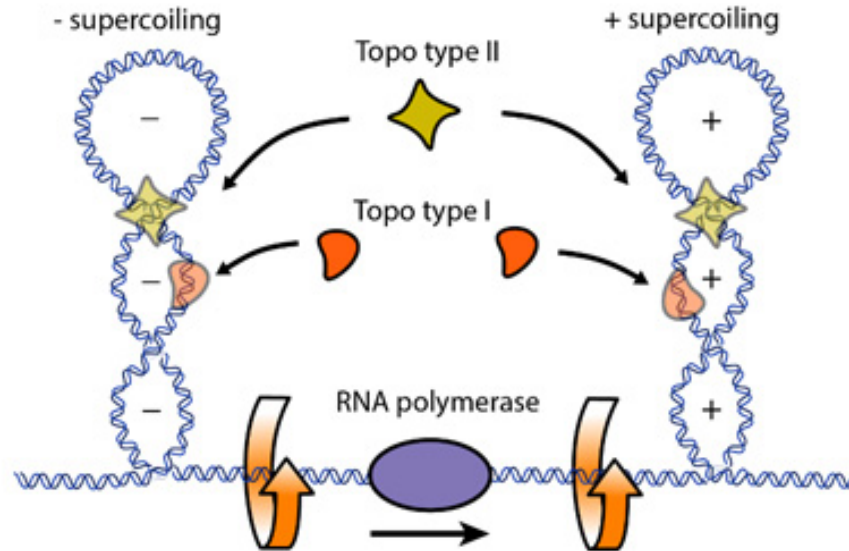


Replication

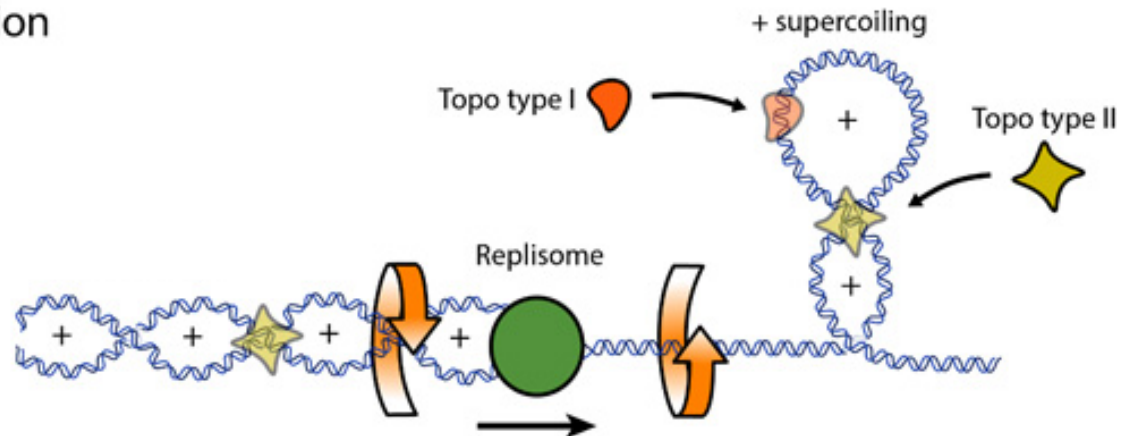


RNA Transcription and DNA Replication cause DNA SUPERCOILING

A Transcription



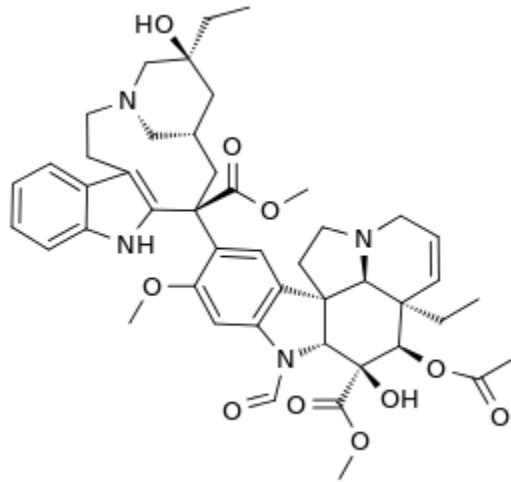
B Replication



Topoisomerase I and II

<https://www.youtube.com/watch?v=EYGrElVyHnU>

Etoposide – Inhibits Topoisomerase II

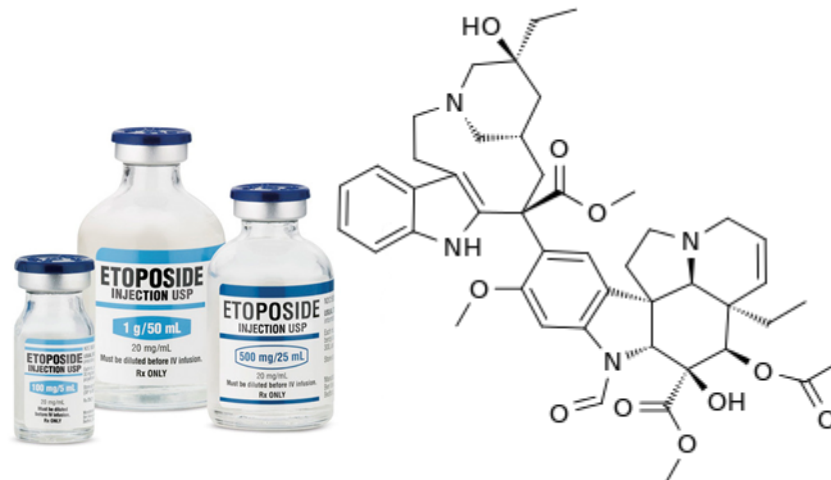
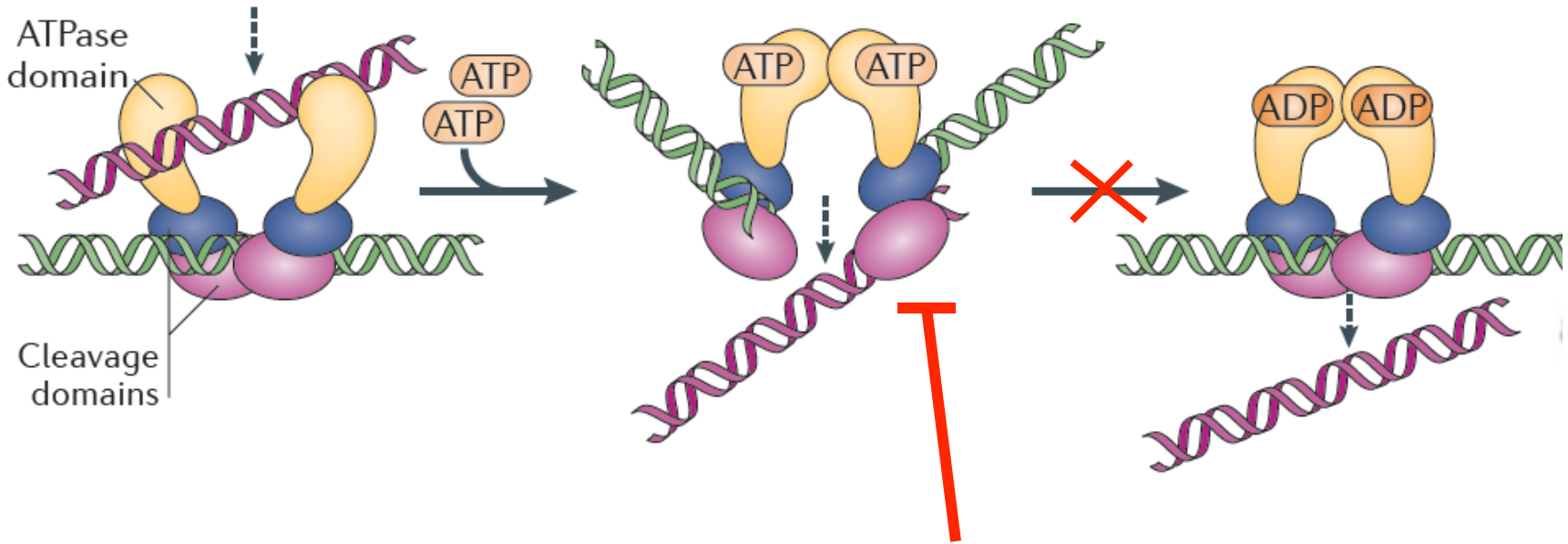


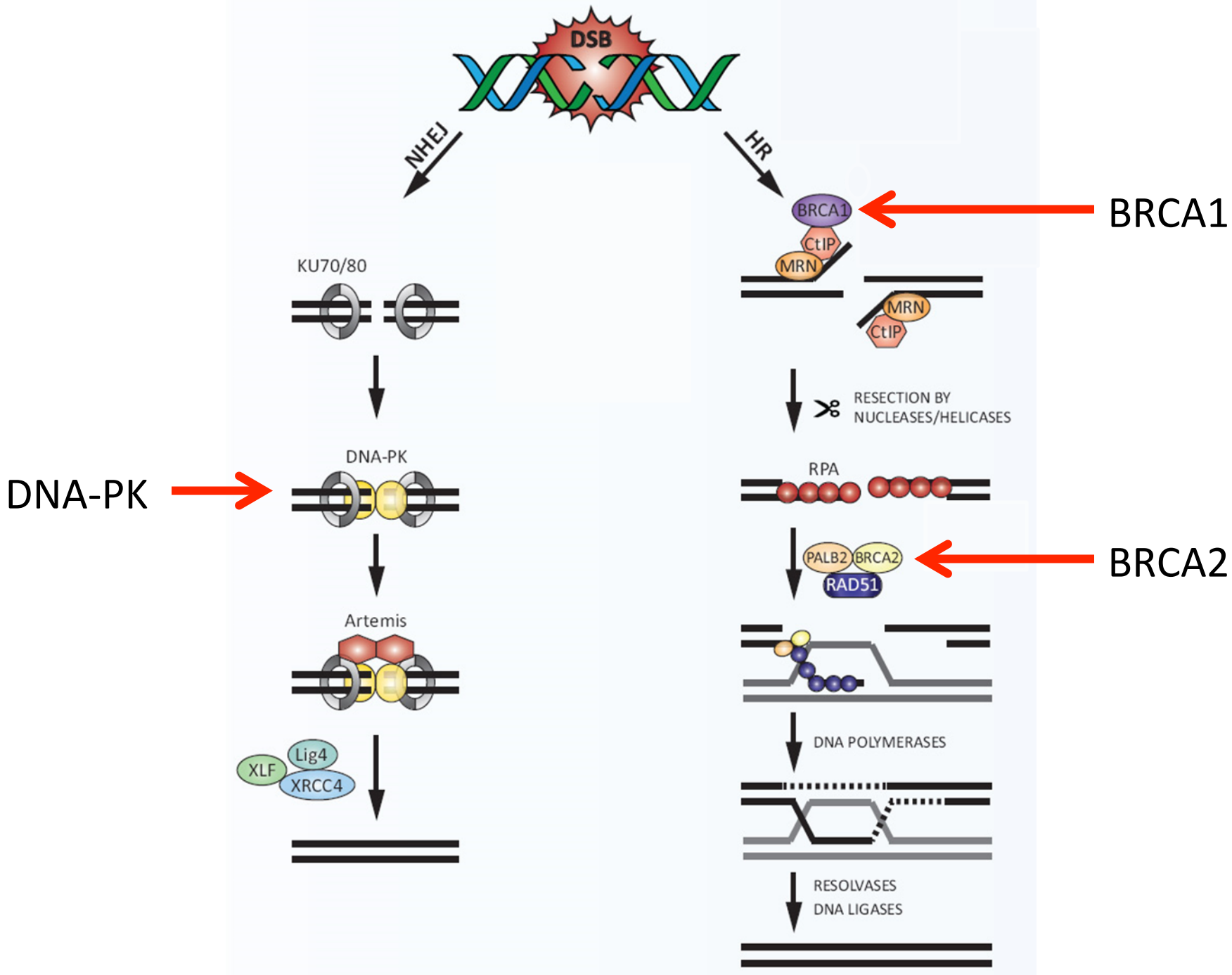
Etoposide is a semisynthetic derivative of PODOPHYLLOTOXIN from the rhizome of the wild mandrake (*Podophyllum peltatum*). Used for **Cancer Chemotherapy**



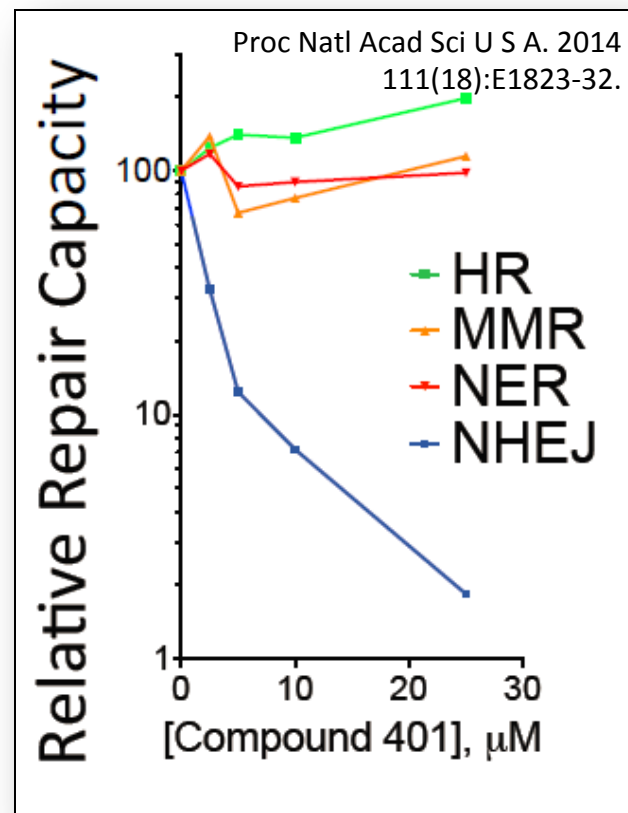
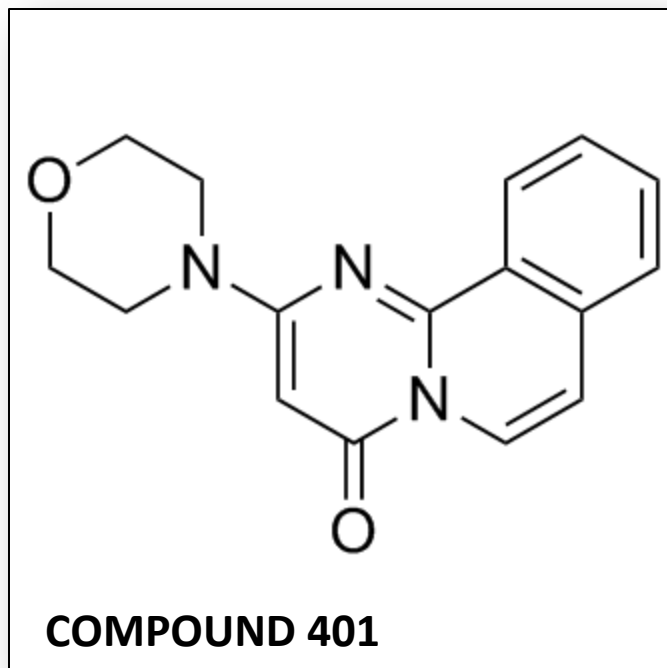
The wild Mandrake, with Rhizome at the bottom (at the bottom)

Etoposide – Inhibits Topoisomerase II





COMPOUND 401 – “Selective” Inhibitor of DNA-PK and NHEJ

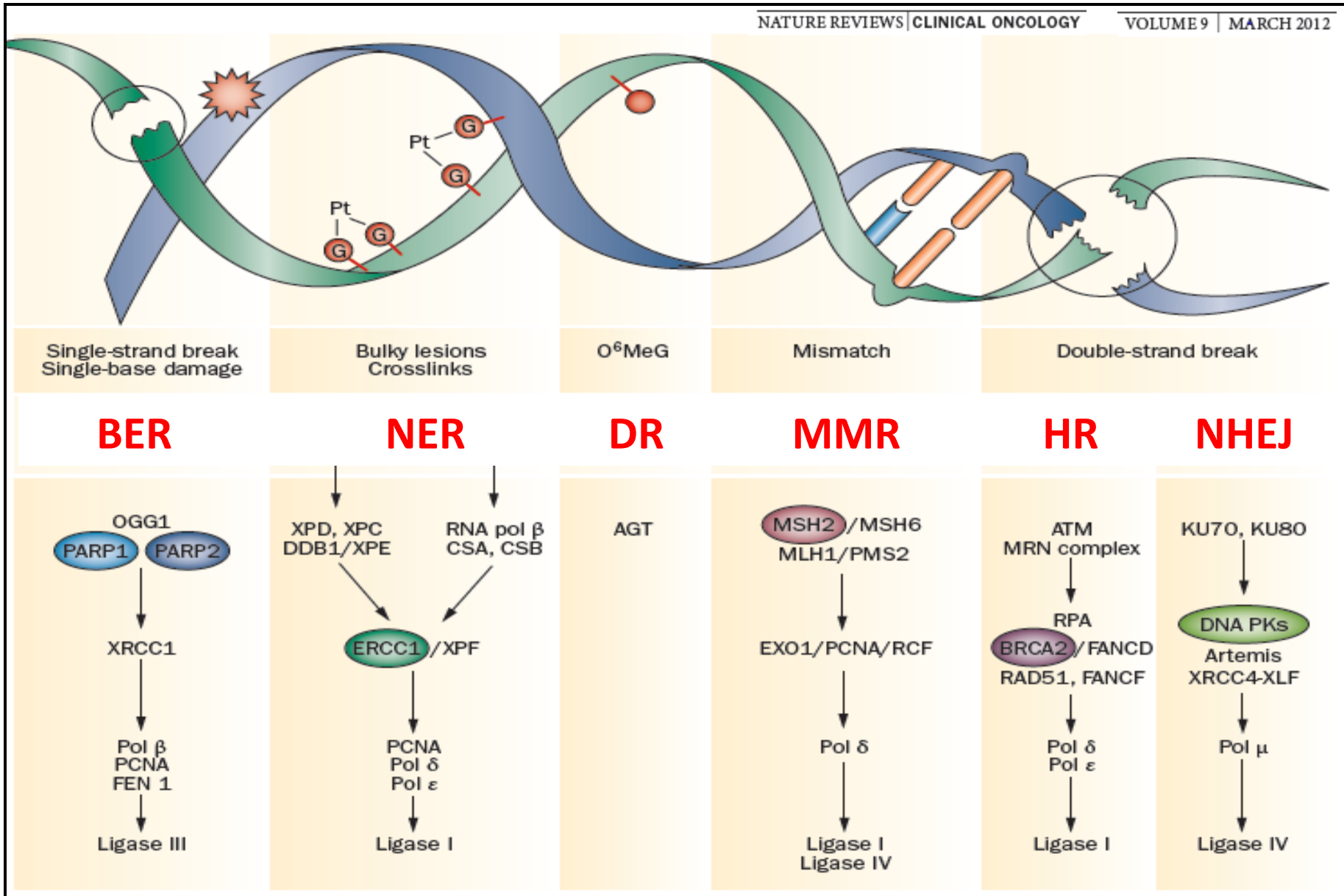


Biological Activity of COMPOUND 104

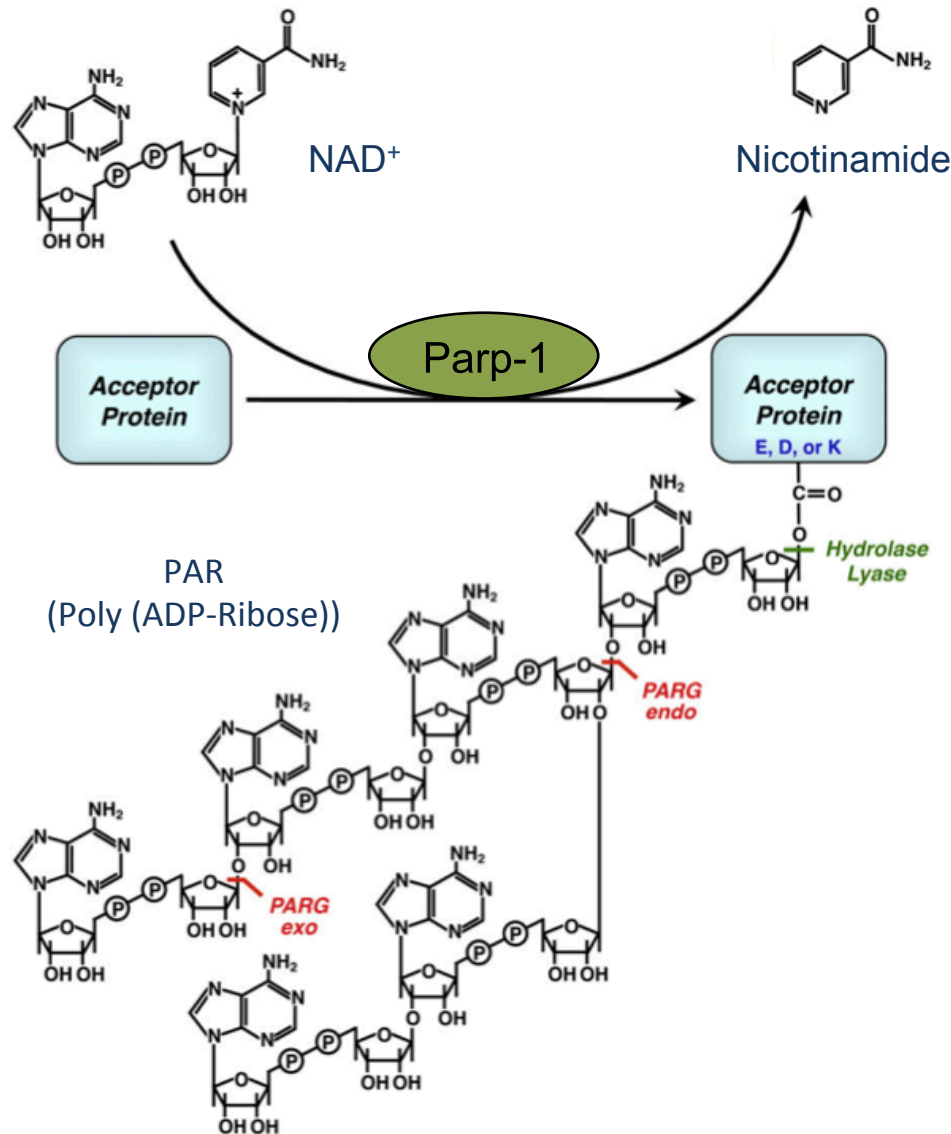
Reversible and selective inhibitor of DNA-dependent protein kinase (DNA-PK) and mammalian target of rapamycin (mTOR) (IC_{50} values are 0.28 and 5.3 μ M respectively). Displays little affinity for other commonly studied kinases including PI 3-K, ATM and ATR (IC_{50} values are all > 100 μ M).

Six Major DNA Repair Pathways

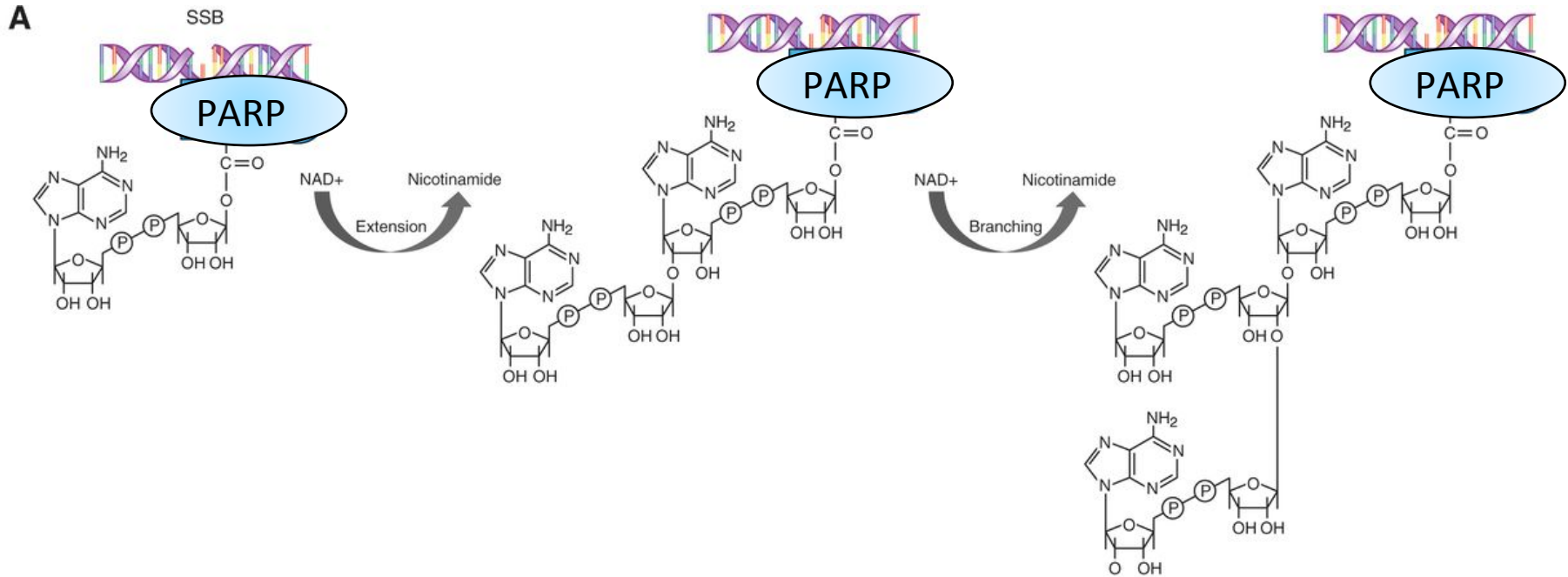
NATURE REVIEWS | CLINICAL ONCOLOGY | VOLUME 9 | MARCH 2012



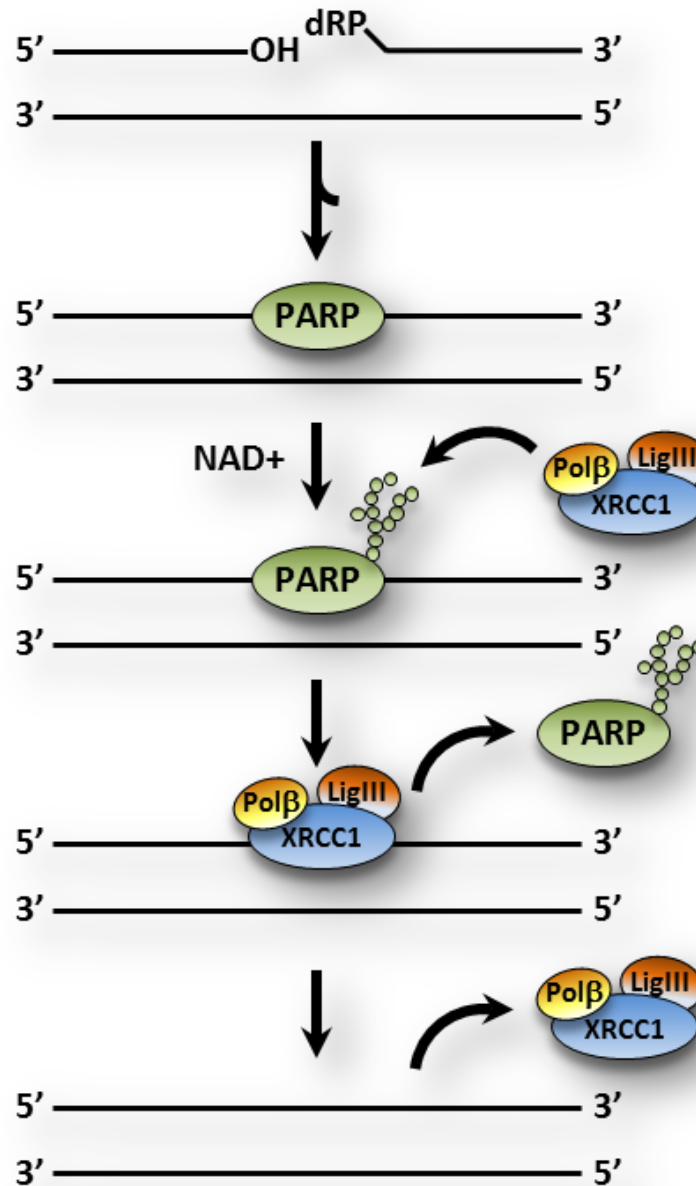
Poly (ADP-Ribose) Polymerase - PARP



Poly (ADP-Ribose) Polymerase – PARP is activated at DNA Single Strand Breaks (SSBs)



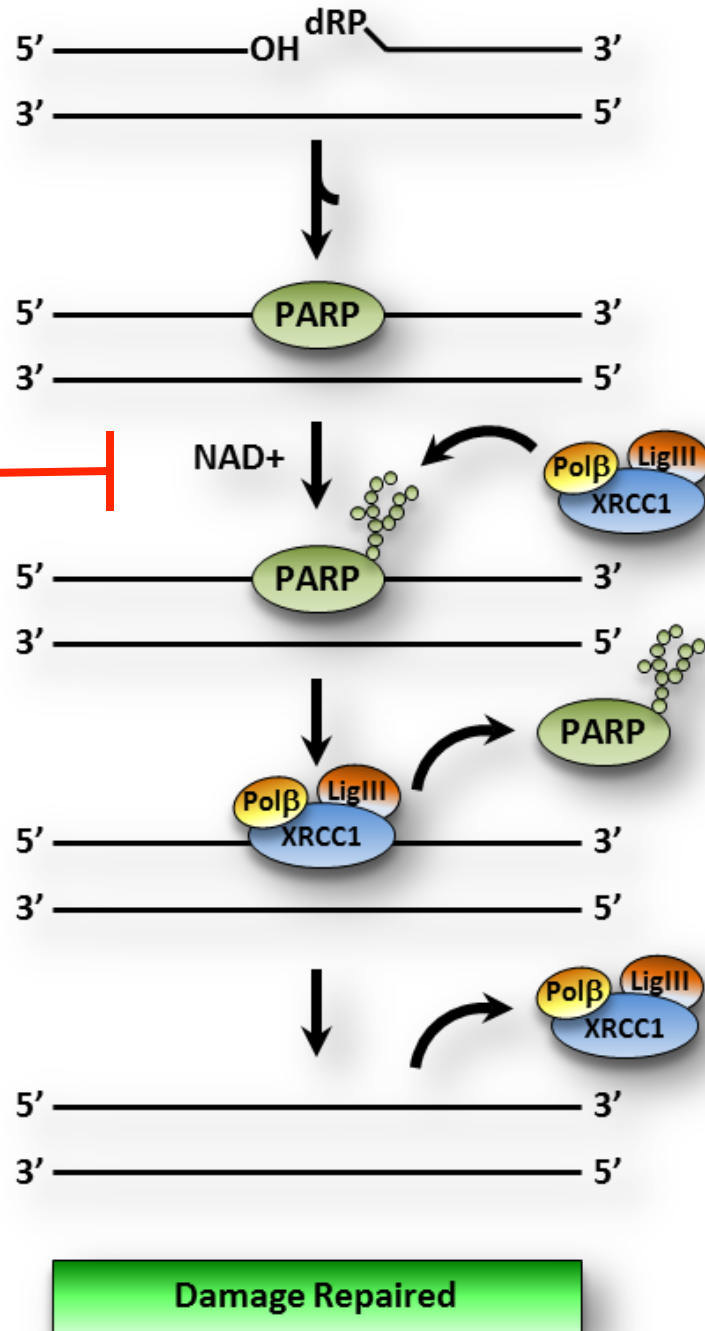
DNA Single Strand Break (SSB) Repair



Damage Repaired

DNA Single Strand Break (SSB) Repair

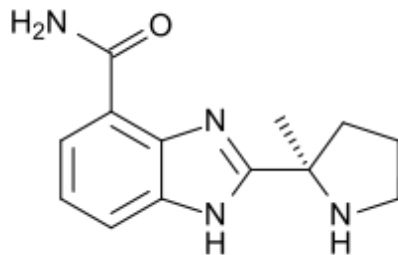
Inhibit this step



Parp Inhibitors in Clinical Trials

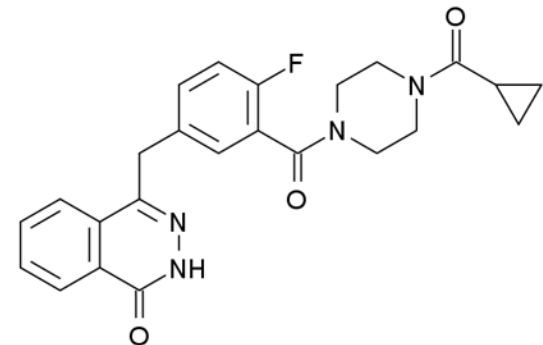
Veliparib

- ABT-888 (AbbVie)
- Competes for NAD⁺ binding site
- Less DNA trapping

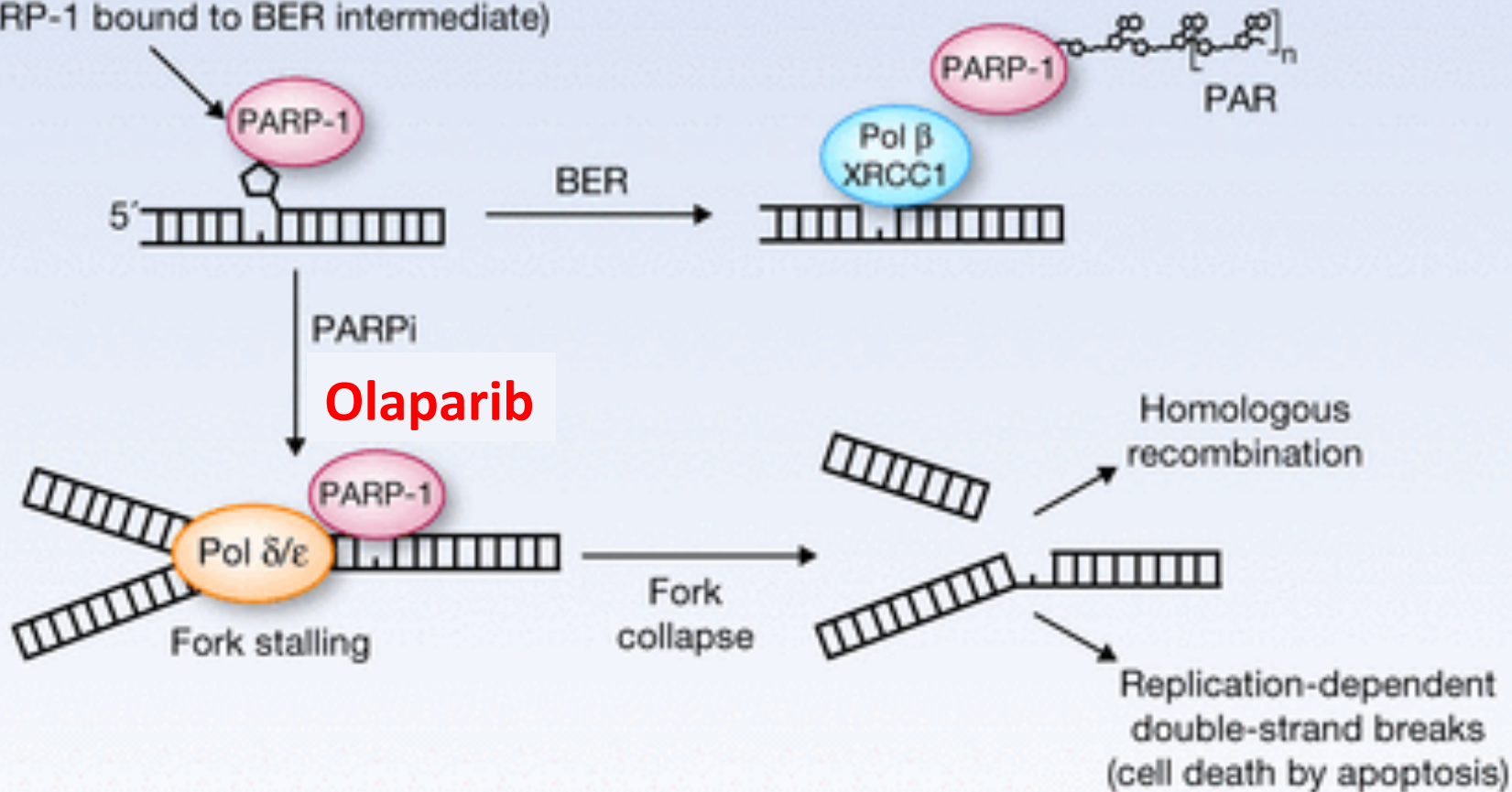


Olaparib

- AZD-2281 (Astra Zeneca)
- Competes for NAD⁺ binding site
- More DNA trapping



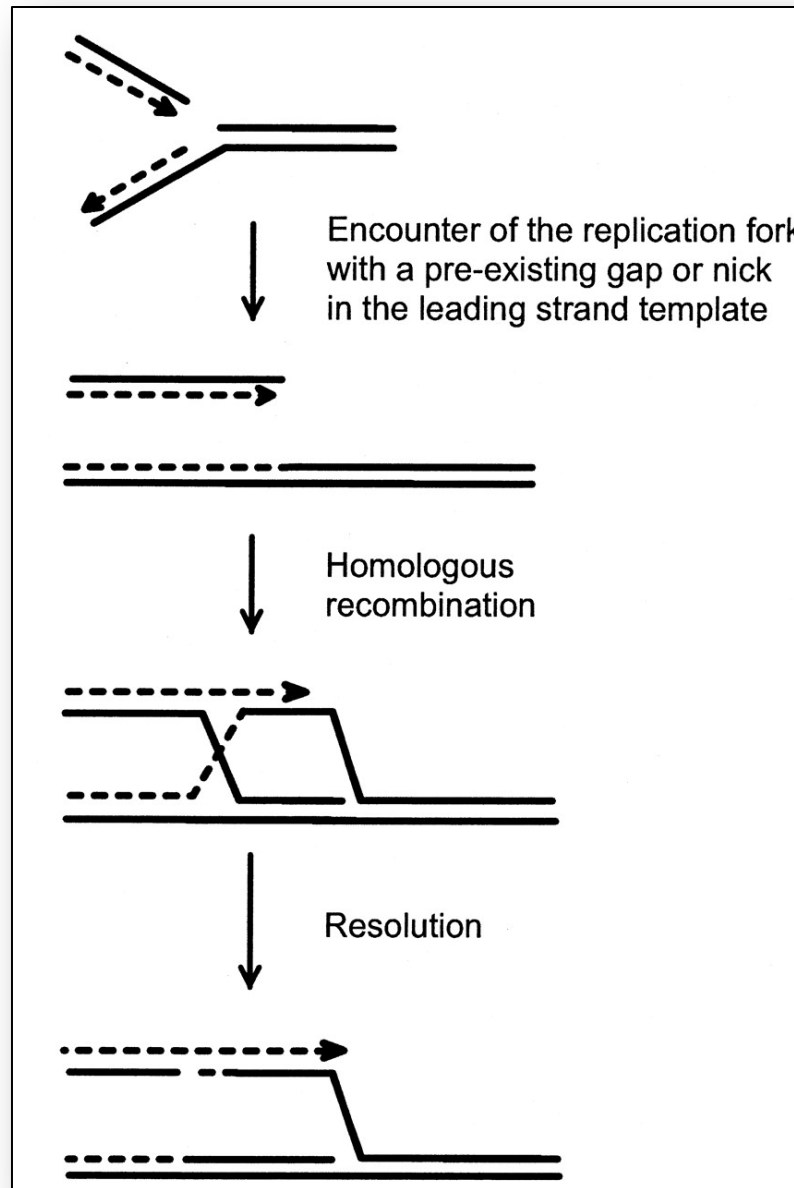
Endogenous DNA Damage
(PARP-1 bound to BER intermediate)



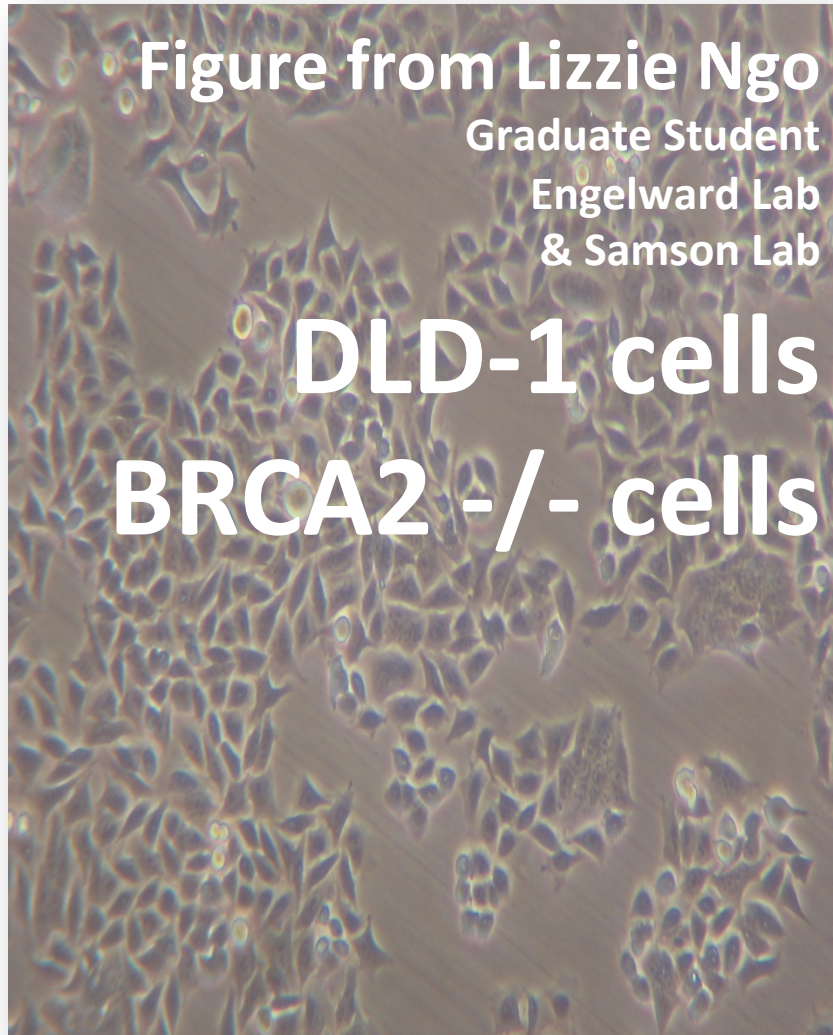
© 2014 American Association for Cancer Research

Homologous Recombination can Rescue collapsed Replication Forks – what happens in BRCA2-/- cells?

Collapsed replication fork



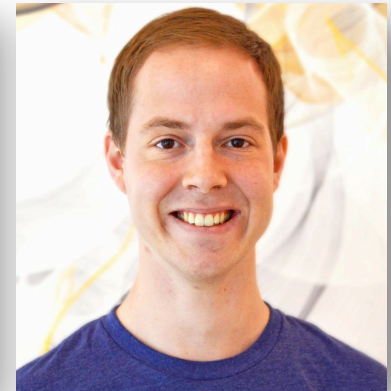
Here's how you will treat your cells today



- **Etoposide** to inhibit TopoII leading to DNA DSBs
- **Compound 401** to inhibit Non Homologous End Joining (NHEJ)
- **Etoposide + Compound 401**
- **Olaparib** to inhibit PARP1 to stabilize DNA SSBs leading to DNA DSBs at collapsed replication forks

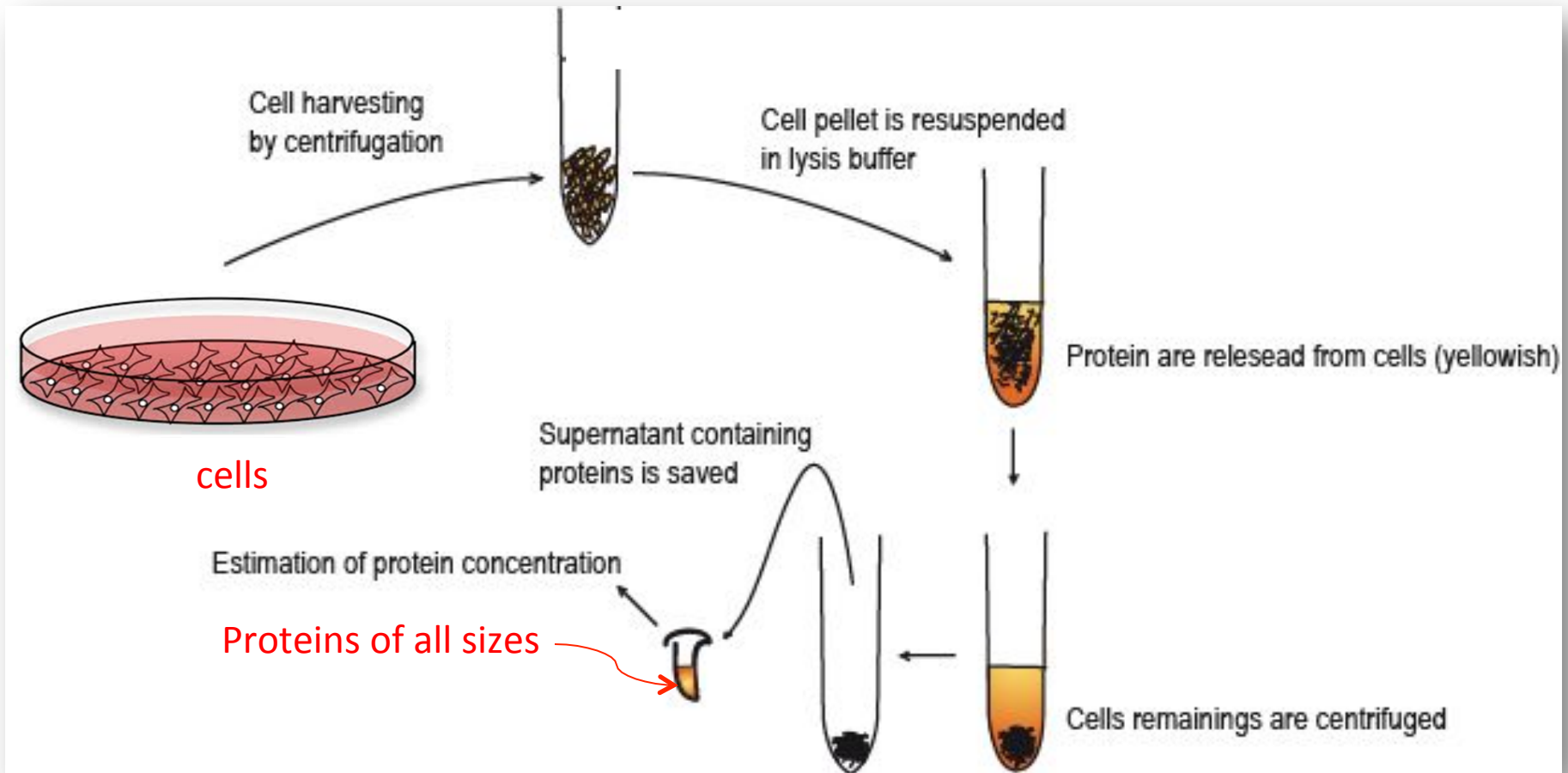
20.109 Spring 2017 Module 2 – Lecture 2

Gene Expression Engineering (March 14th 2017)

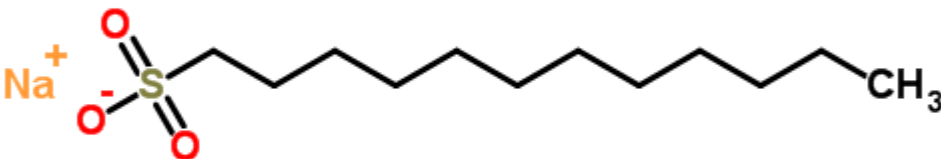


Noreen Lyell
Leslie McLain
Maxine Jonas
Rob Wilson
Leona Samson (Lectures)

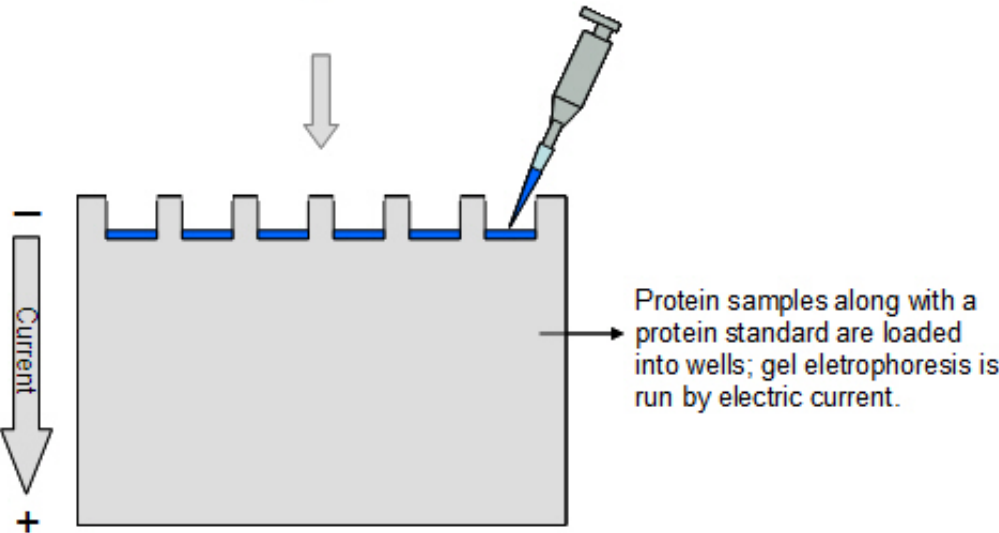
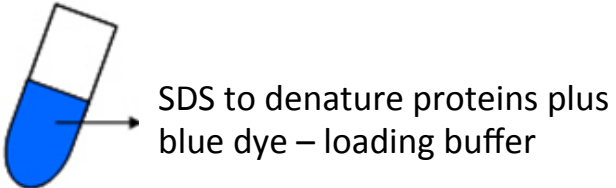
Prepare Cell Extract for Western Blot (Immunoblot) Analysis



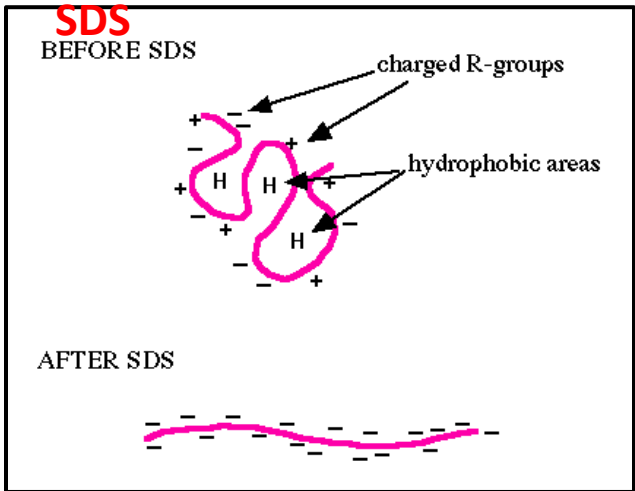
Western Blot or Immunoblot Analysis



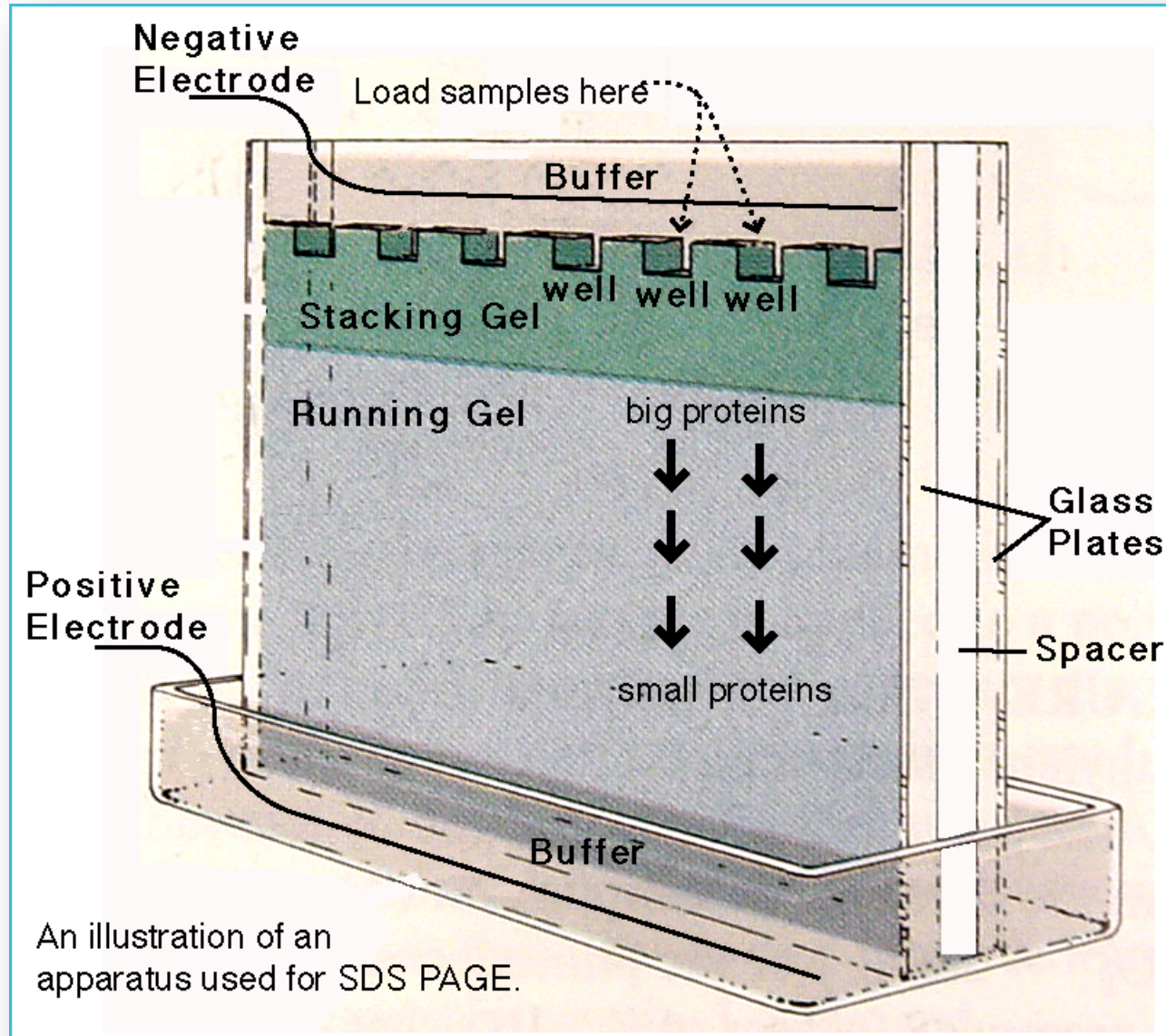
SDS – sodium dodecyl sulfate

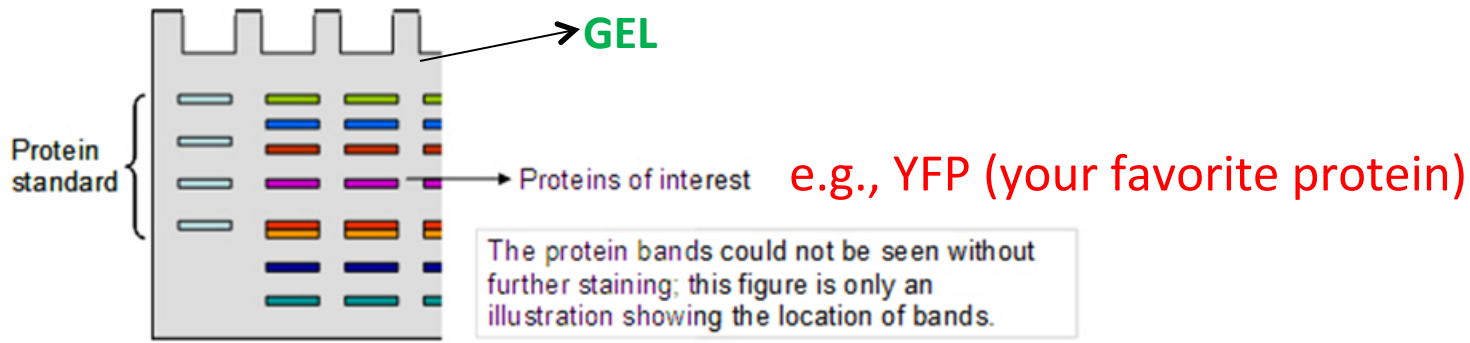


Proteins denatured by SDS

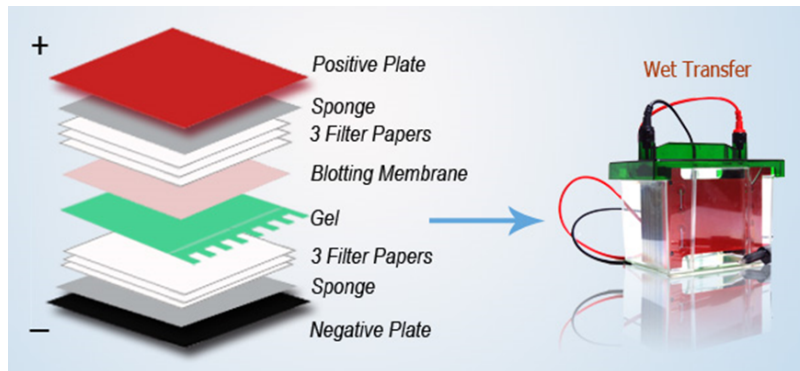


Separating proteins on an SDS polyacrylamide gel

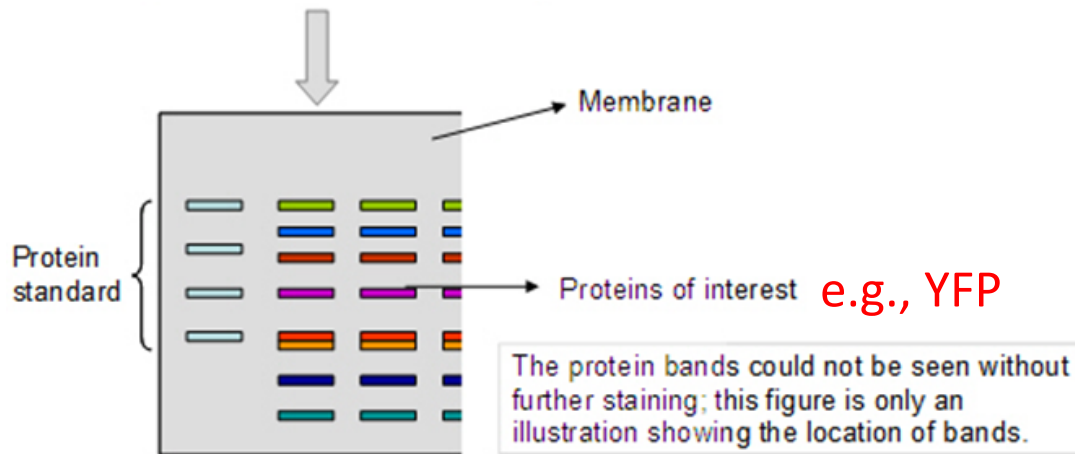


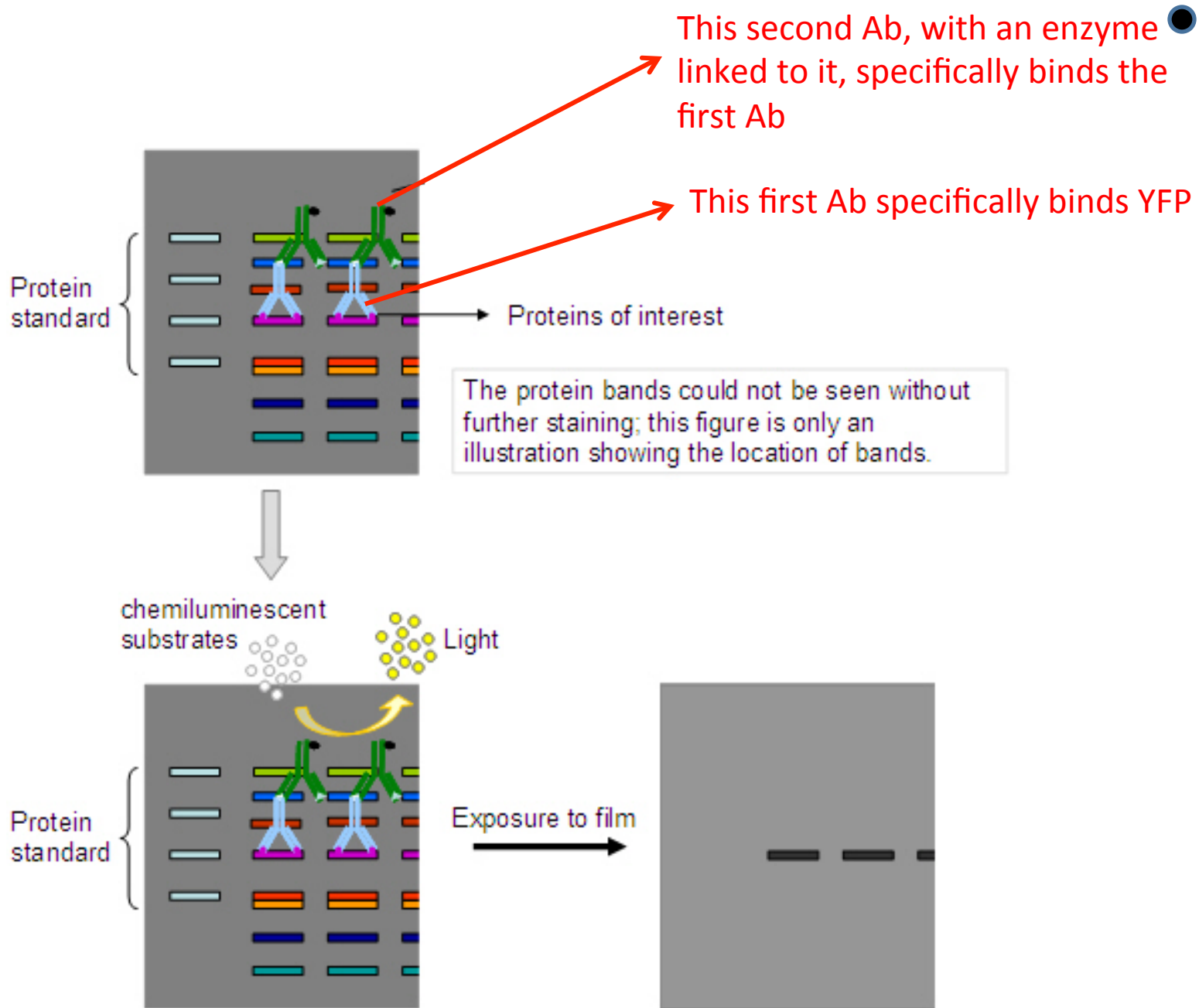


Transfer (Semi-dry transfer as an example)



The Separated proteins are transferred out of the gel and **BLOTTED** onto a nitrocellulose or other membrane





Western Blot Analysis

