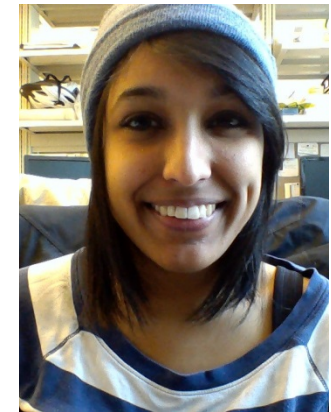


20.109 Spring 2014 Mod 2 – Lecture 5

System Engineering and Protein Foundations



Agi Stachowiak

Shannon Hughes

Aneesh Ramaswamy

Suhani Vora (TA)

Leona Samson (Lectures)

Zachary Nagel (help with development)



What experimental question will you ask in Module 2?

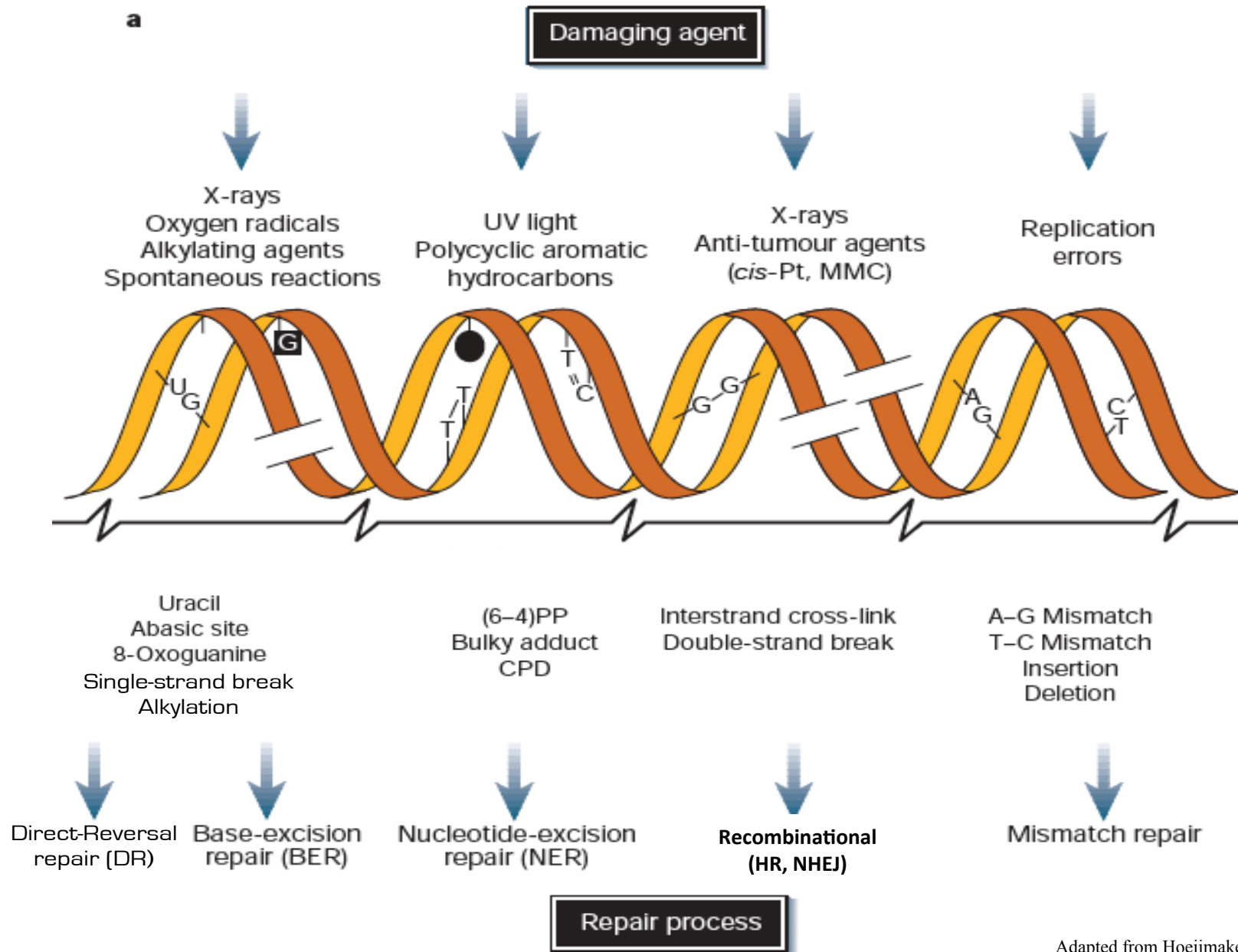
How efficiently does DNA repair by the Non Homologous End Joining (NHEJ) pathway act on DNA damage with different topologies?



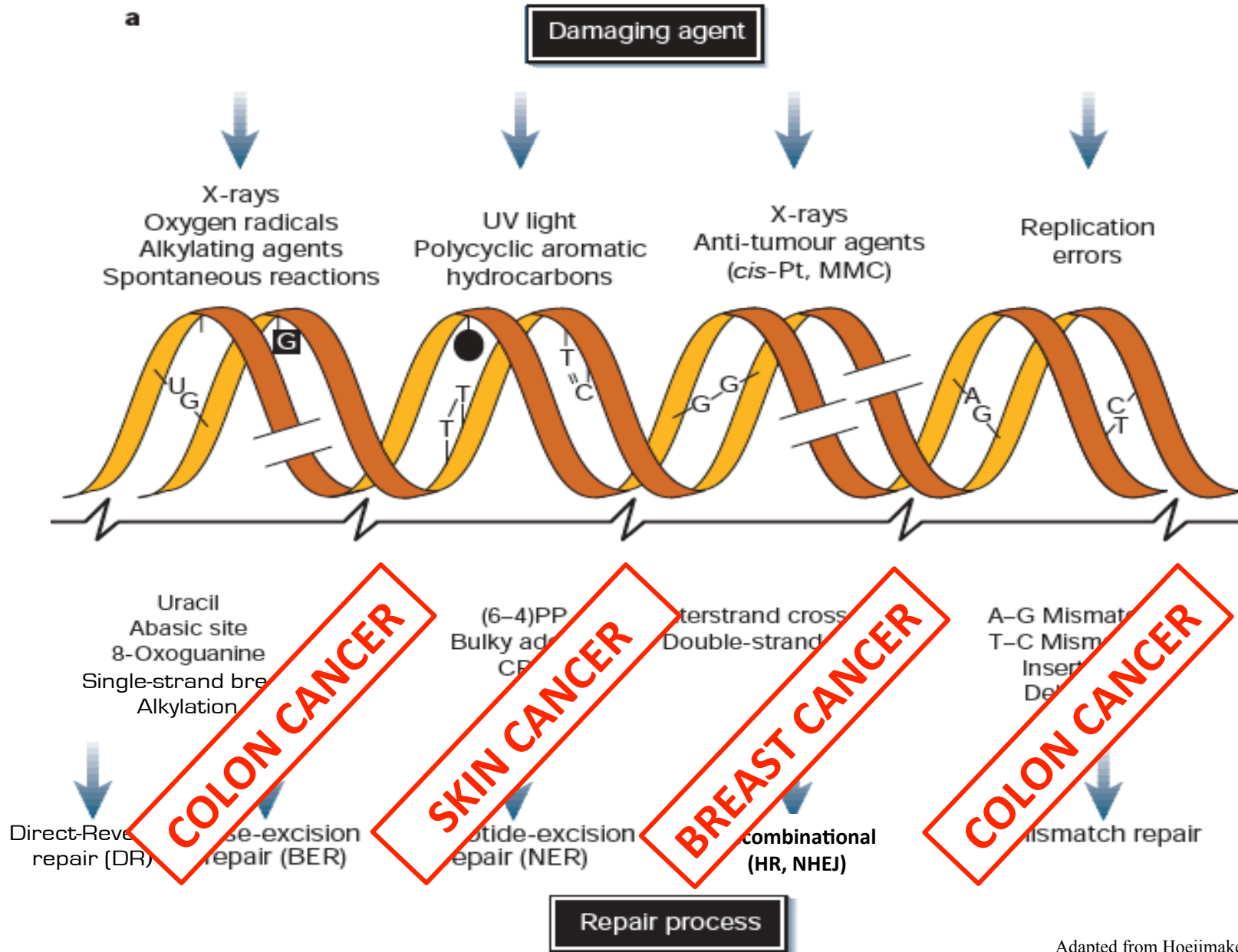
This raises the following questions

- How does DNA get damaged?
- What is DNA repair?
- Why does DNA repair exist?
- Why do we care about how efficient DNA repair is?
- How does one actually measure DNA repair efficiency?

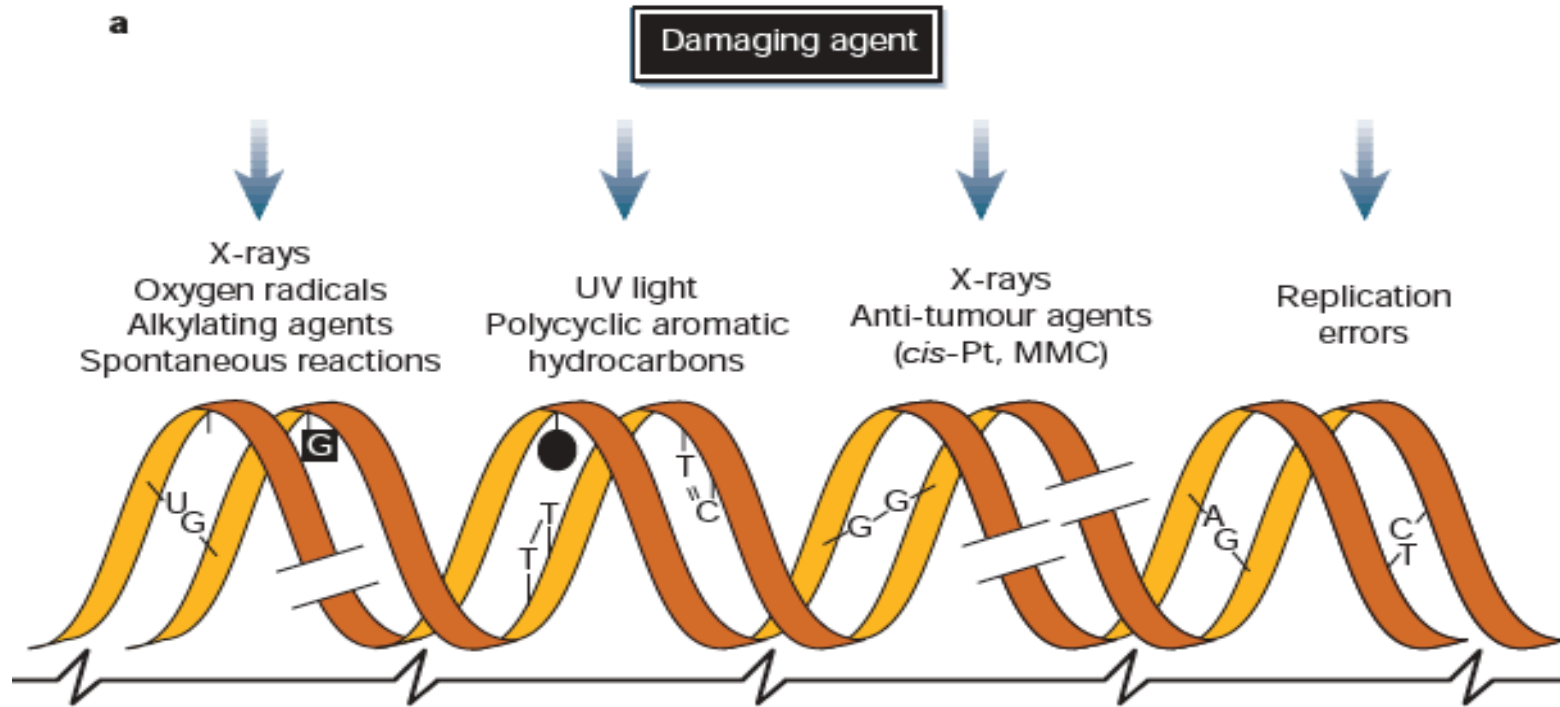
DNA Damage and Repair



DNA Damage and Repair



DNA Damage and Repair



RESPONSES of TUMOR and NON-TUMOR CELLS to CANCER RADIOTHERAPY and CHEMOTHERAPY



Key Experimental Methods for Module 2

- Mammalian tissue cell culture
- Monitoring protein level by Western blot
- Generating plasmids with DNA damage
- Transfecting plasmids into mammalian cells
- Using fluorescent proteins as reporters of biological processes
- Flow cytometry to measure DNA repair
- Statistical analysis of biological data

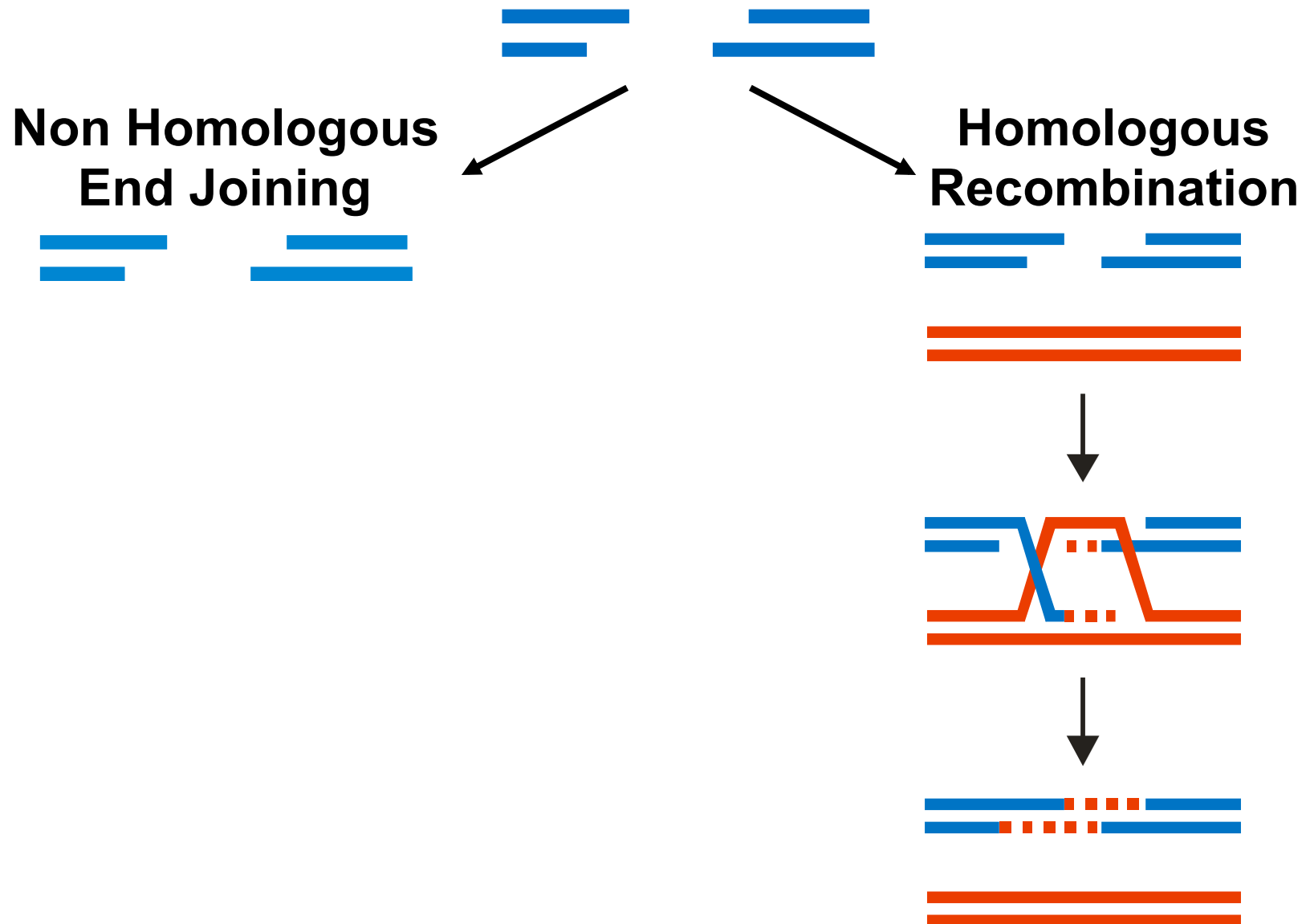


Key Experimental Methods for Module 2

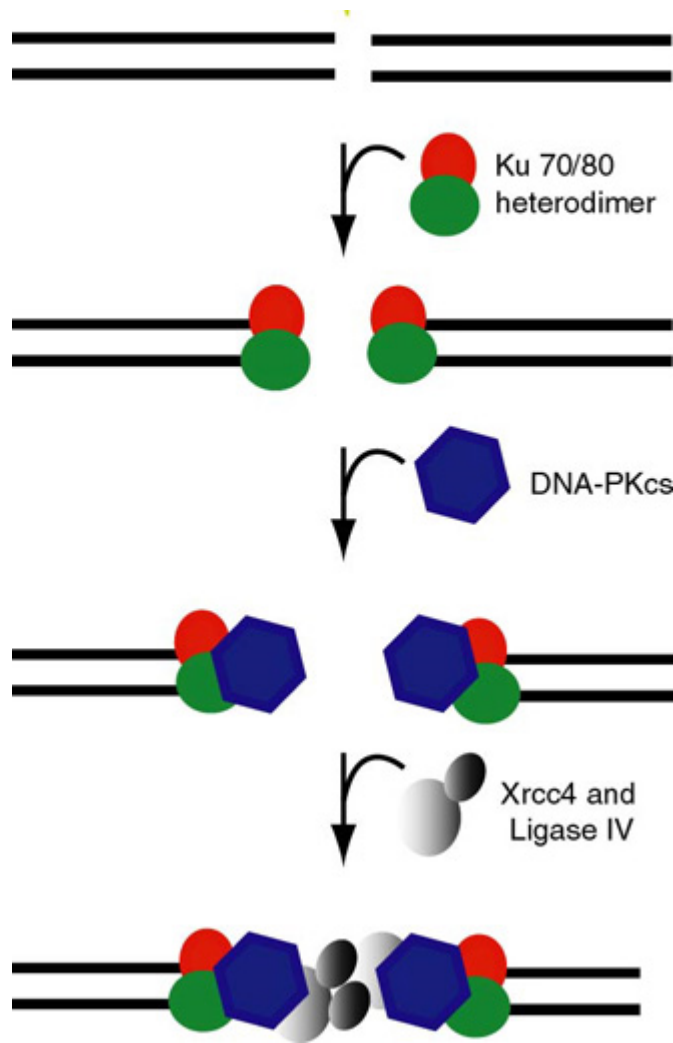


- Mammalian tissue cell culture
- Monitoring protein level by Western blot
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- **Transfecting plasmids into mammalian cells**
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- **Flow cytometry to measure DNA repair**
- Statistical analysis of biological data

DNA double-strand break repair



Non-Homologous End Joining (NHEJ)



Ku70

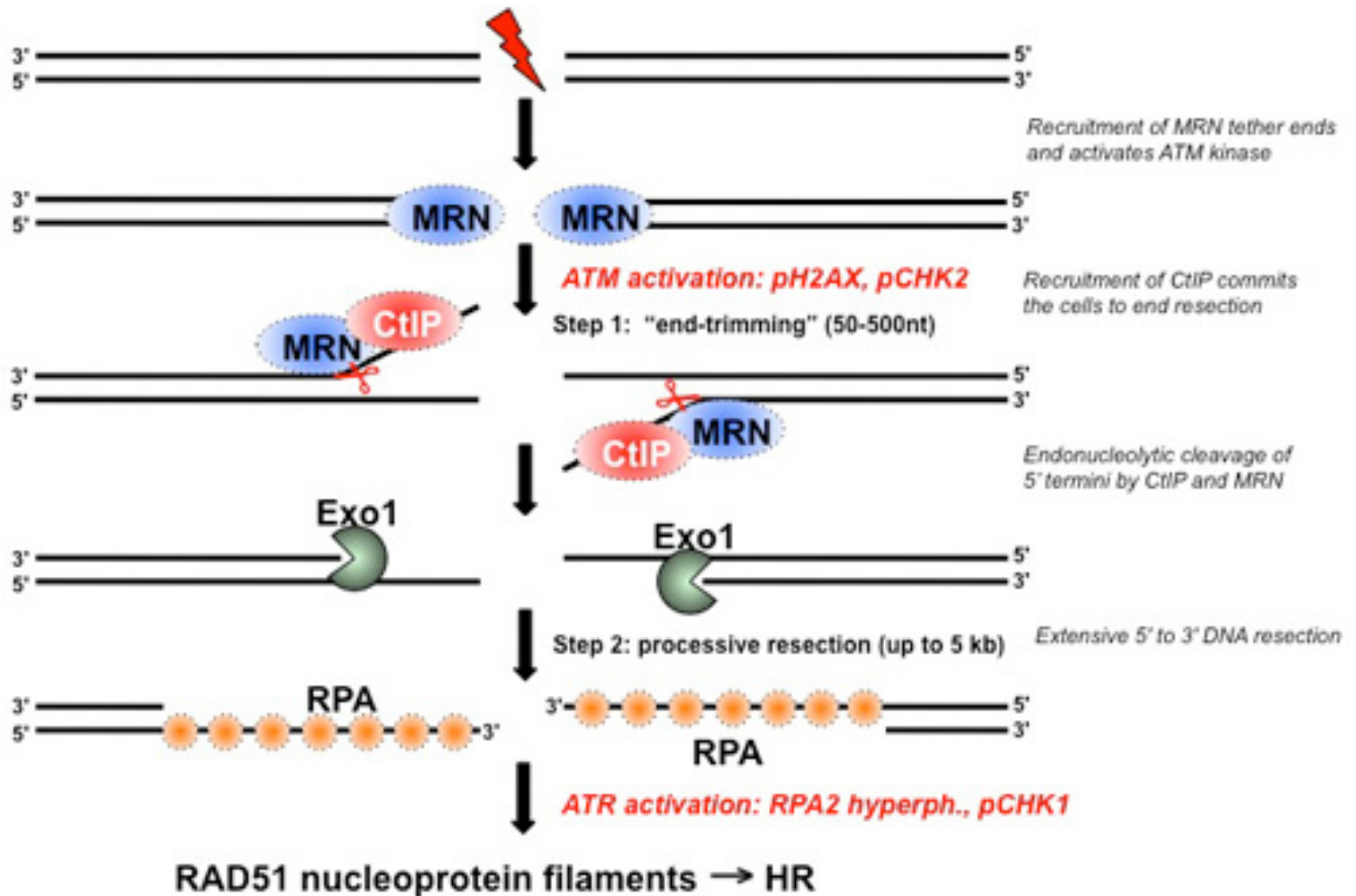
Ku80

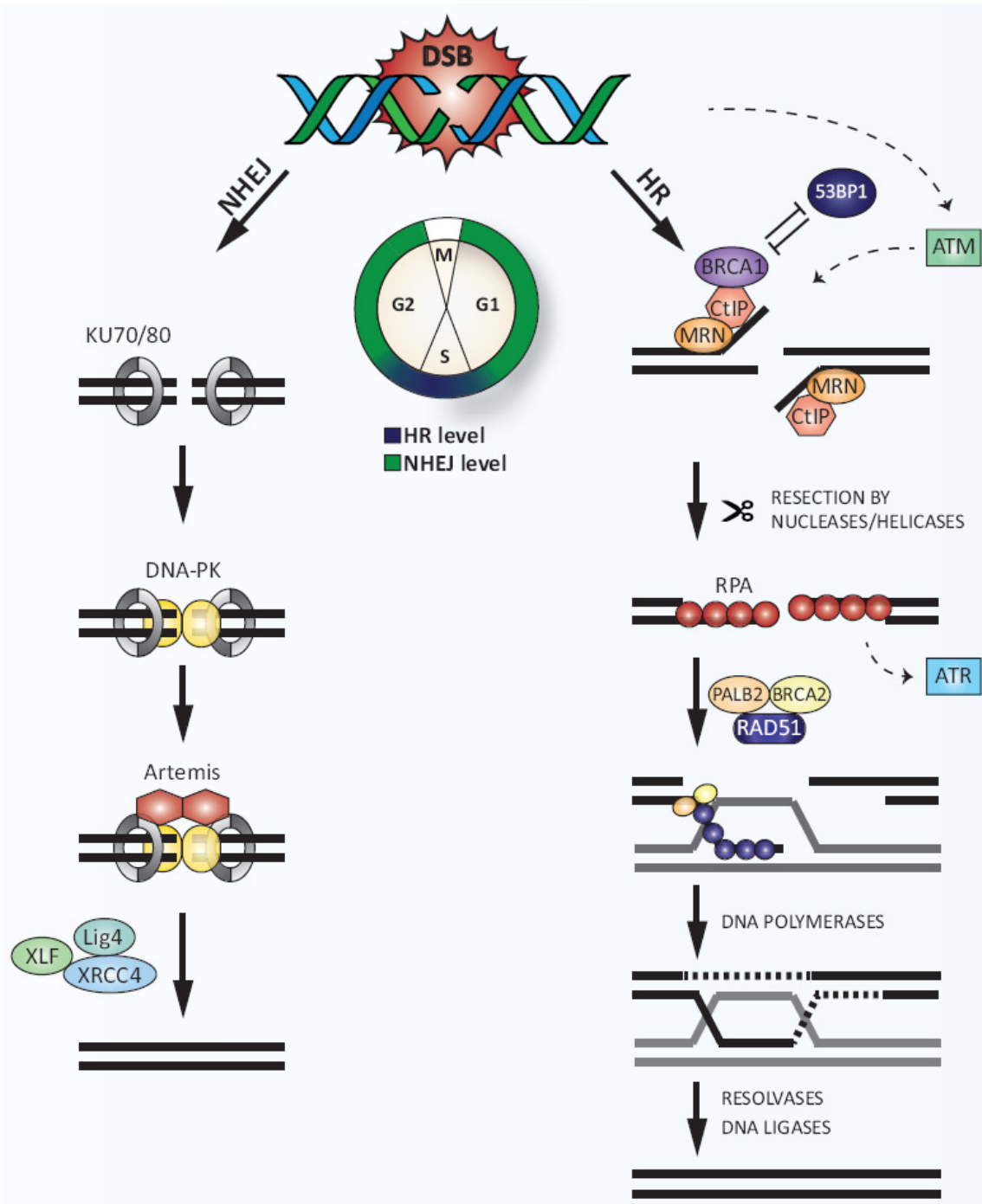
DNA-PKcs

Xrcc4

Ligase IV

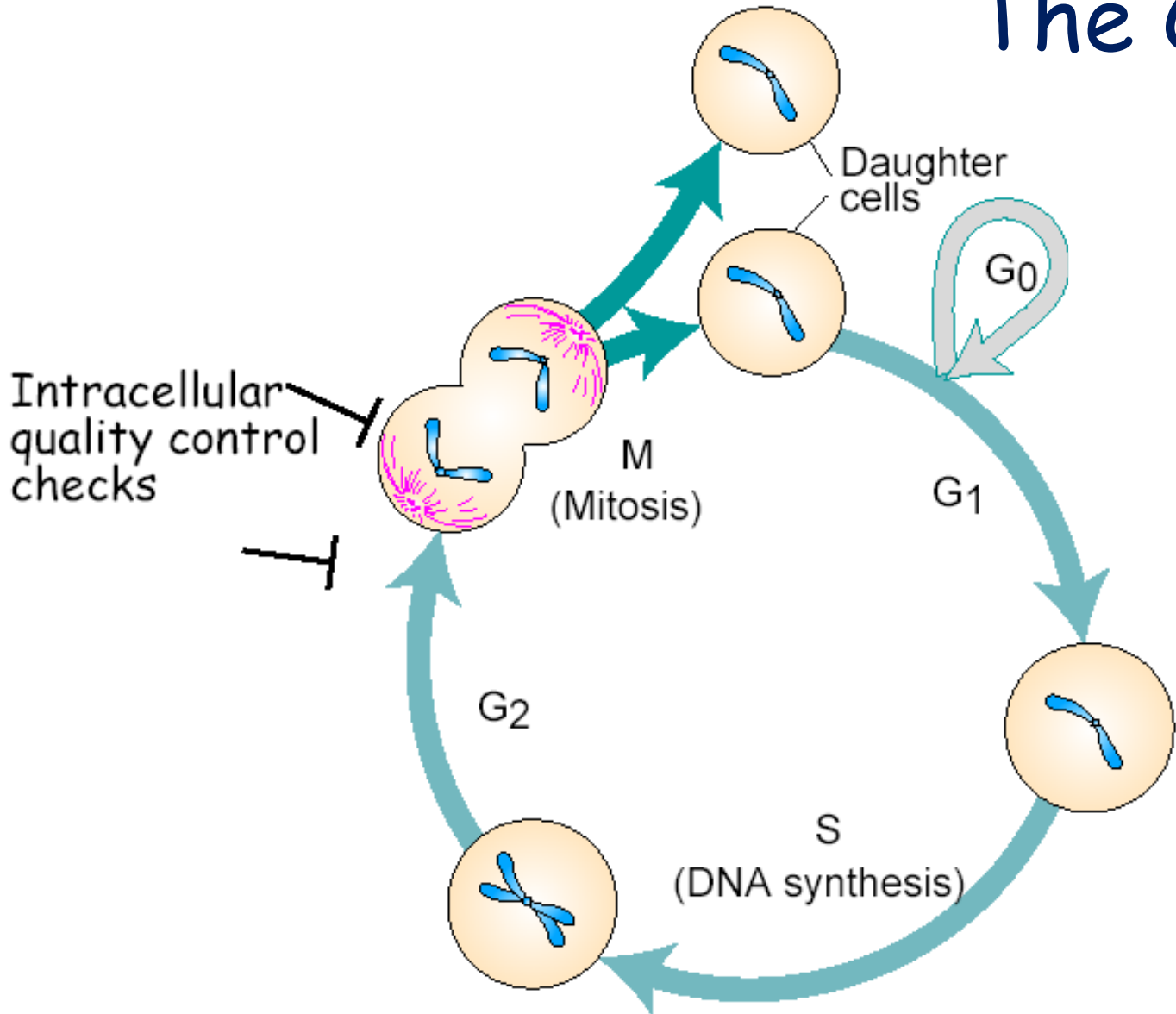
The MRN complex (Mre11/Rad50/Nbs1) competes with Ku for binding DSBs





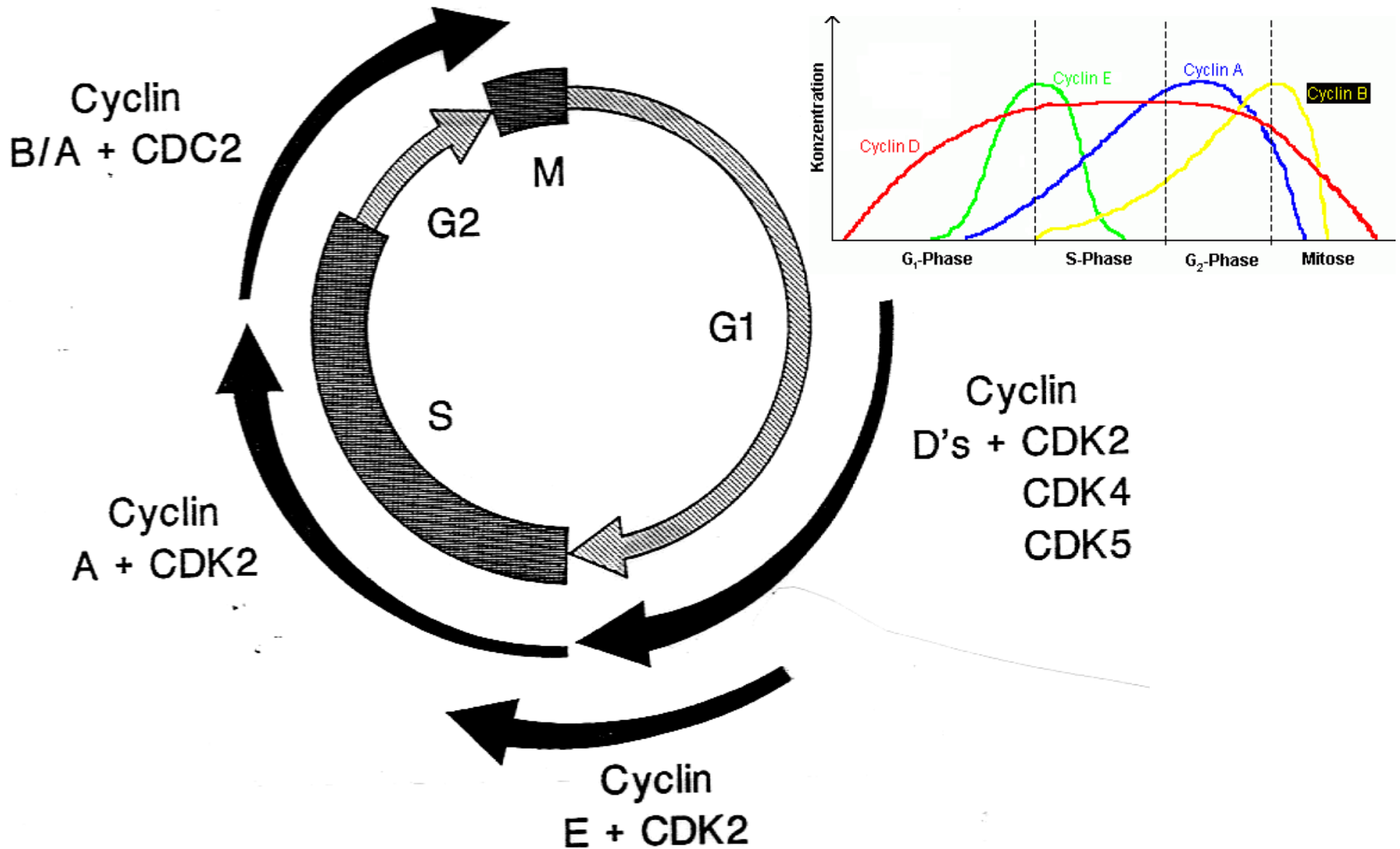
How does the cell decide which pathway to use?

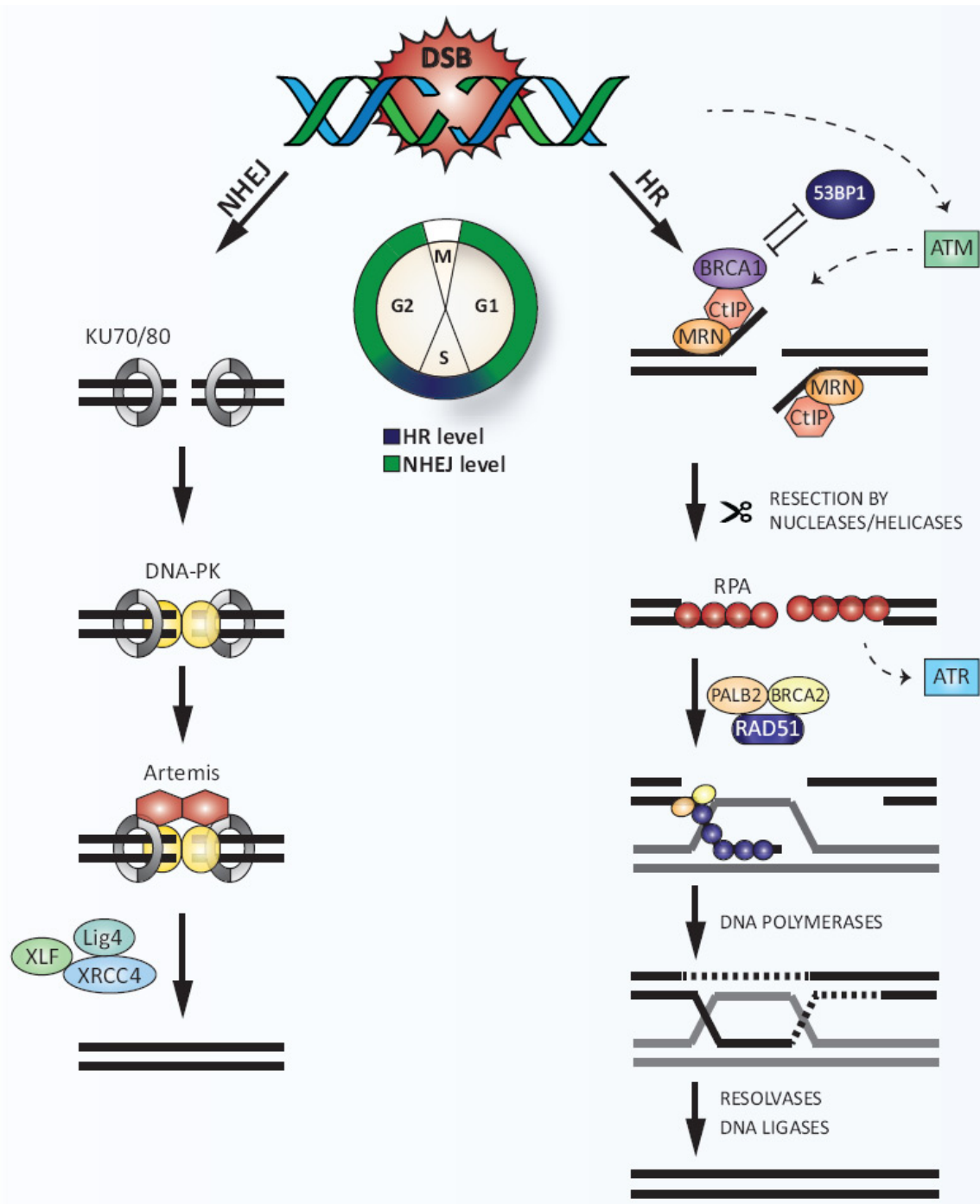
The Cell Cycle



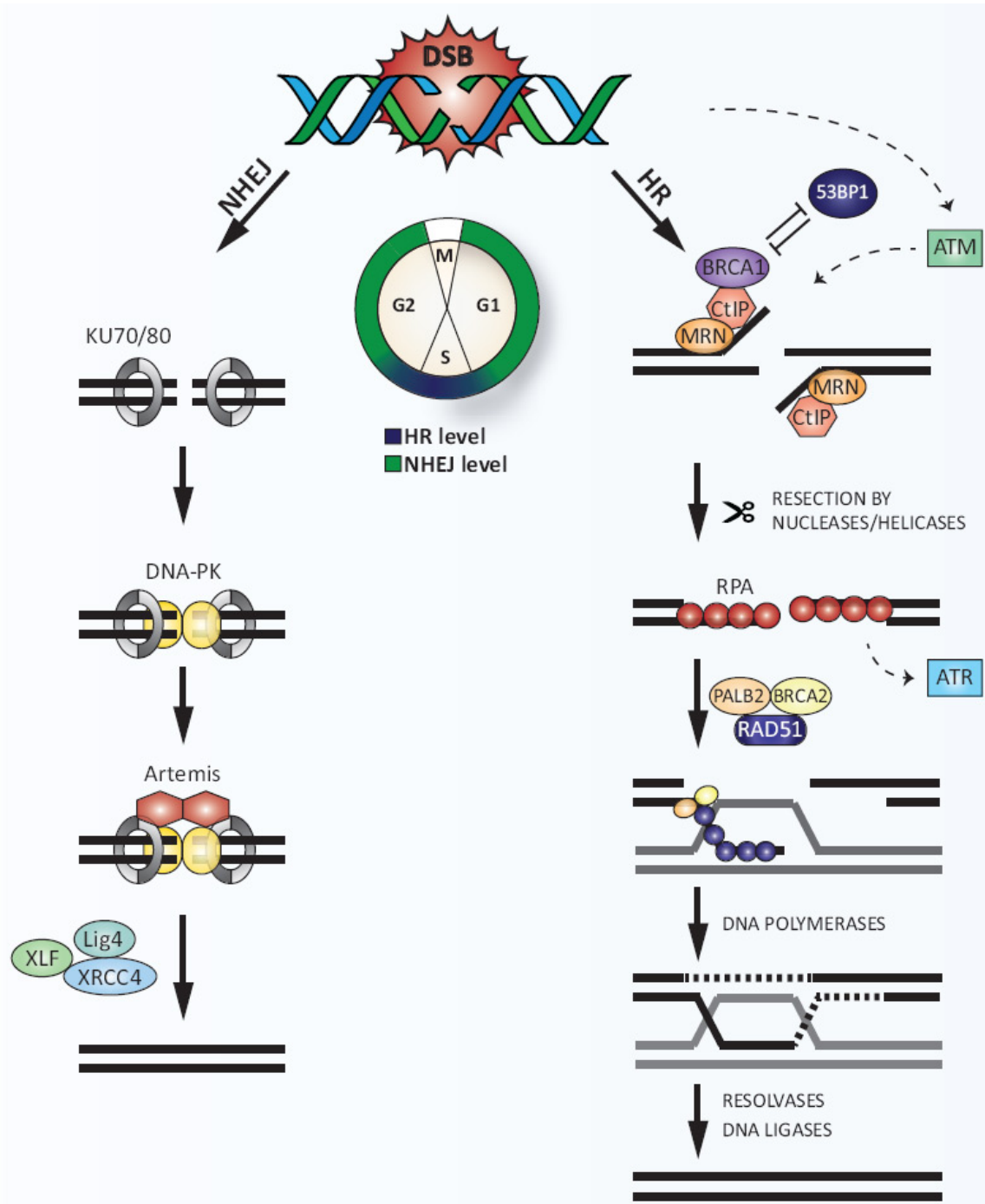
Duplication of chromosomes
DNA Replication

Progression through the Cell Cycle REQUIRES a series of cyclins and cyclin-dependent-kinases (CDKs)

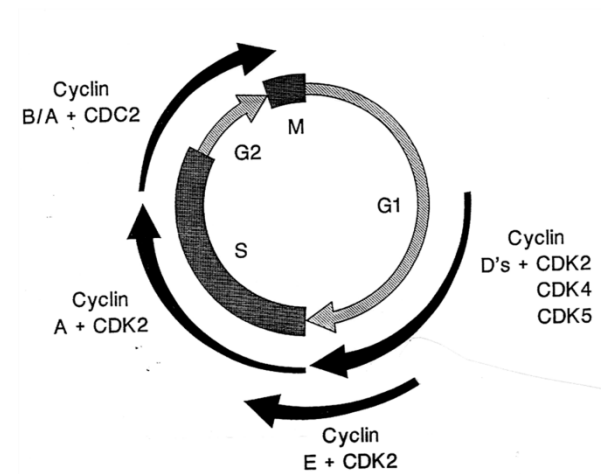




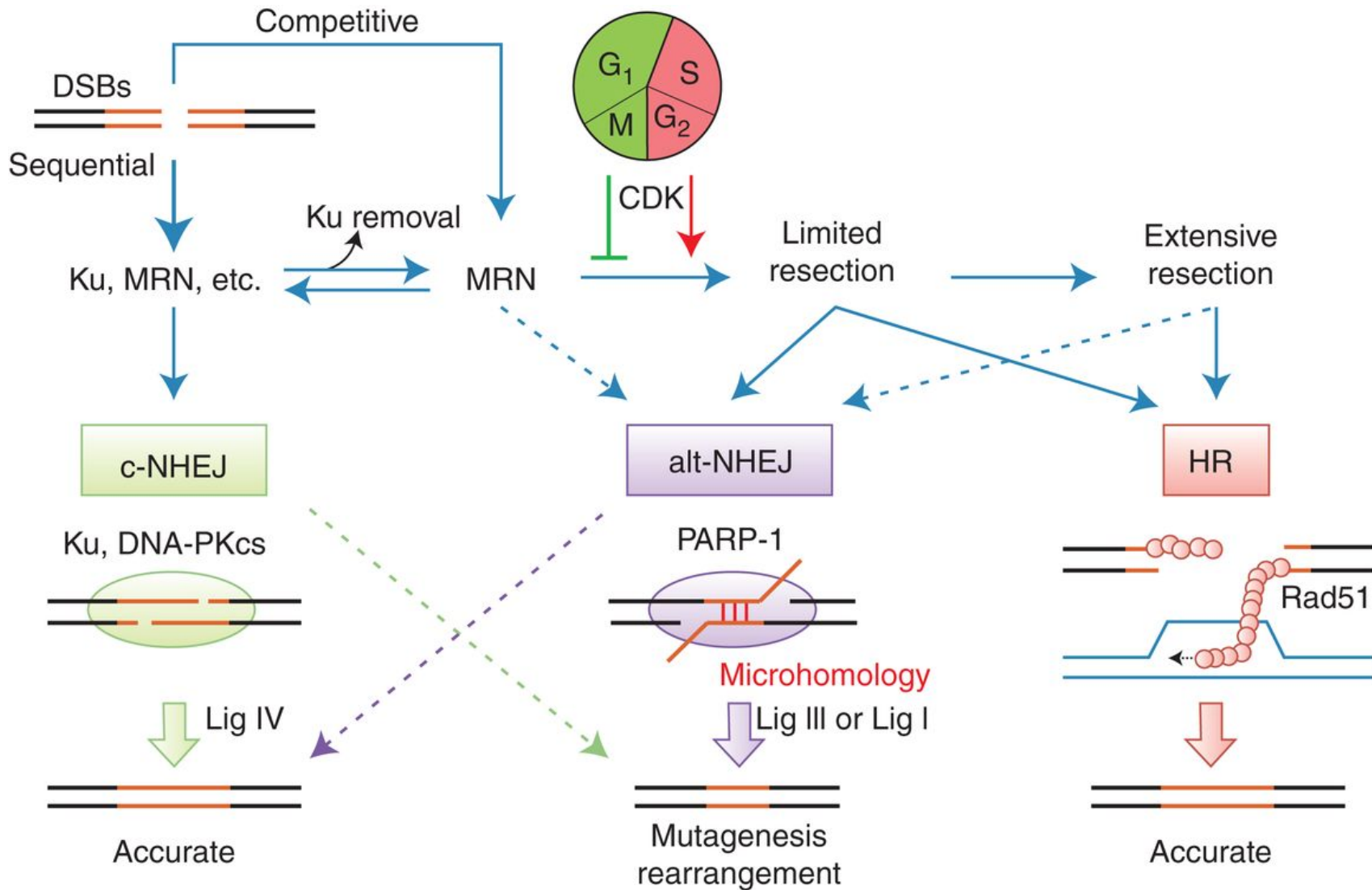
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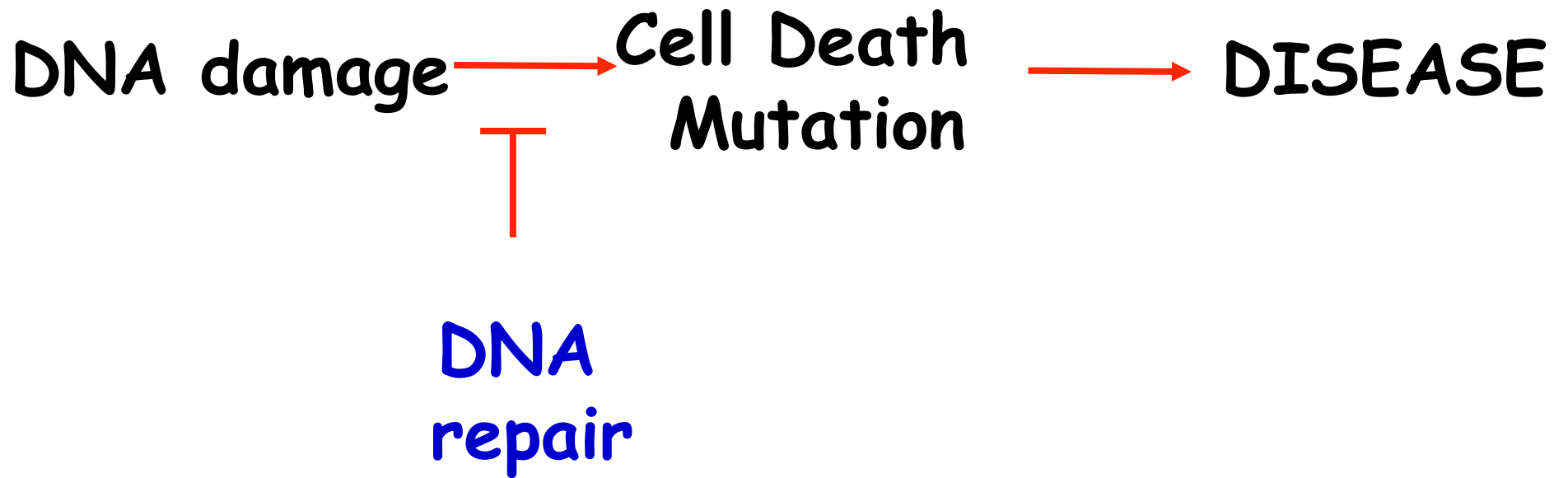


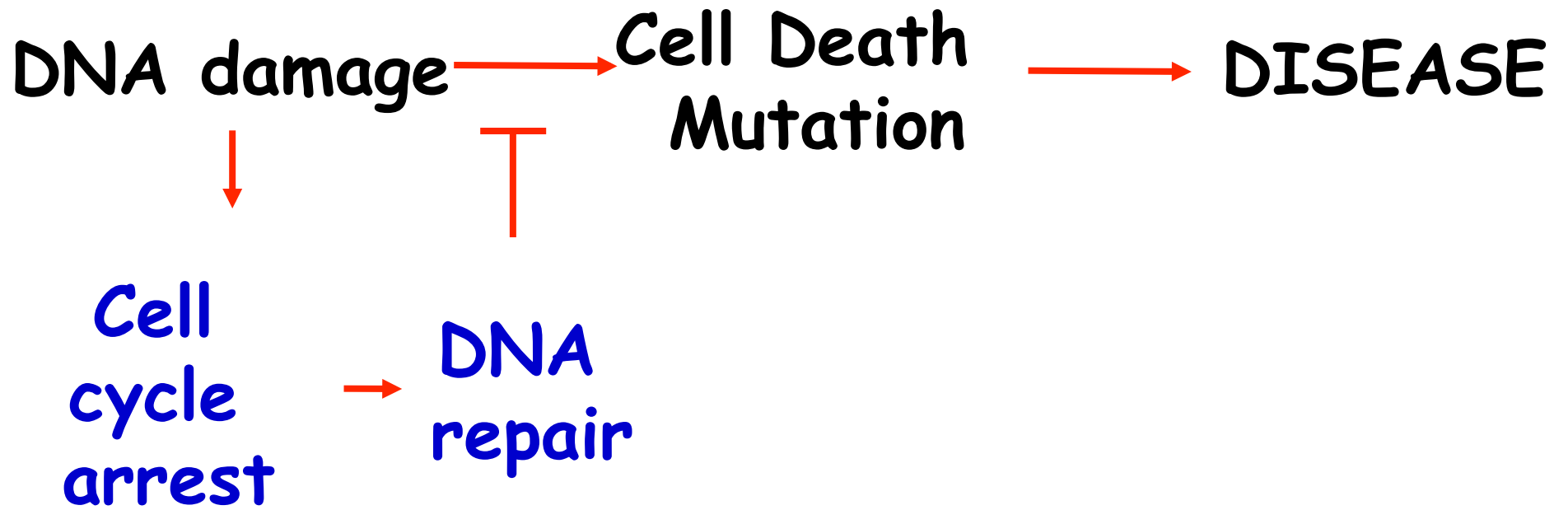
CyclinA-
CDK2
targets the
CtIP/BRCA1
complex

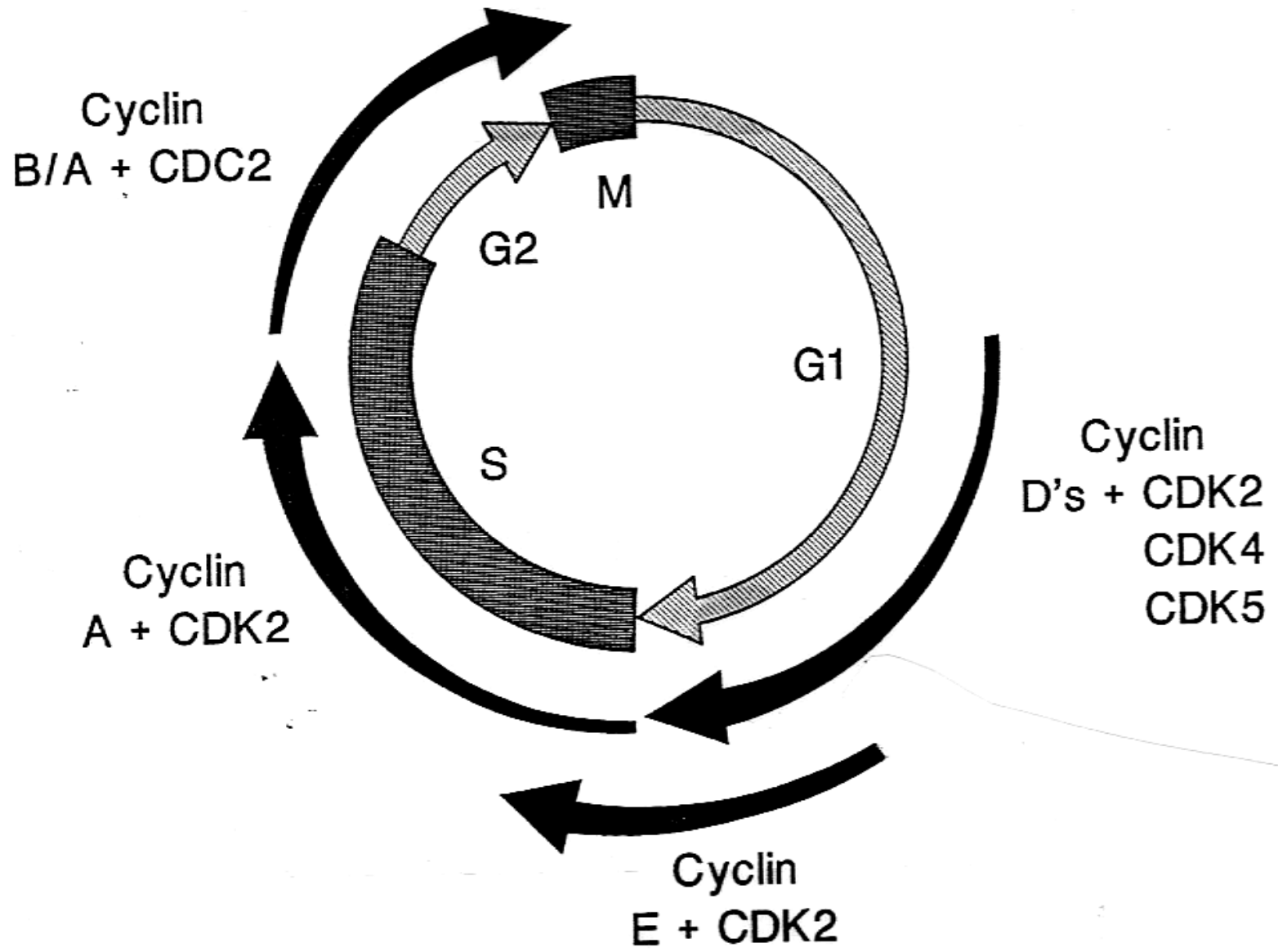


Disposition of DSBs between repair pathways.

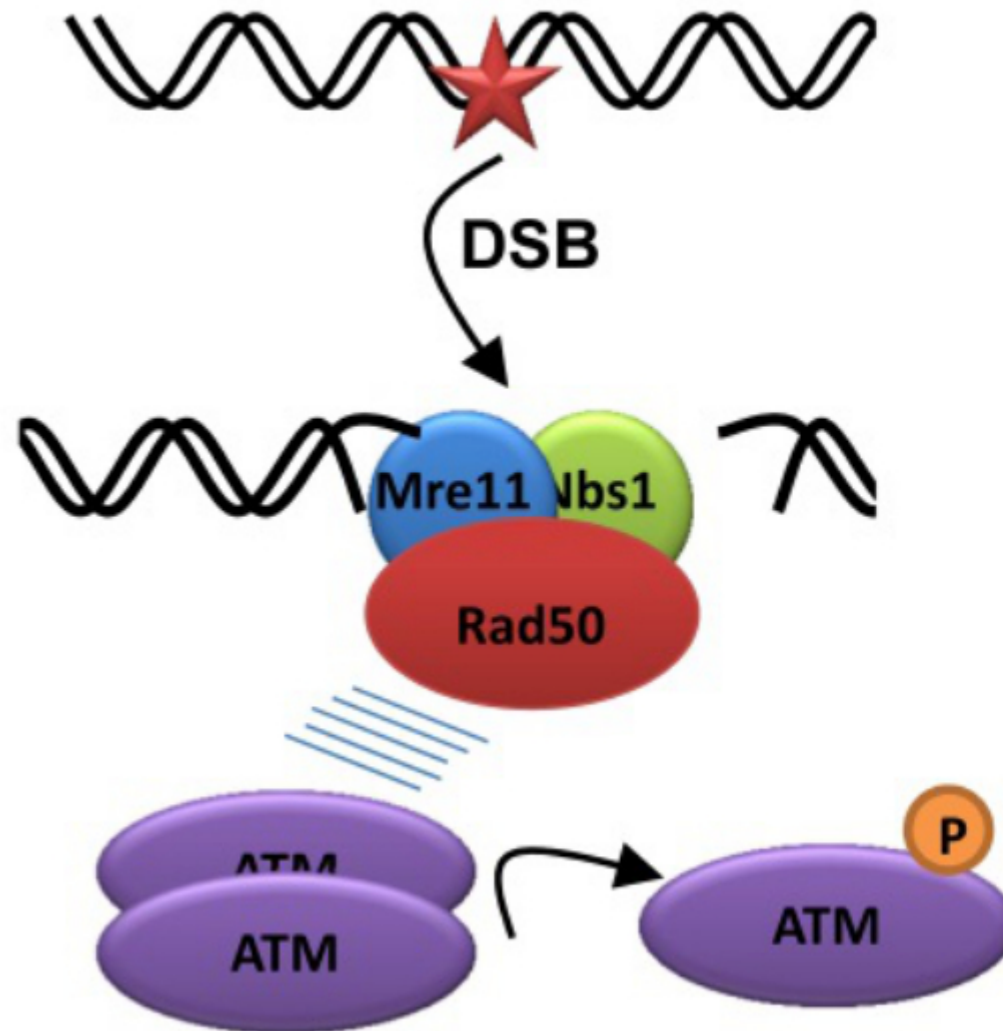


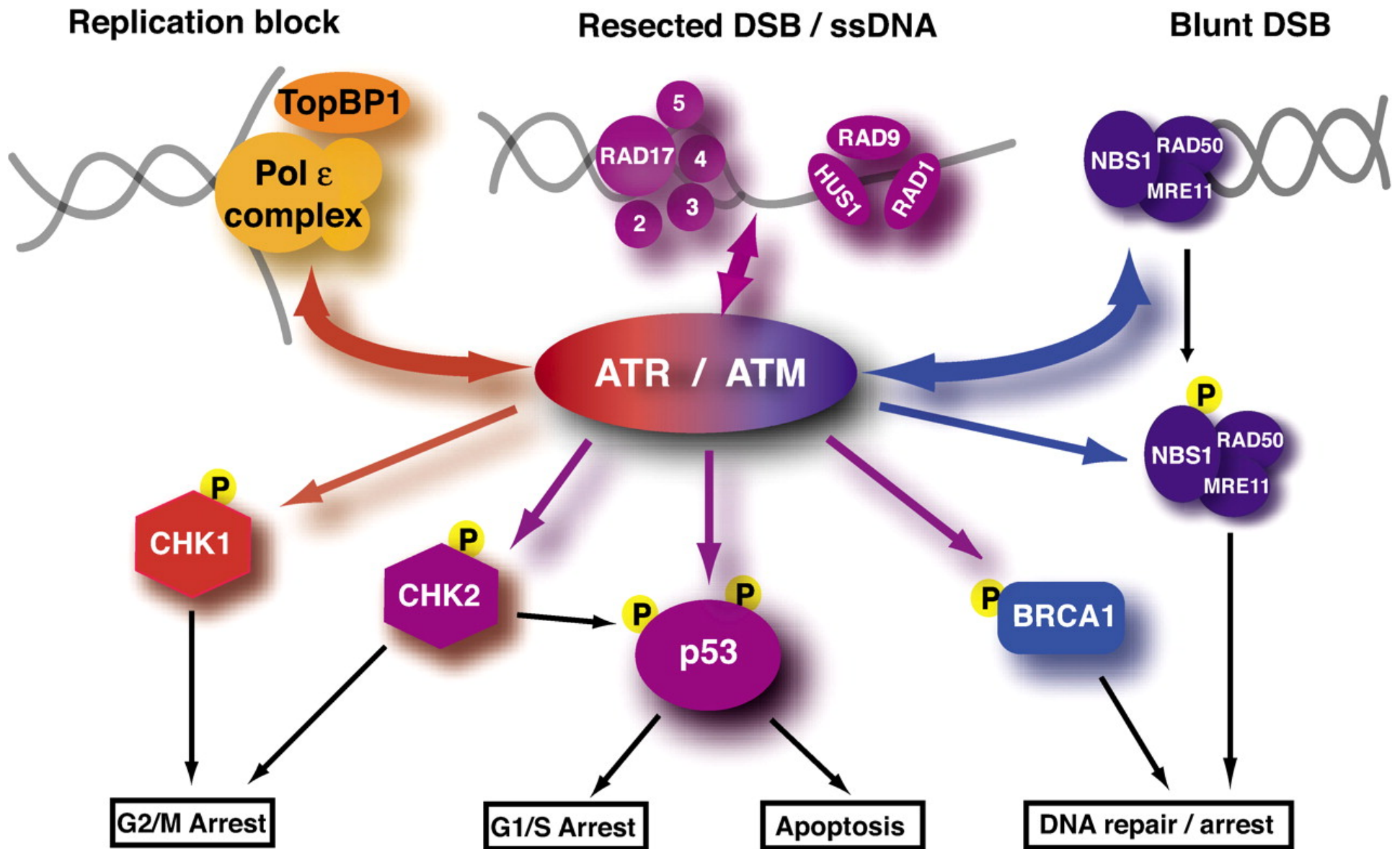




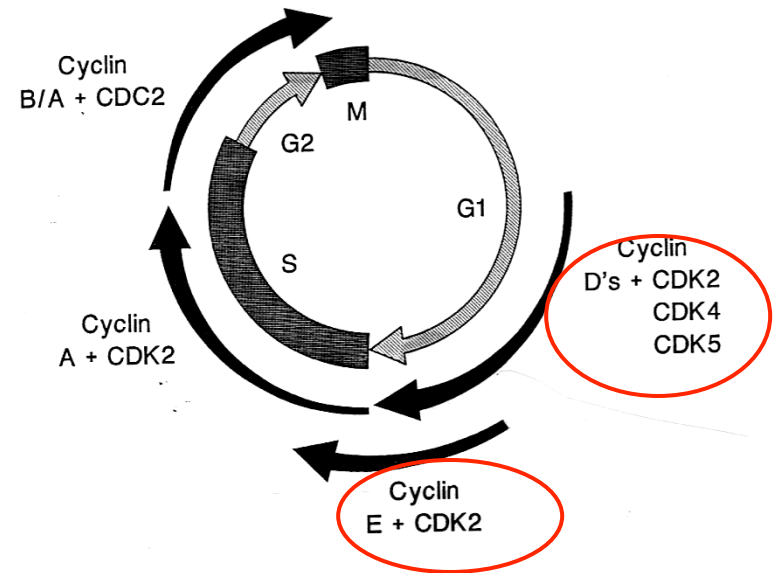
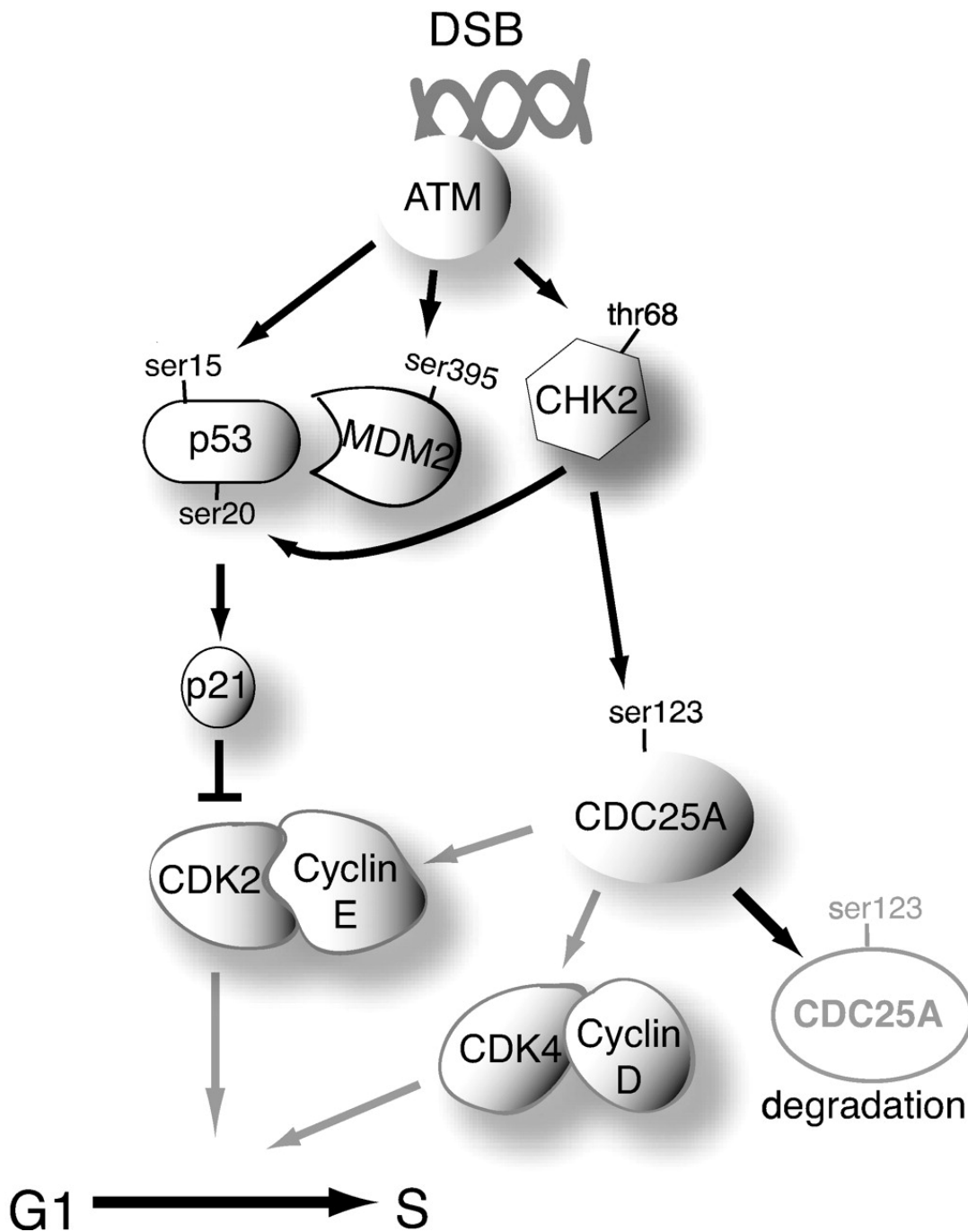


Signaling at DSBs – ATM kinase activated

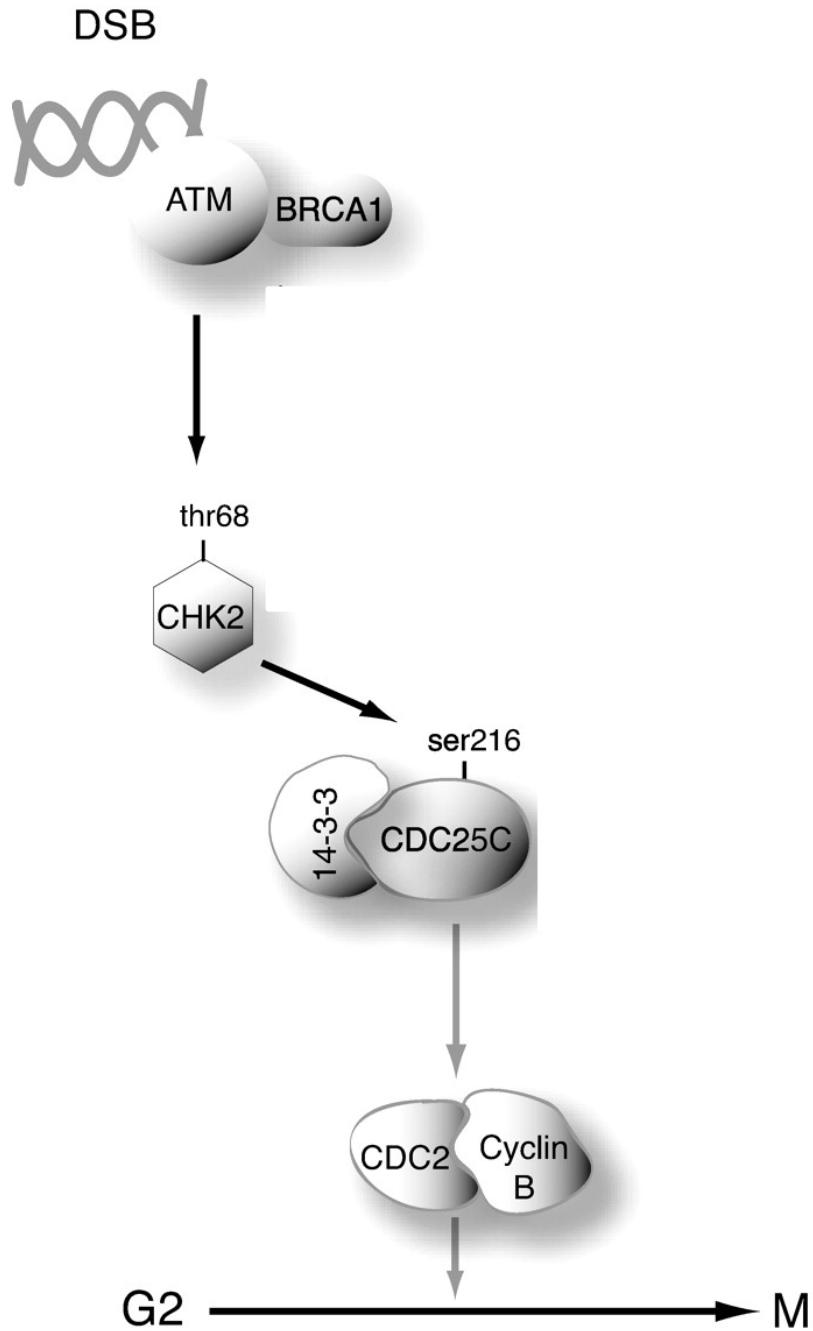




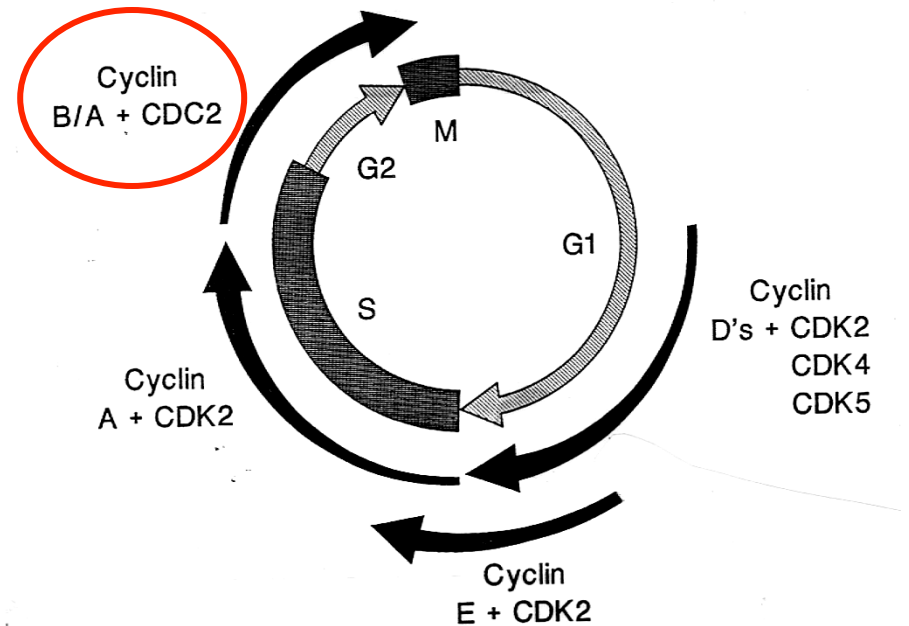
Two routes to activate a G1 arrest



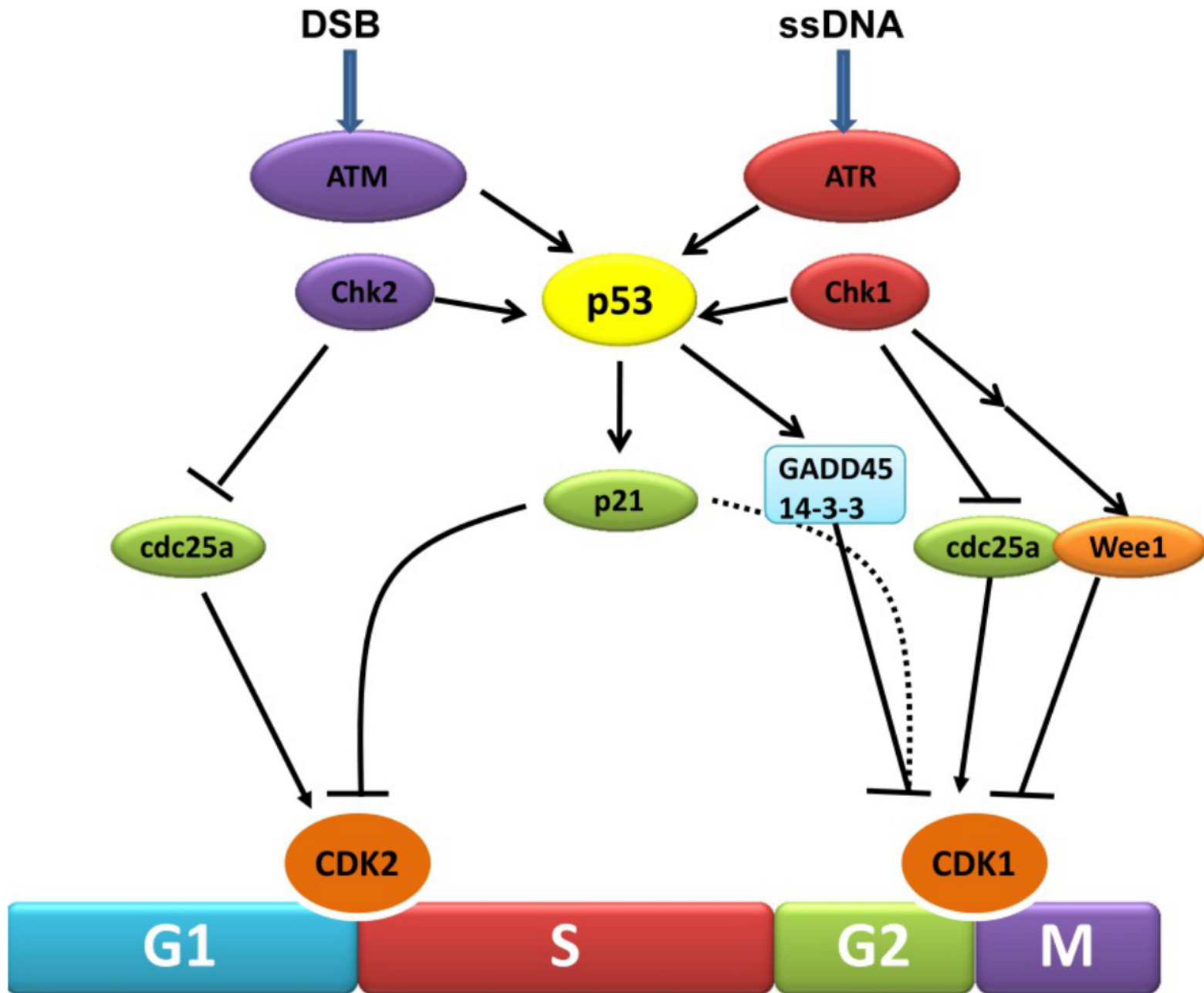
CDC25A is a phosphatase that must act on CDK2 and CDK4 to activate the complexes

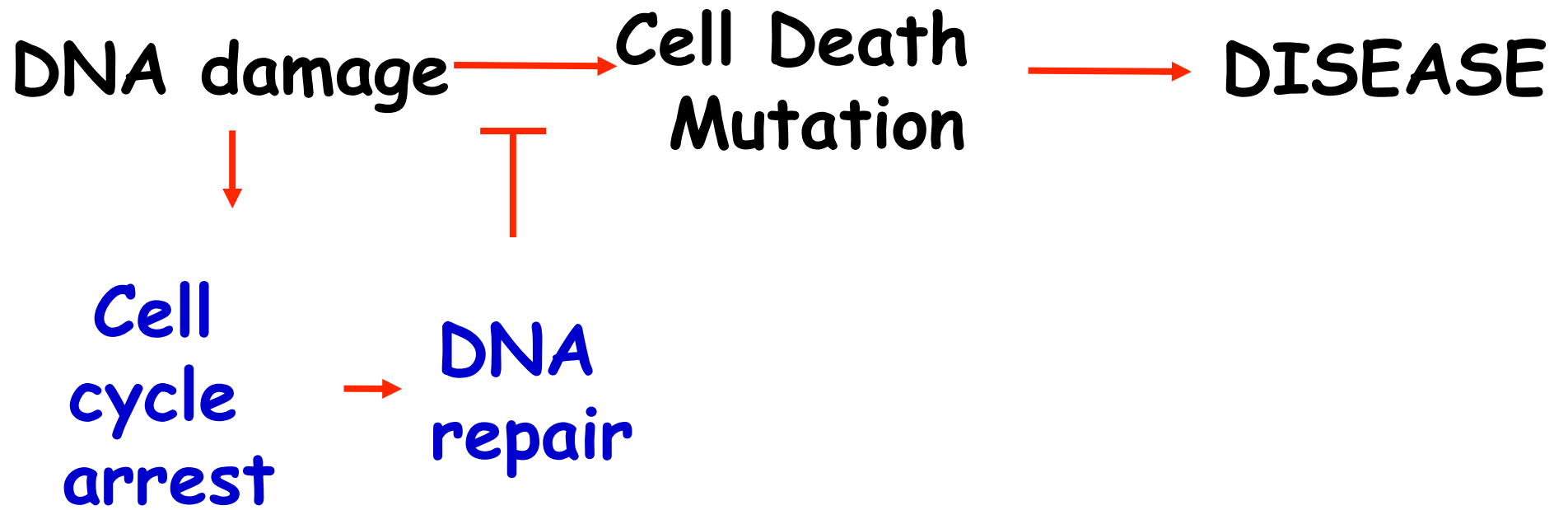


G2/M Arrest



CDC25C phosphatase has different substrate..Cdc2 to target G2/M transition





Ataxia Telangiectasia – Cancer Prone

Defective DNA
Damage Responses
can affect both
neurodegeneration
and cancer
susceptibility



Ataxia Telangiectasia

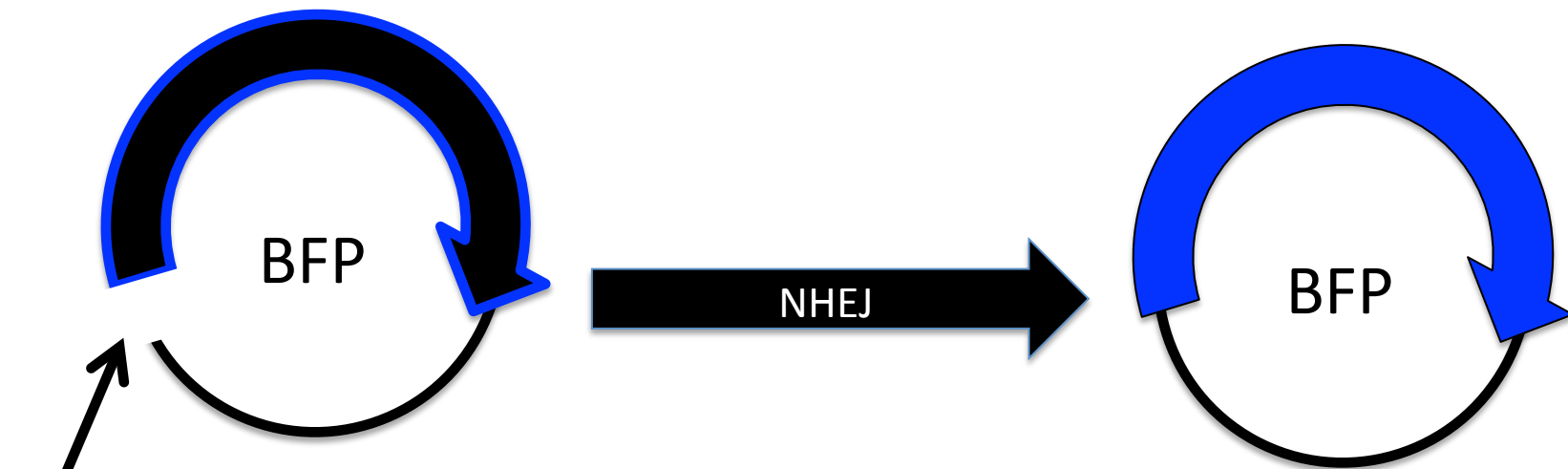
- Staggering gait
- Muscular un-coordination
- Mental retardation
- Dilation of small blood vessels
- Immune dysfunction
- Cancer prone...lymphomas
- Cells from AT patients have lost cell cycle checkpoints

Key Experimental Methods for Module 1

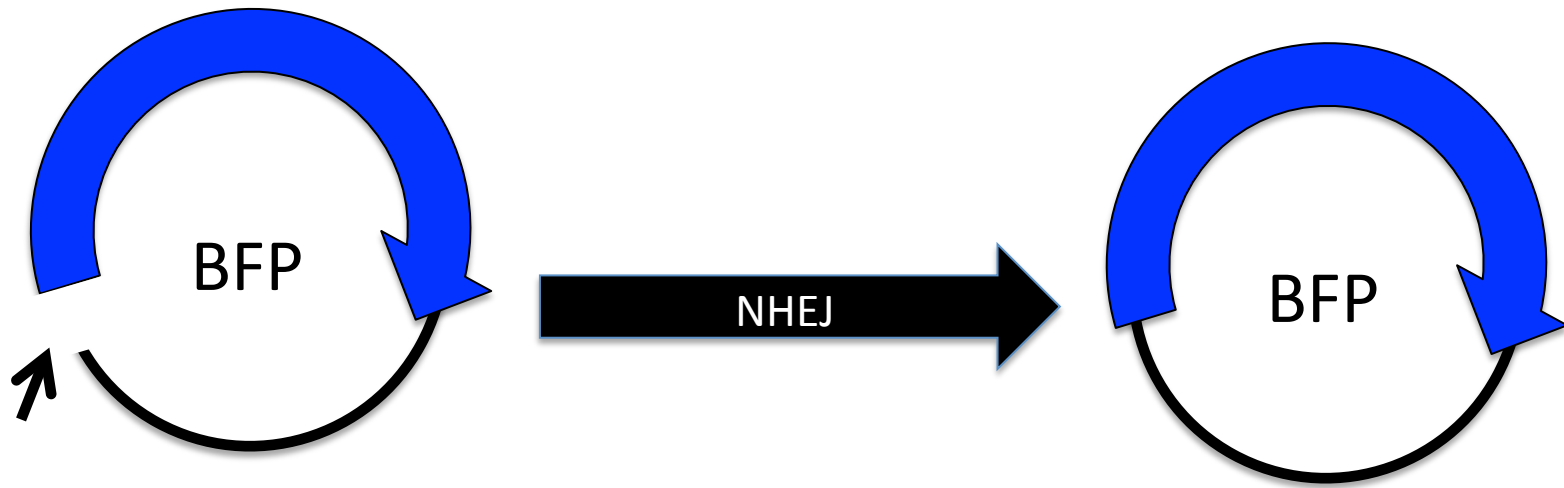


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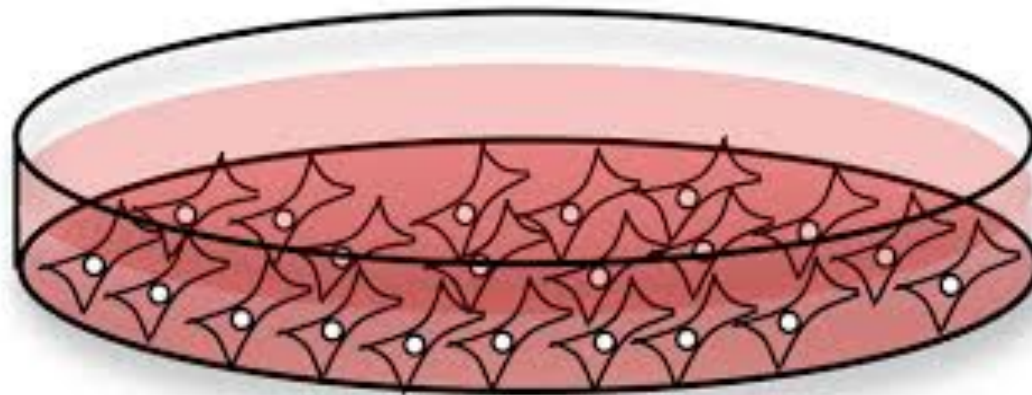
Basis for the fluorescent reporter assay:

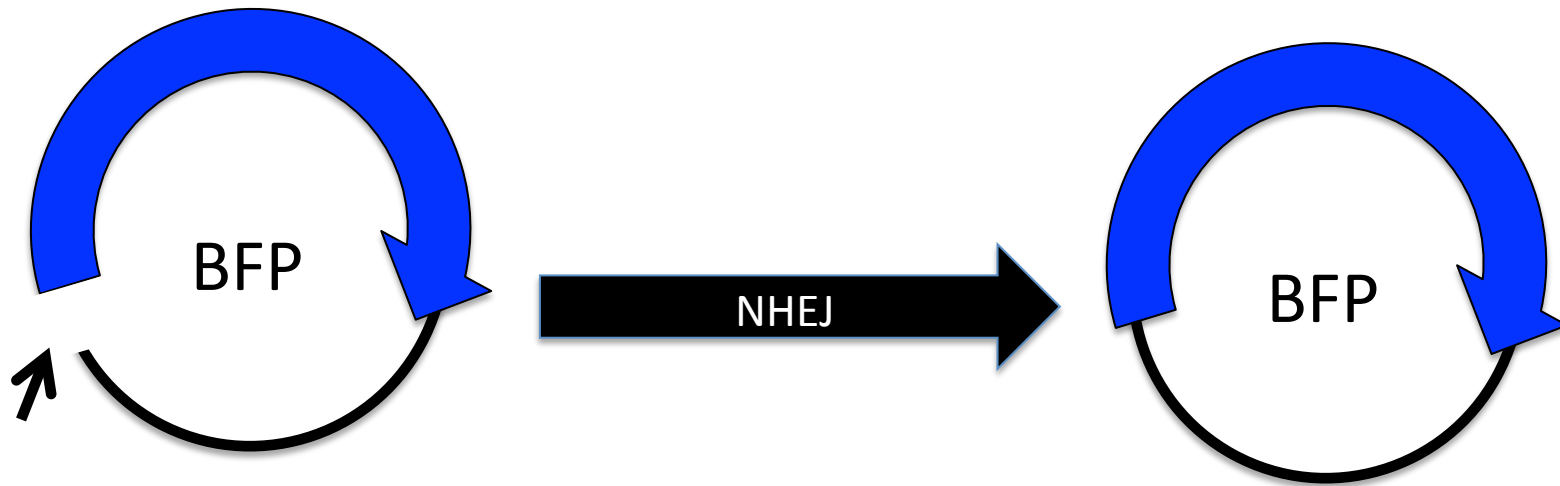


Following digest, the substrate contains a DSB in the 5' UTR that prevents fluorescent reporter expression

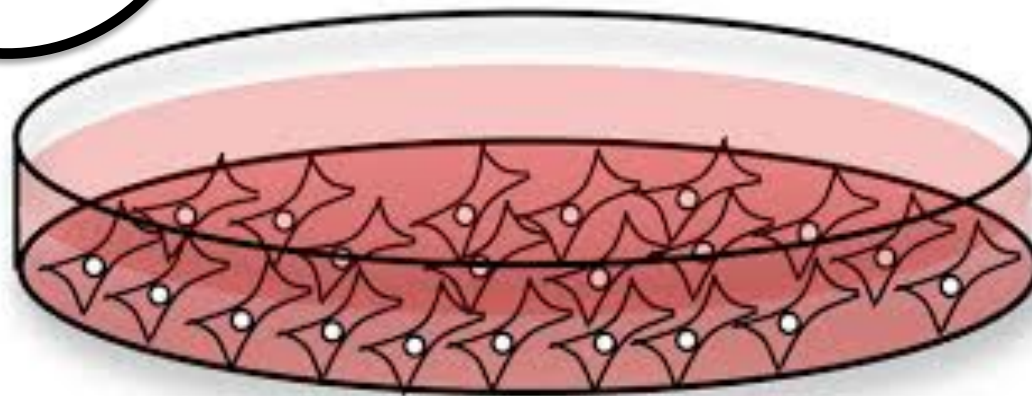
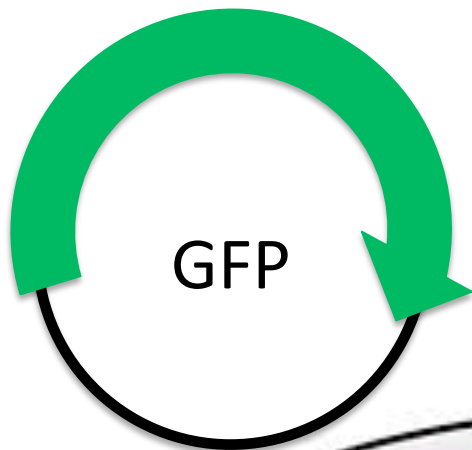


Substrate contains a DNA double strand break

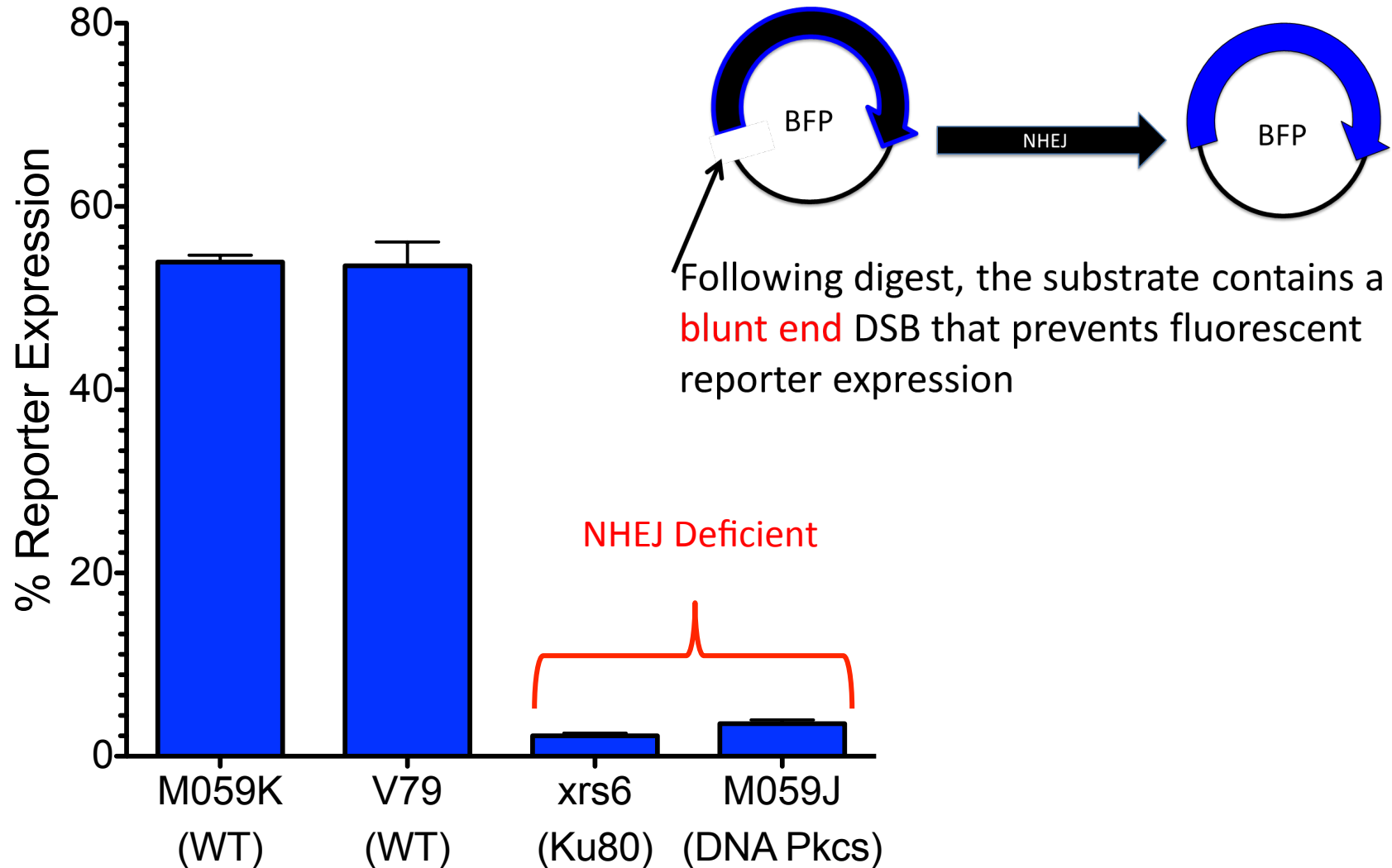




Substrate contains a DNA double strand break

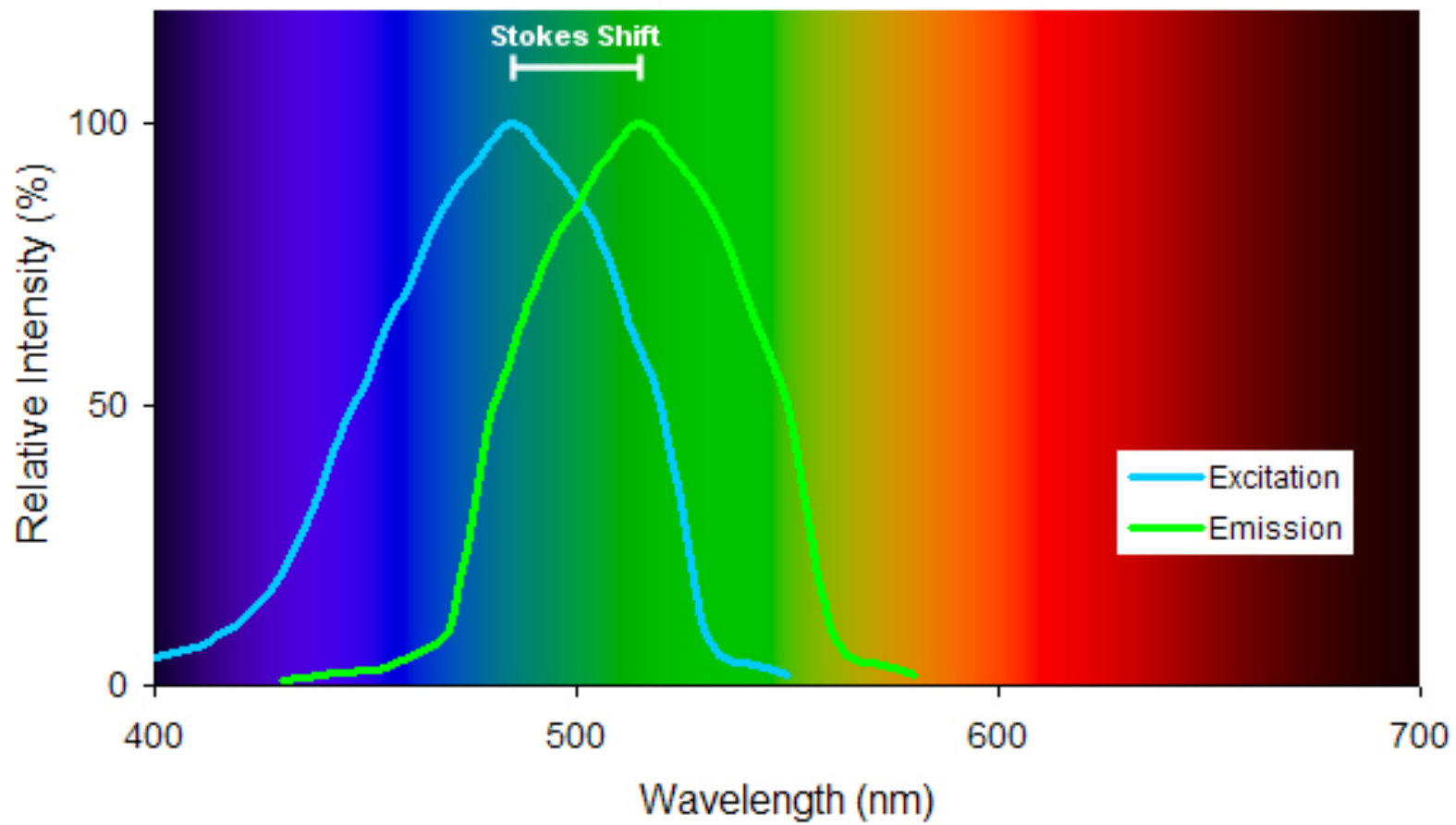
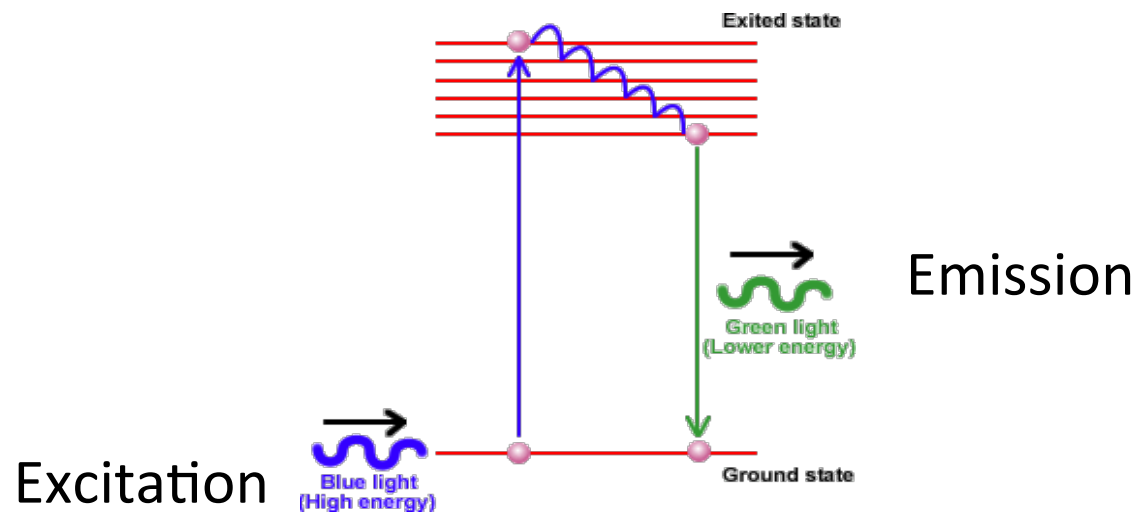


NHEJ HCR in WT and NHEJ defective cells at 18 hours post-transfection:

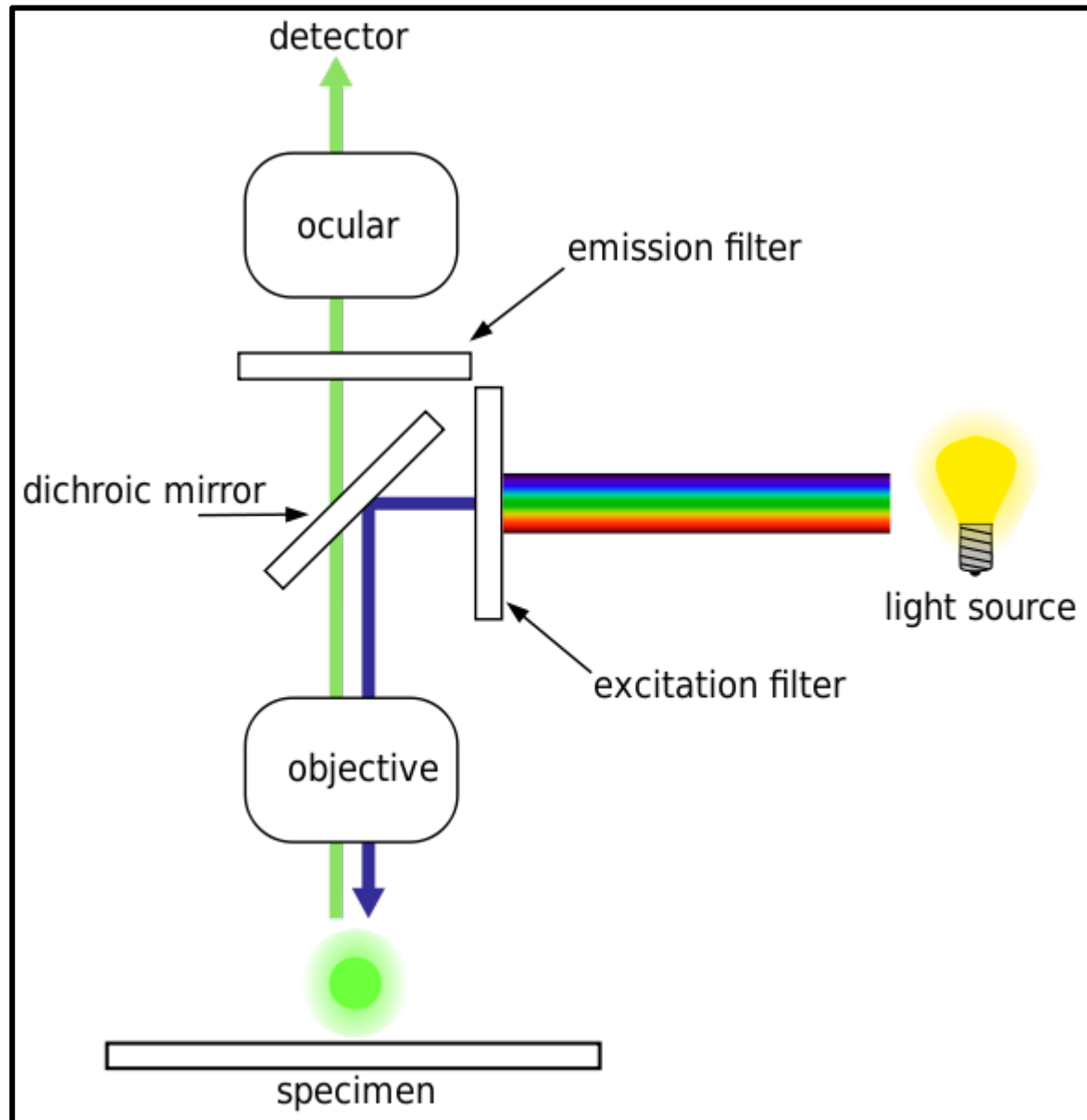


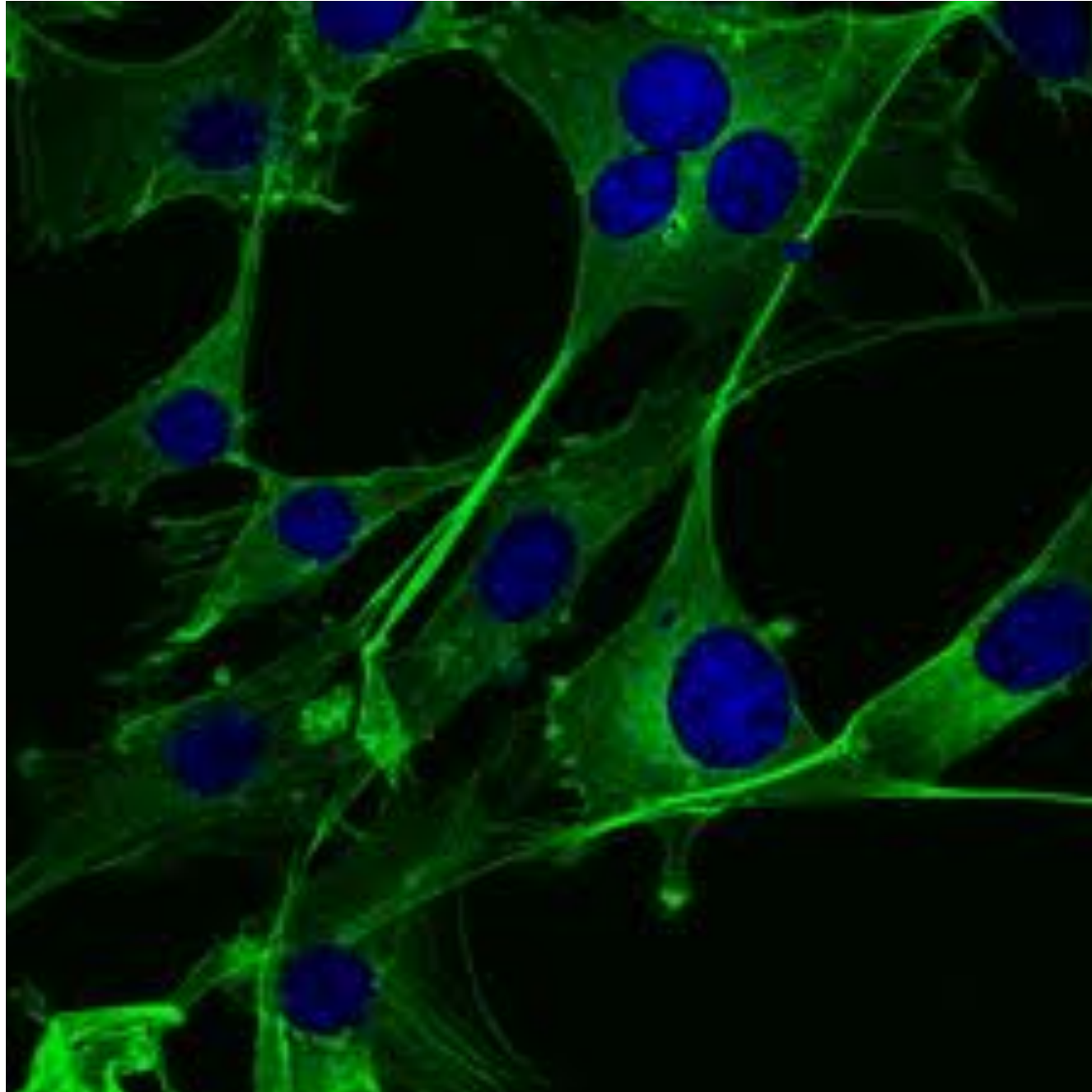
Some restriction enzymes

| Enzyme | Source organism | Restriction recognition site in double-stranded DNA | Structure of the cleaved products | |
|-------------|----------------------------|---|--|---|
| (a) | <i>EcoRI</i> | <i>Escherichia coli</i> | | |
| | | | <p style="text-align: center;">5' overhang</p> | |
| | <i>PstI</i> | <i>Providencia stuartii</i> | | |
| | | | <p style="text-align: center;">3' overhang</p> | |
| <i>SmaI</i> | <i>Serratia marcescens</i> | | | |
| | | | <p style="text-align: center;">Blunt ends</p> | |
| (b) | <i>HaeIII</i> | <i>Haemophilus aegyptius</i> | | |
| | | | | <p style="text-align: center;">Blunt ends</p> |
| | <i>HpaII</i> | <i>Haemophilus parainfluenzae</i> | | |
| | | | <p style="text-align: center;">5' overhang</p> | |



Fluorescence Detection



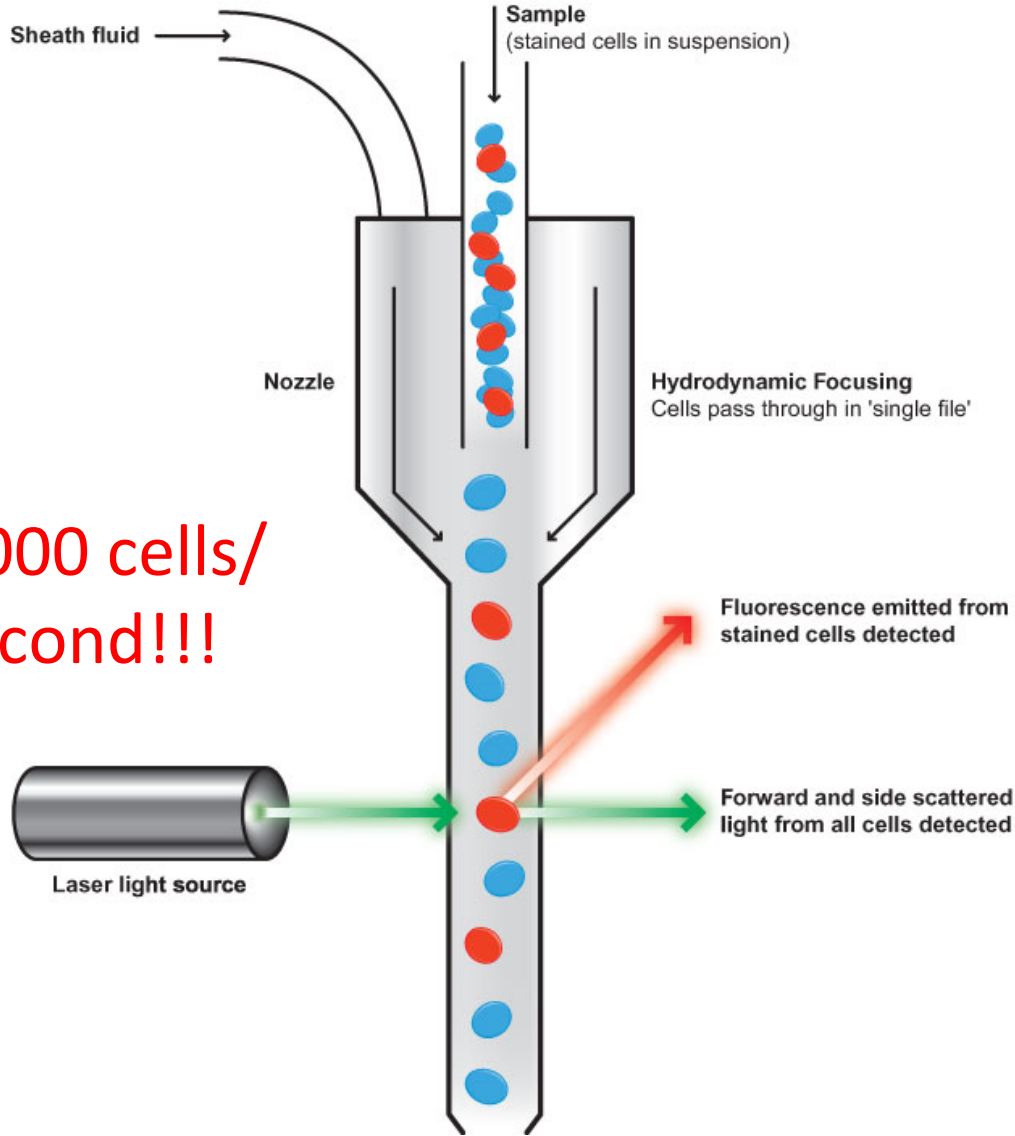


DNA – Blue

GFP - Green

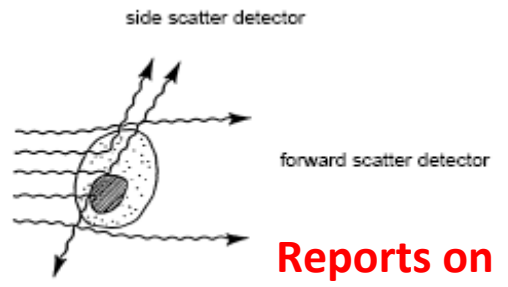
Flow Cytometry

Flow Cytometry

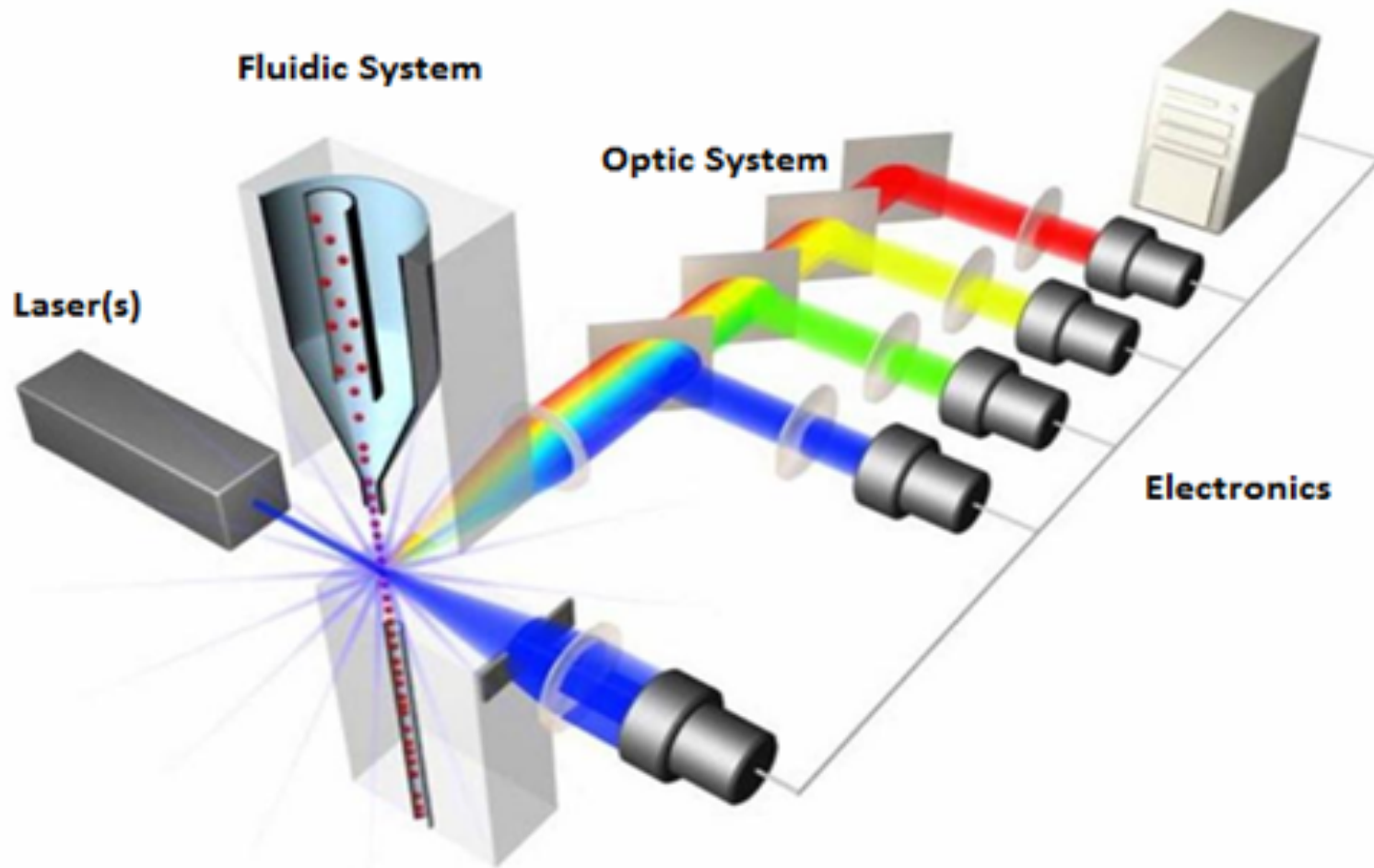


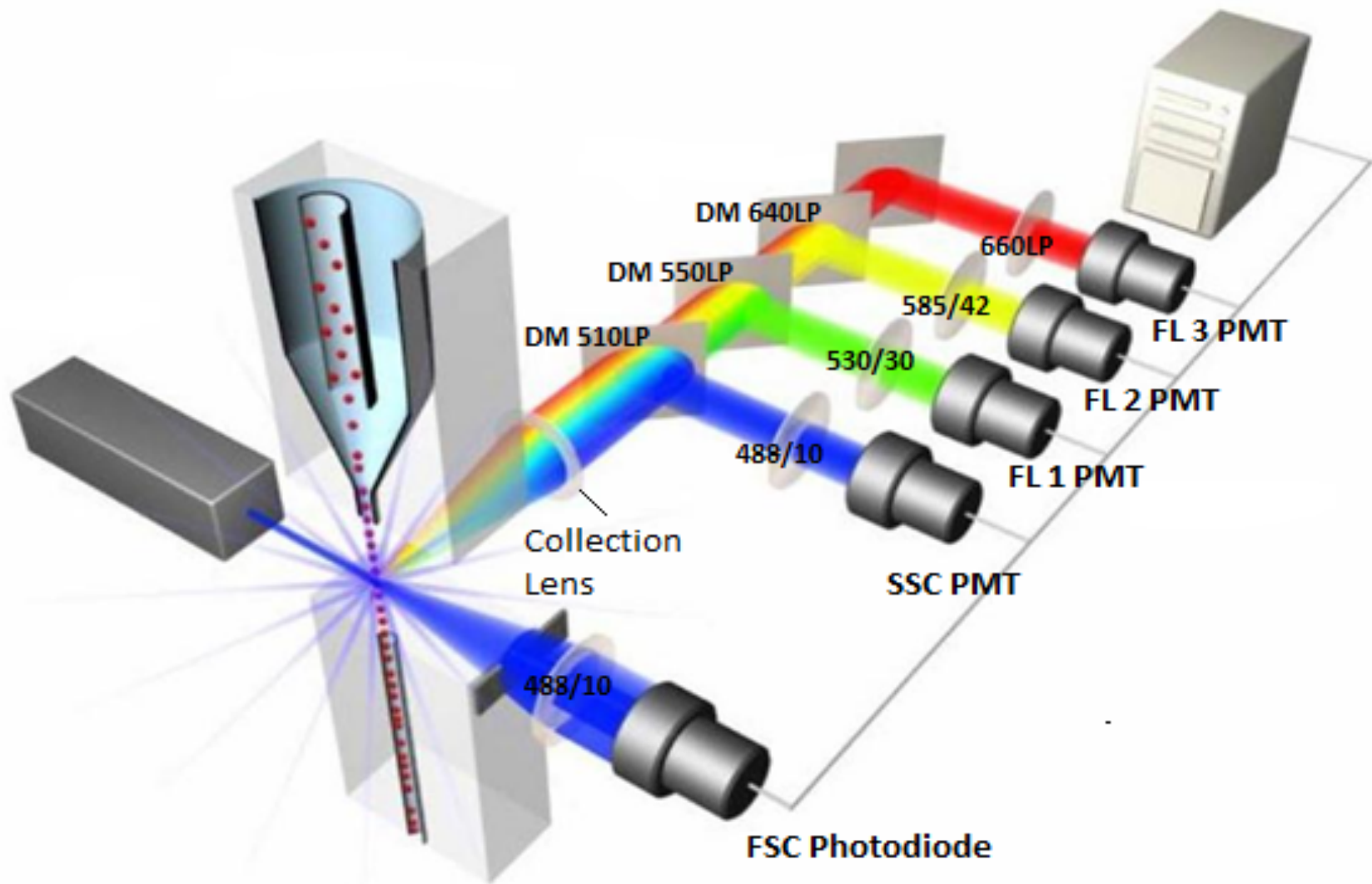
50,000 cells/
second!!!

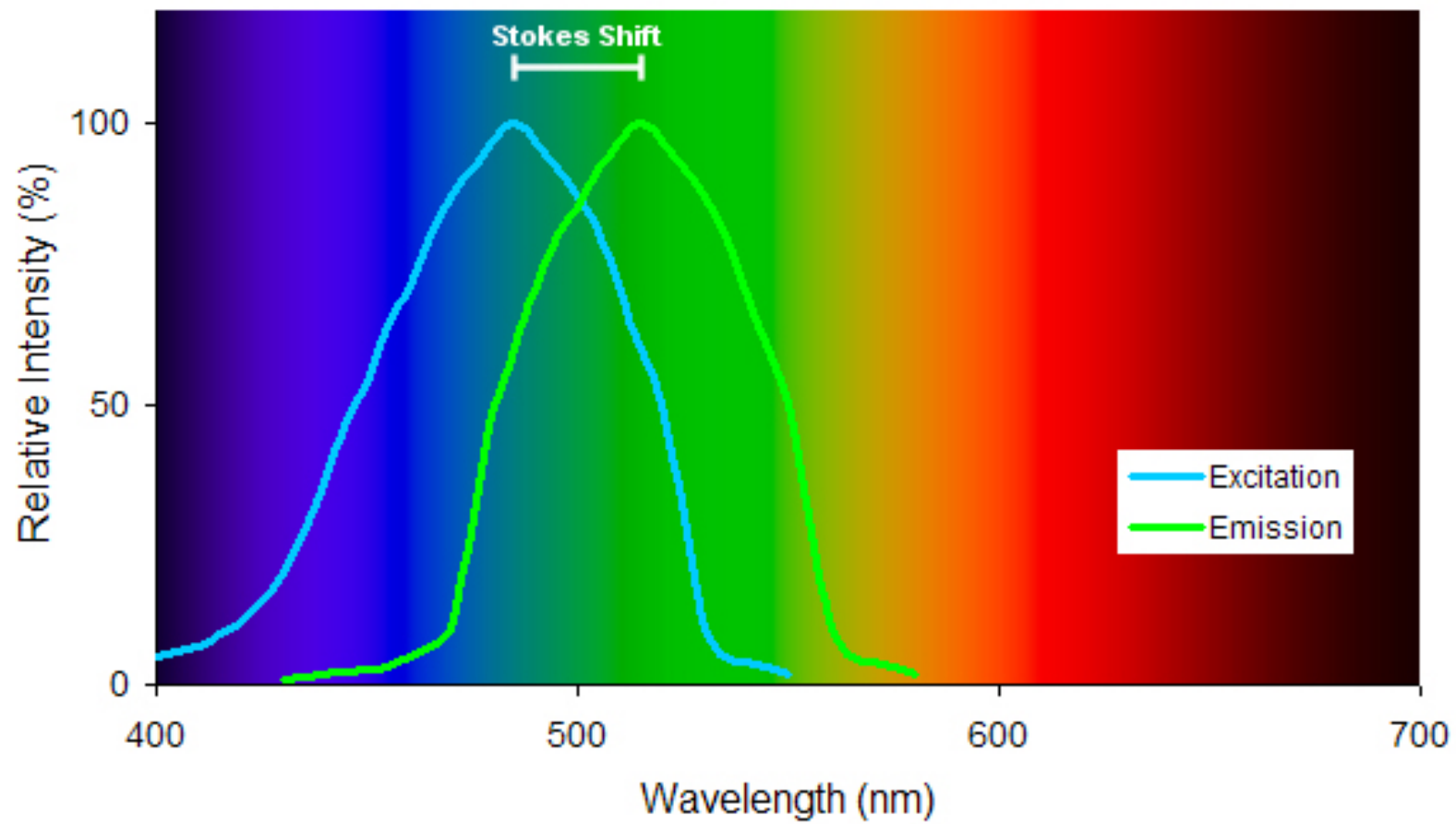
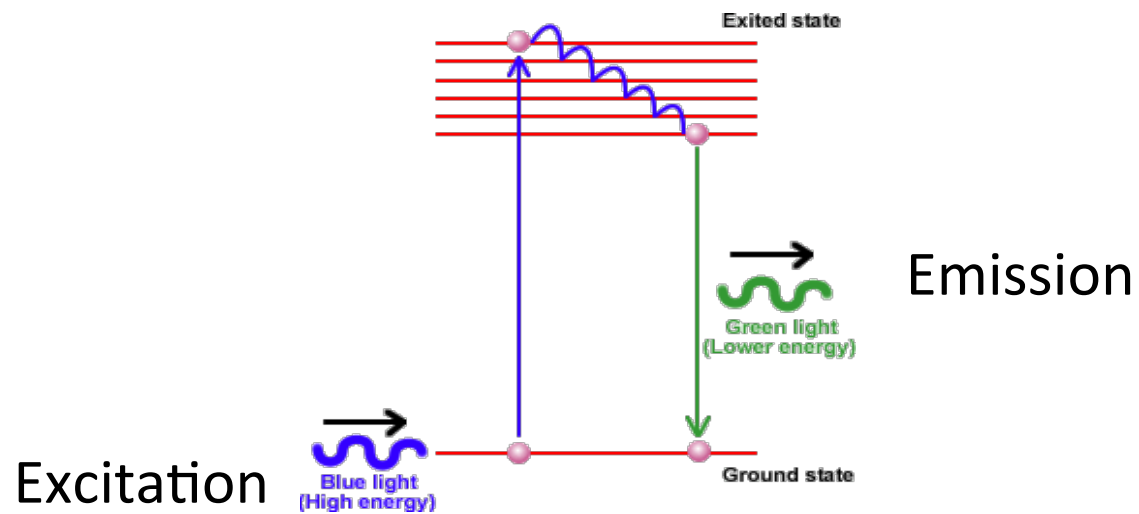
Reports on cell
complexity



Reports on
cell size

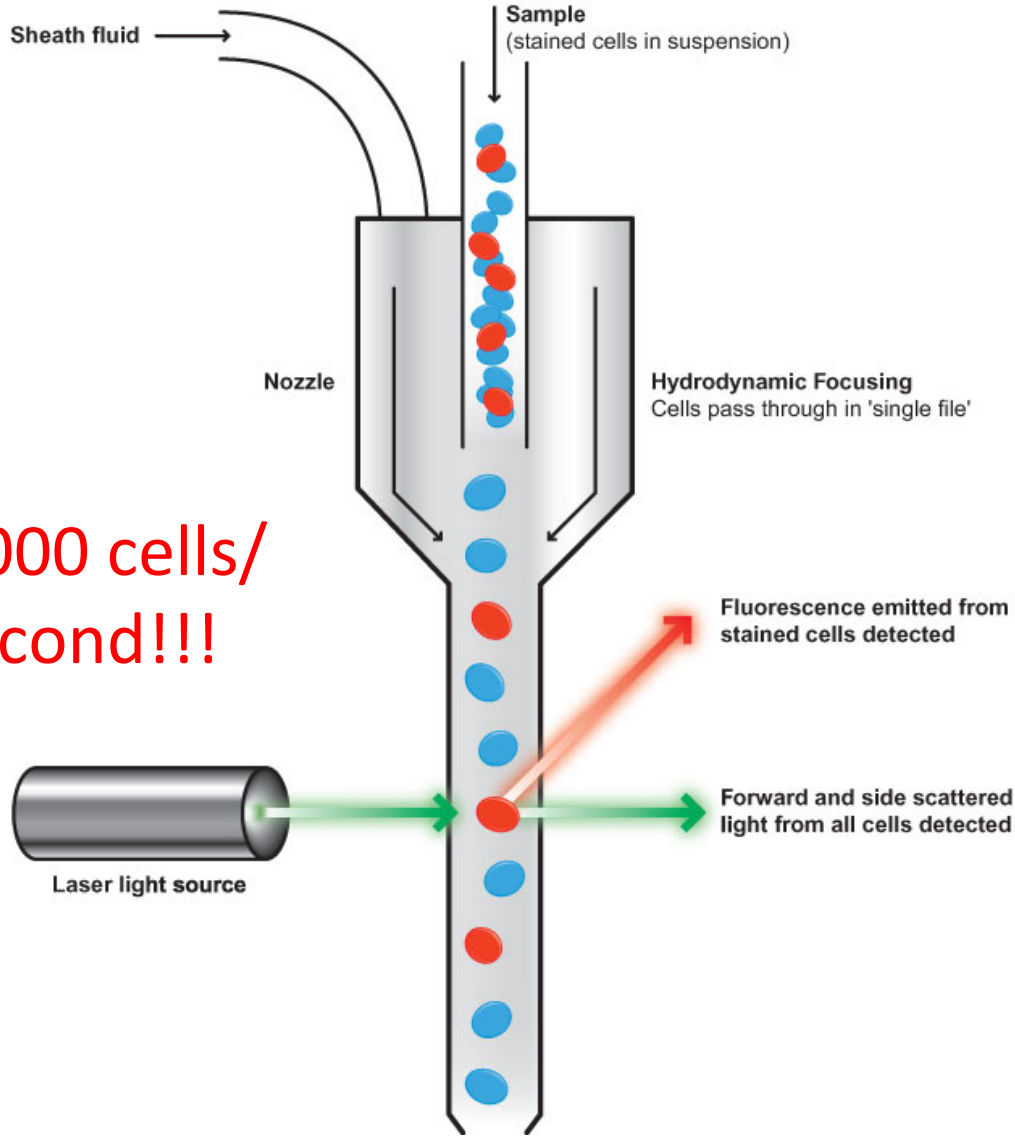






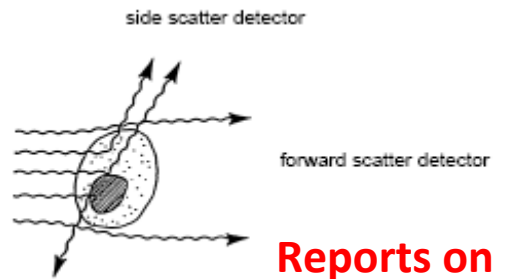
Flow Cytometry

Flow Cytometry



50,000 cells/
second!!!

Reports on cell
complexity



Reports on
cell size

Flow Cytometry

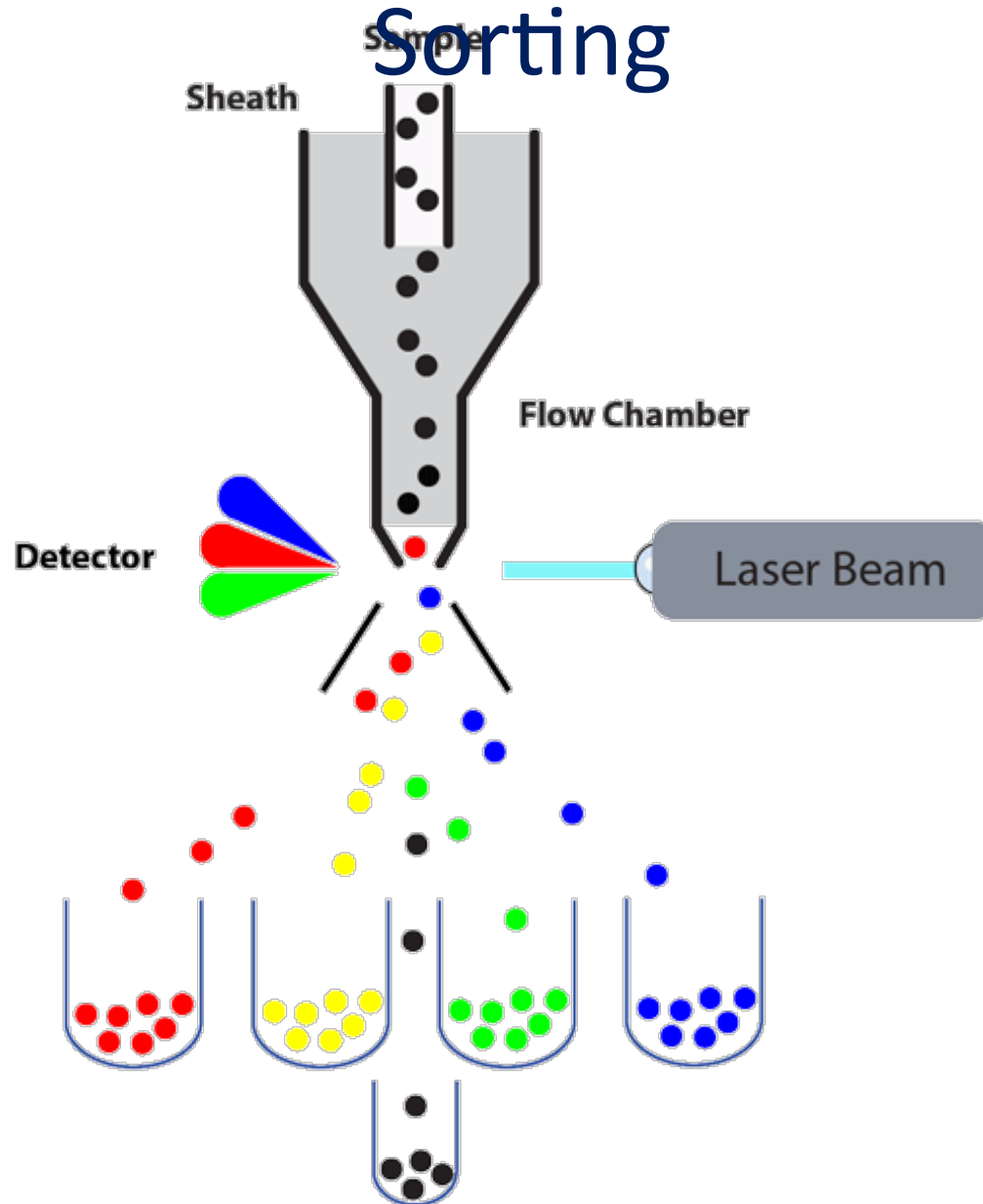
Flow cytometry analyzes cells one by one

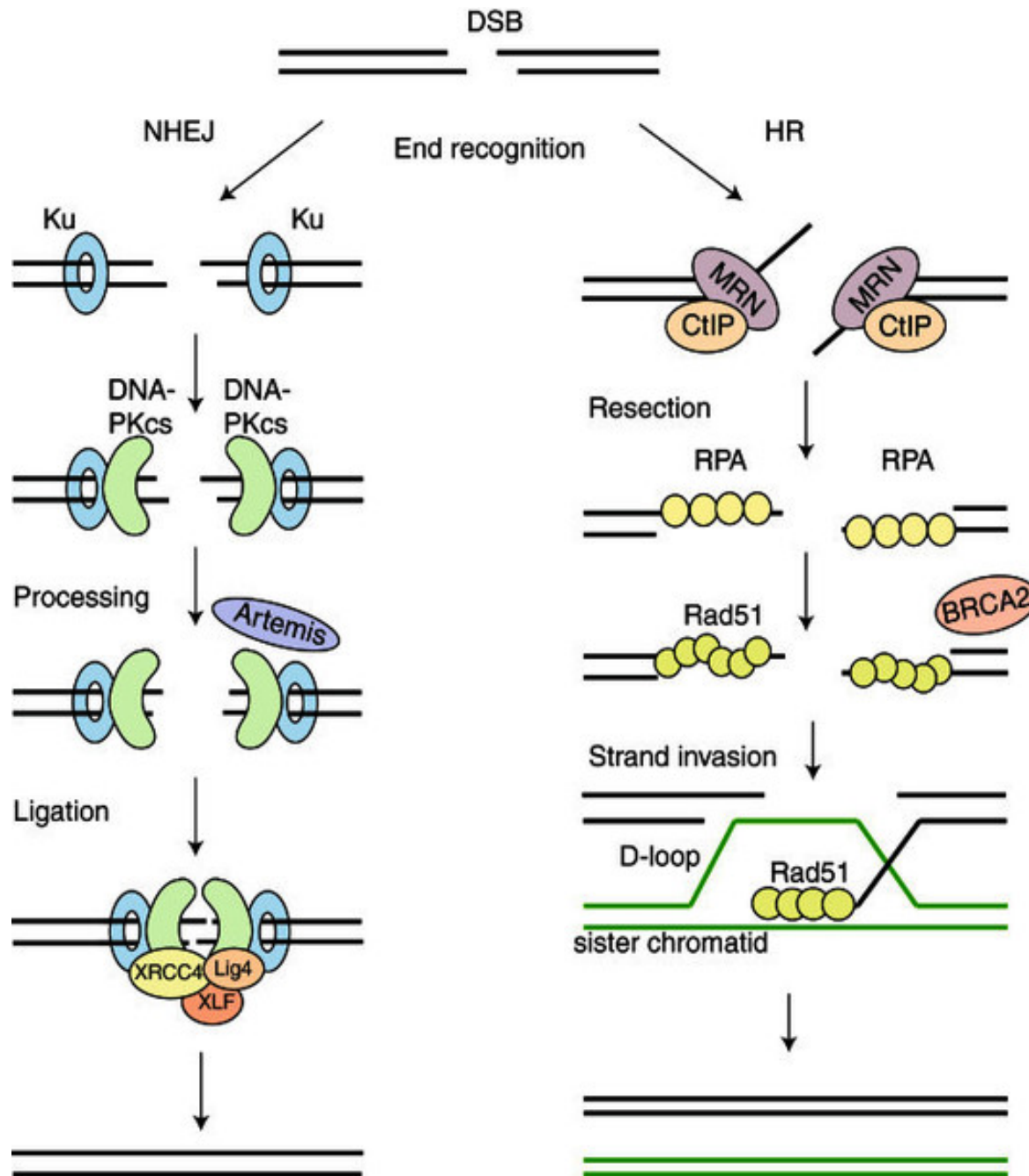
Fluorescence, diffracted and reflected light can be measured for each cell

Multiple lasers and multiple colors can be analyzed at millions of cells per minute

Resulting plots show the relative level of fluorescence of each cell for specific wave lengths.

FACS – Fluorescence Activated Cell Sorting





How does the cell decide which pathway to use?