

20.109 Spring 2015 Module 2 – Lecture 5

System Engineering and Protein Foundations



Shannon Hughes

Noreen Lyell

Leslie McLain

Nova Pishesha (TA)

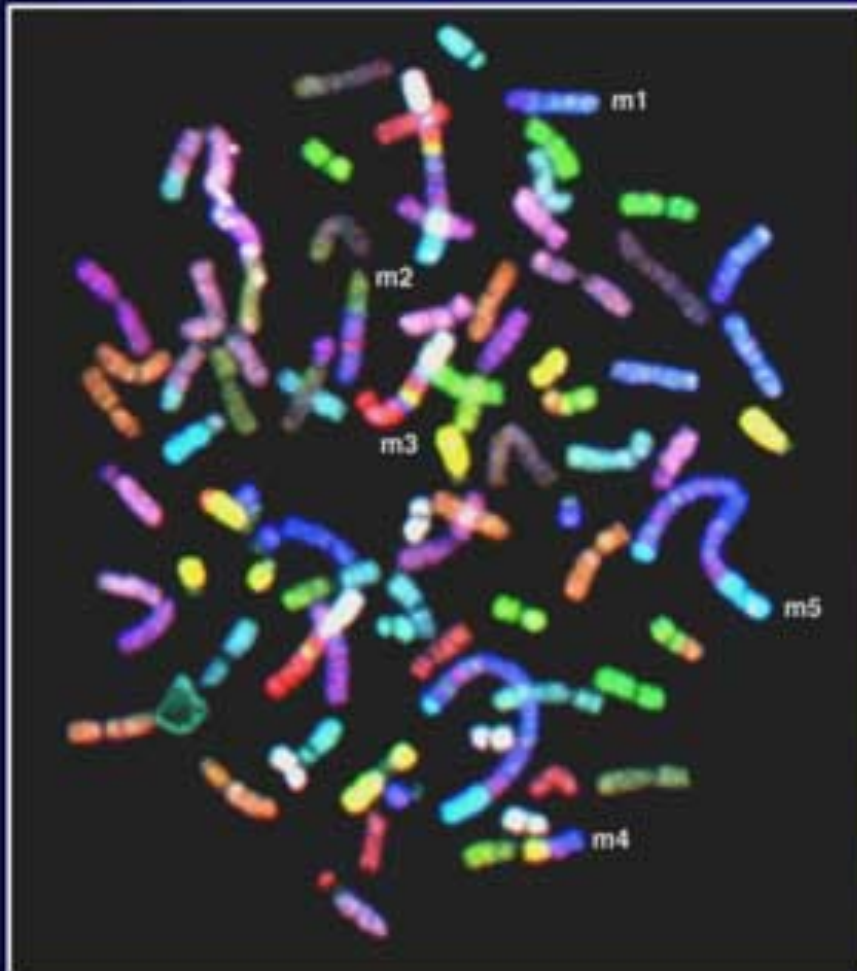
Leona Samson (Lectures)

Zachary Nagel (help with development) Alex Chaim



Large Deletions or Insertions

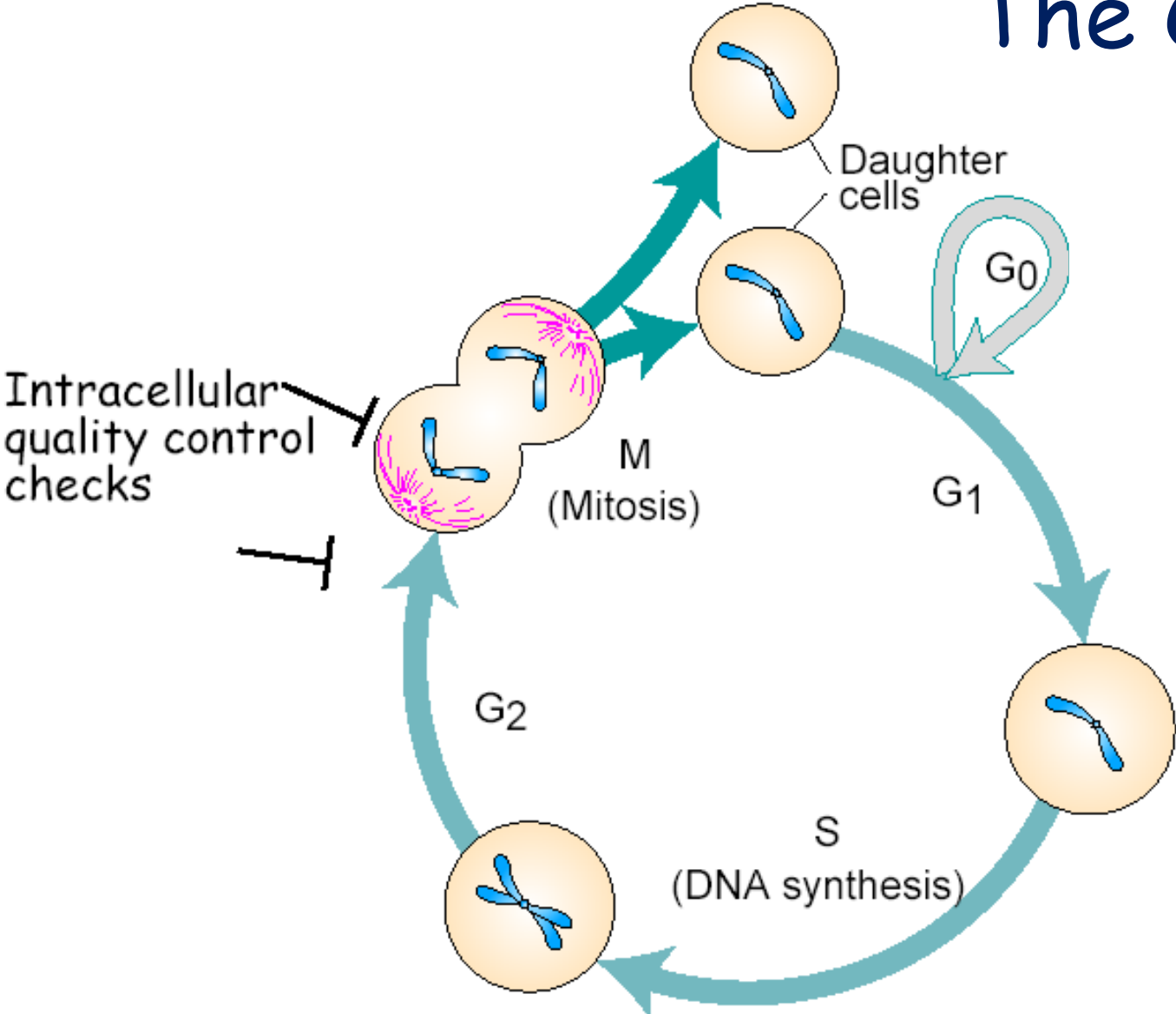
SKY chromosome painting: breast cancer



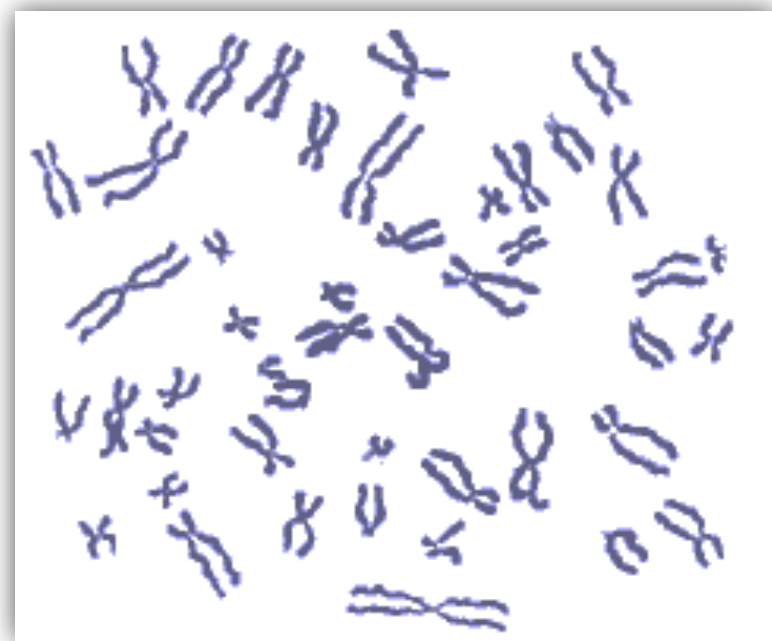
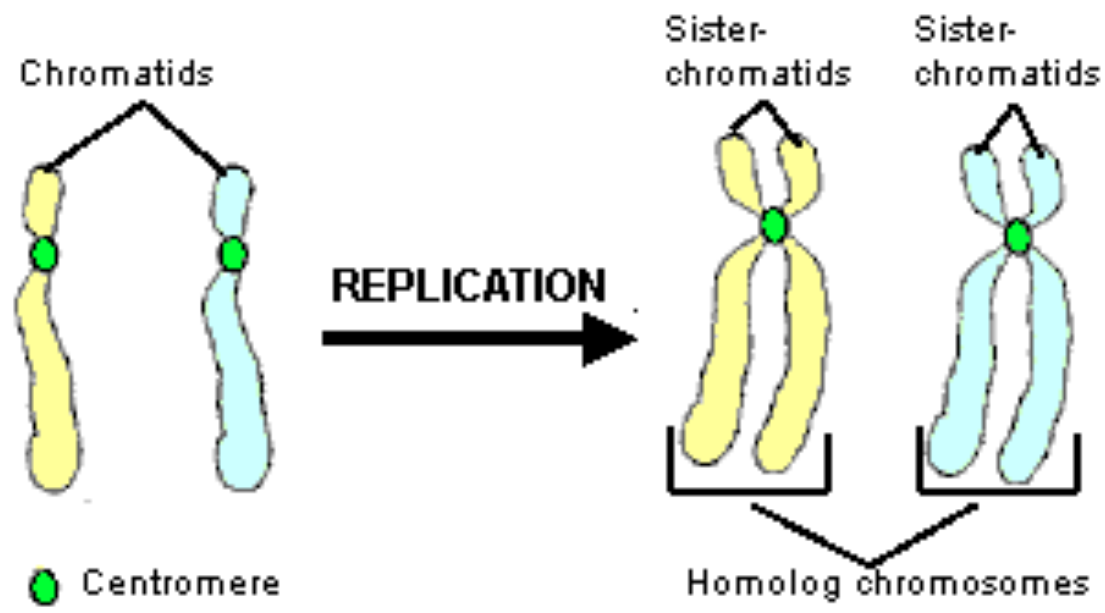
Normal SKY chromosomes are not multicolored.

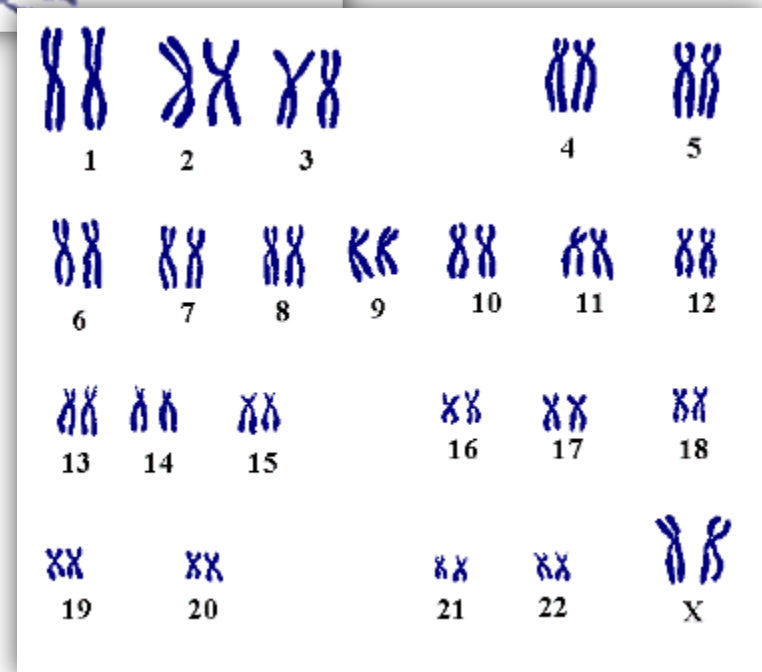
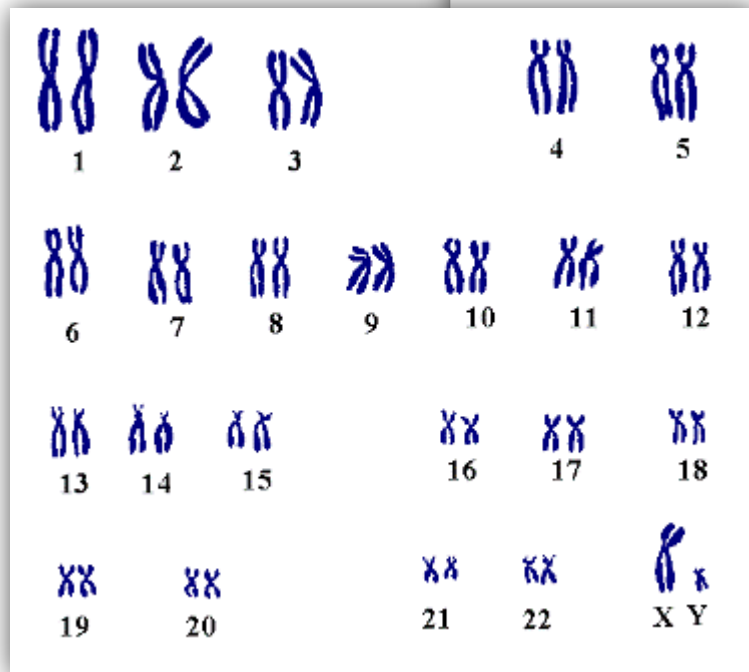
Chromosomes in breast cancer appear multicolored because they have exchanged genetic material.

The Cell Cycle

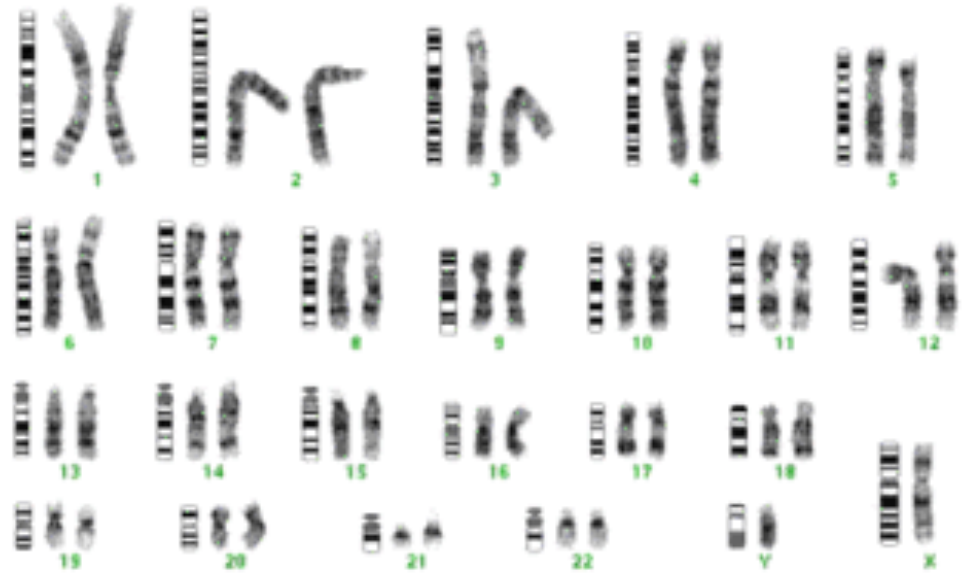
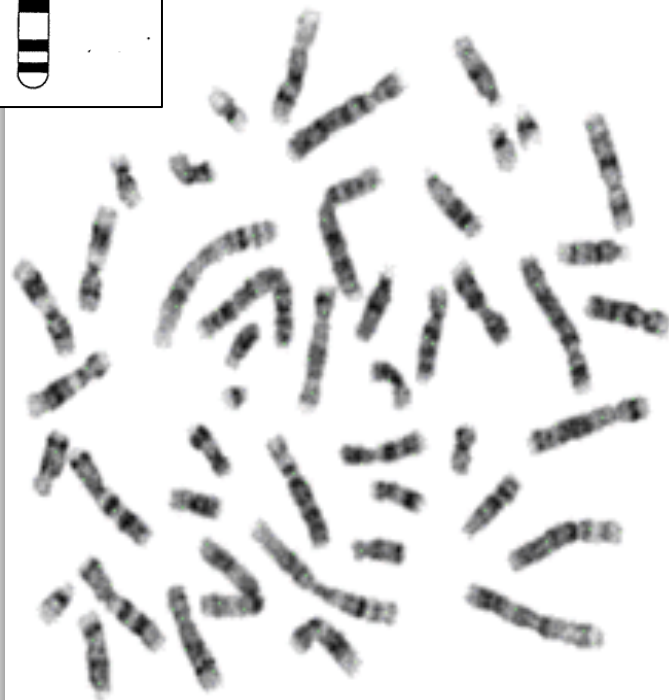
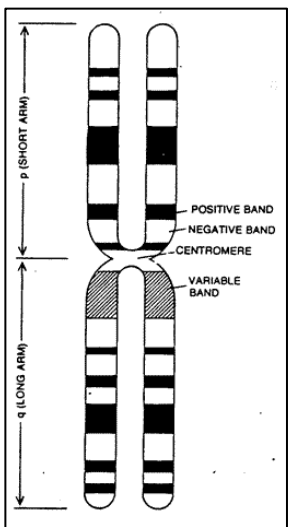


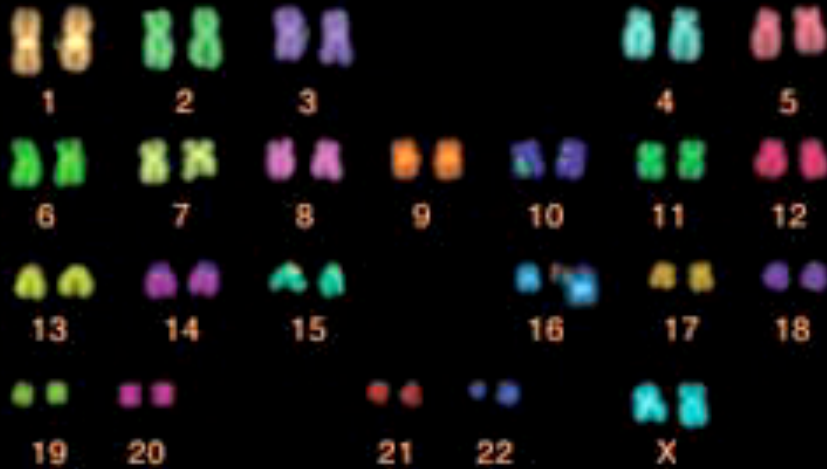
Duplication of chromosomes
DNA Replication





Chromosome Banding - Karyotyping





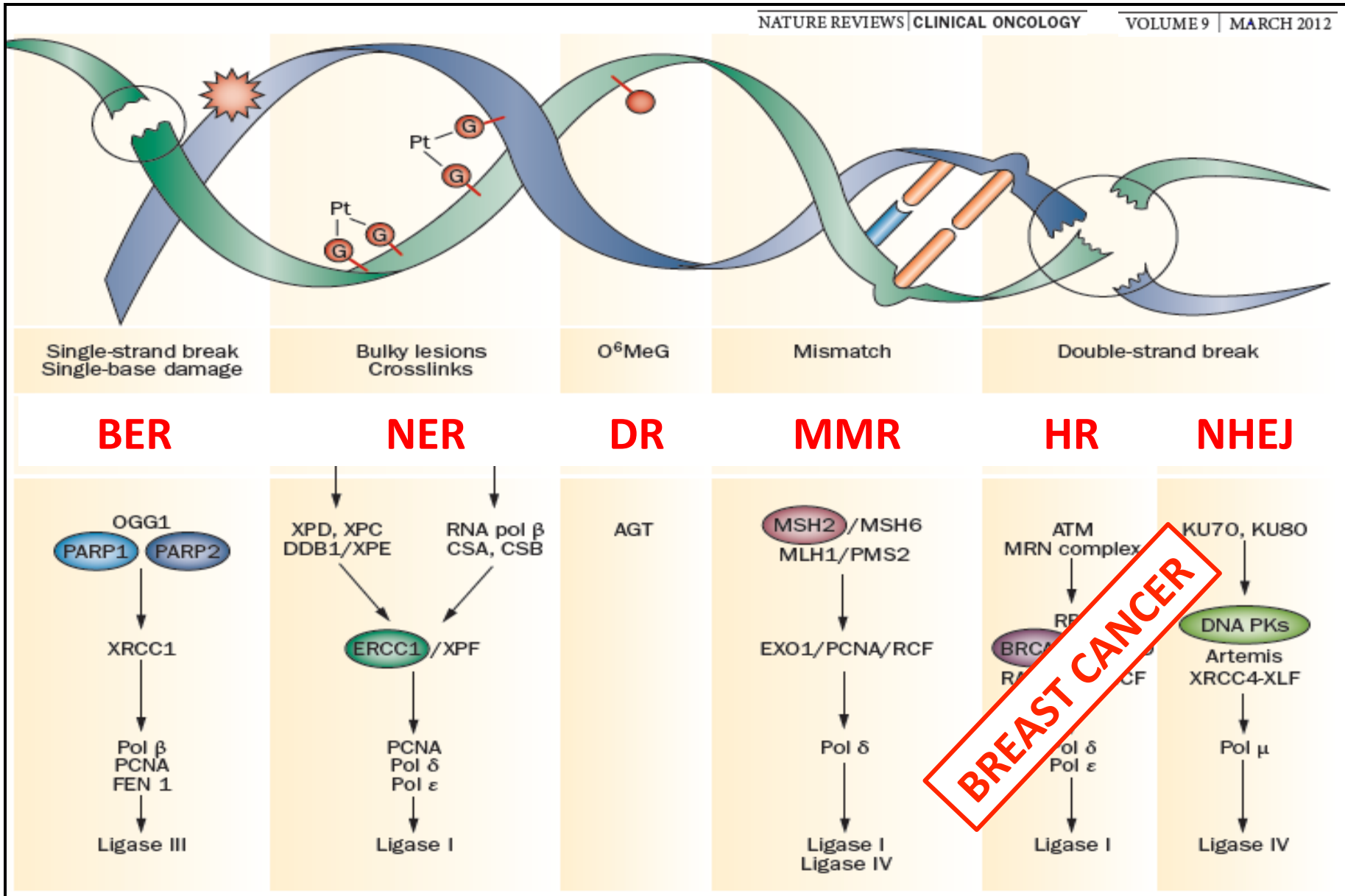
Chromosomes from a Normal cell



Chromosomes from a Tumor cell

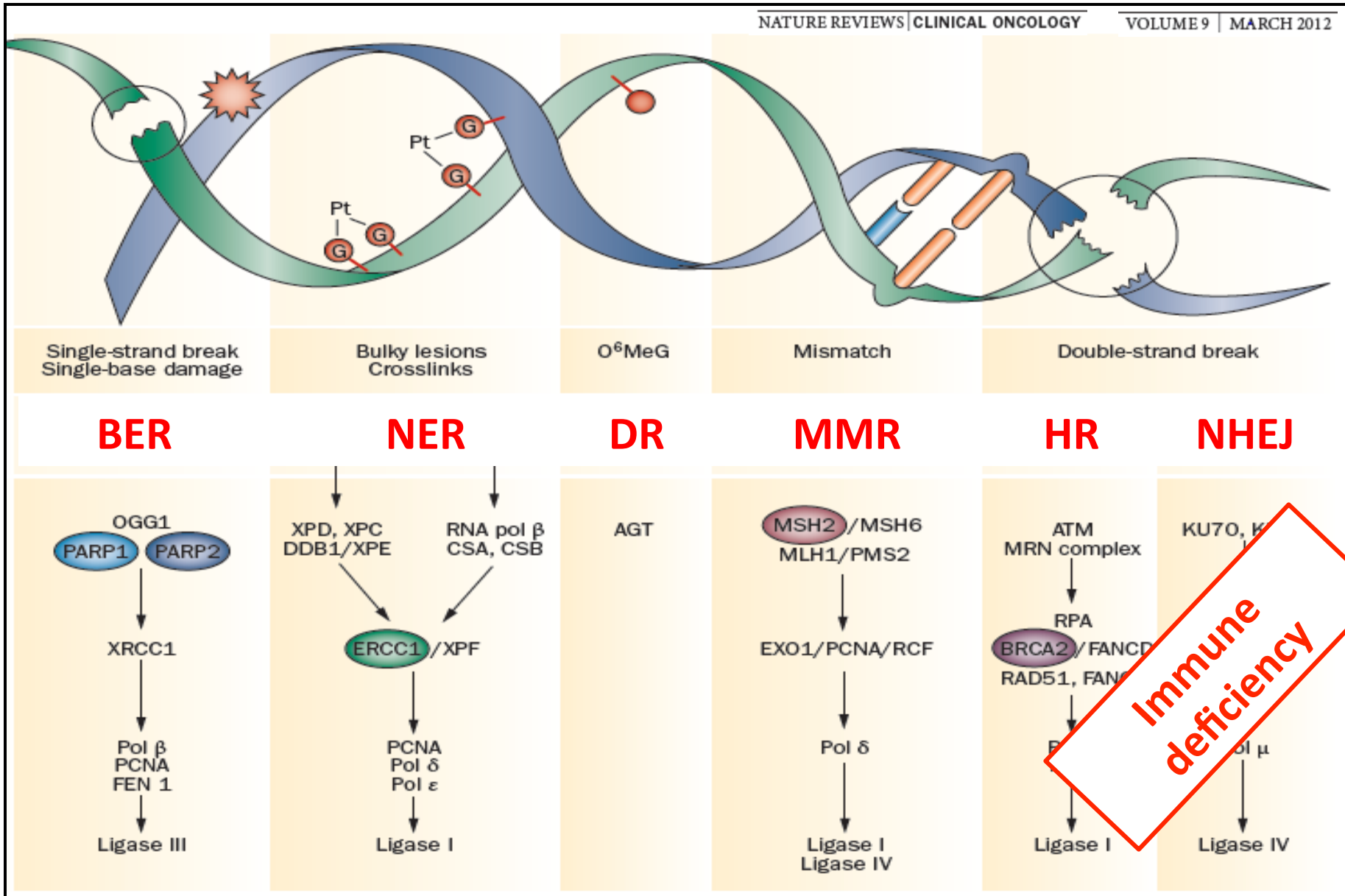
Spectral Karyotyping (SKY)
 "SKY Painted Chromosomes"

Six Major DNA Repair Pathways



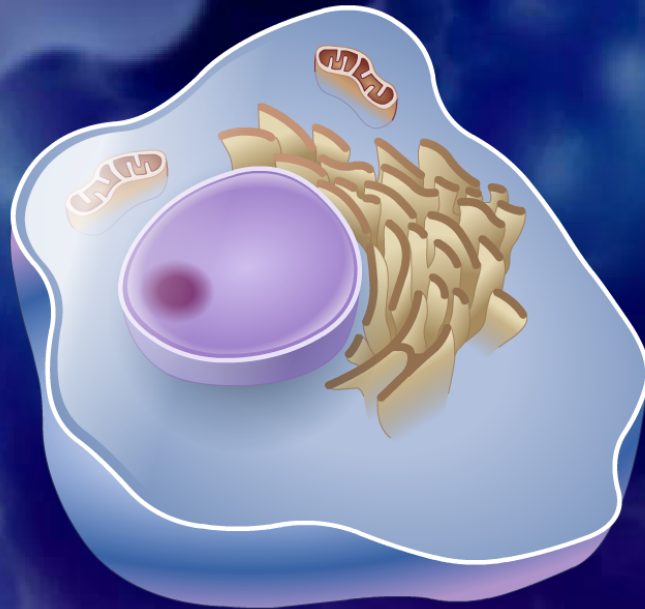
Six Major DNA Repair Pathways

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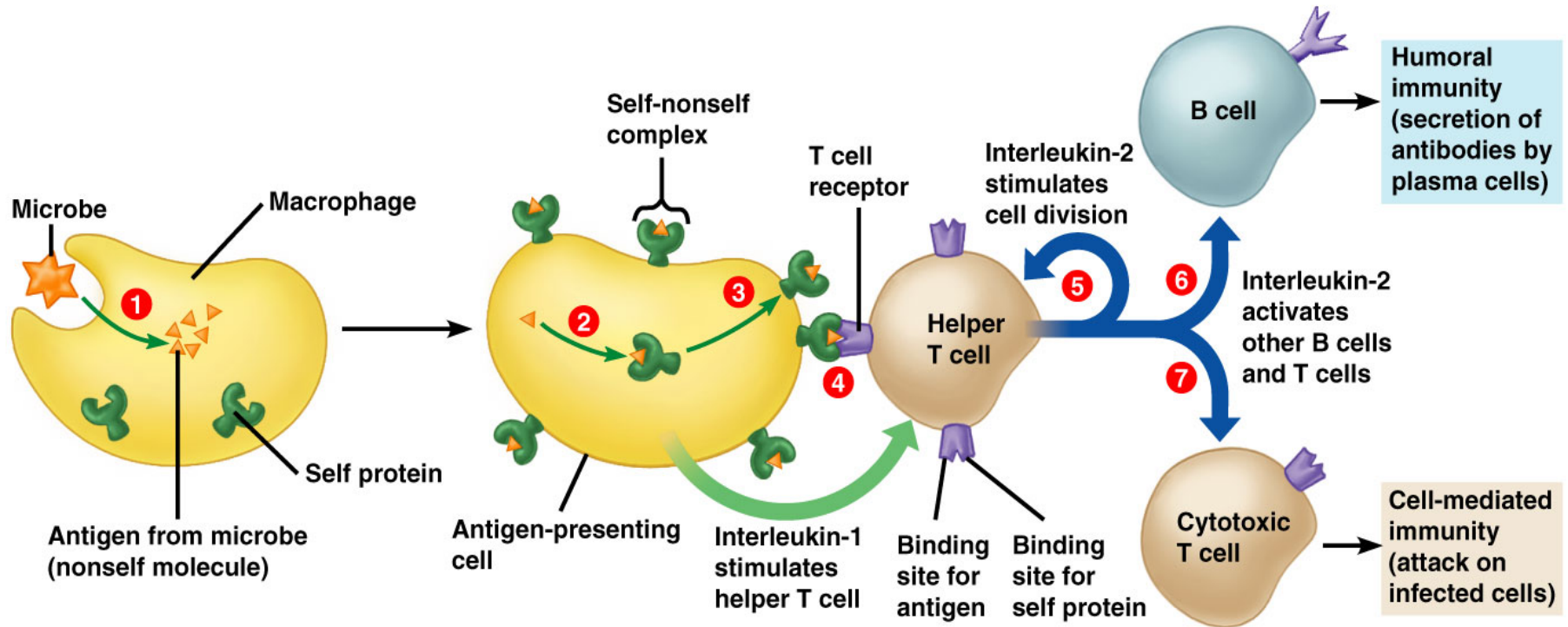
Non Homologous End Joining
is **REQUIRED** for a functional
immune system!

The Immune Response



Activation of the immune response typically begins when a pathogen enters the body. Macrophages that encounter the pathogen ingest, process and display the antigen fragments on their cell surfaces.

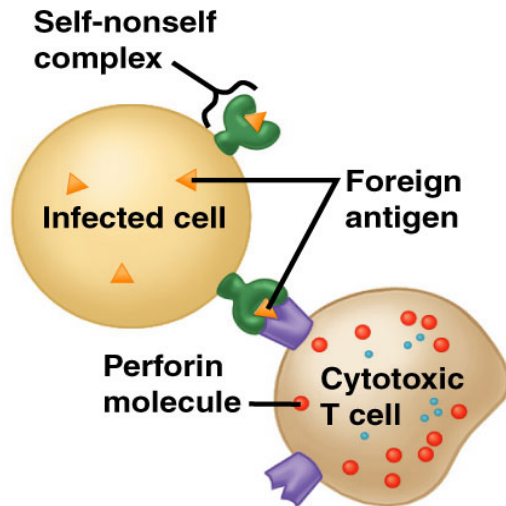
The body contains millions of different T-cells and B-cells, each able to respond to one specific antigen.



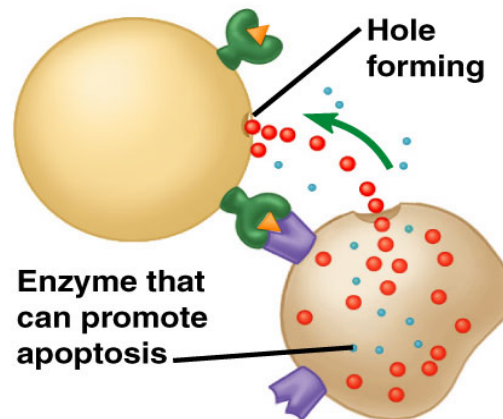
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The body contains millions of different T-cells and B-cells, each able to respond to one specific antigen.

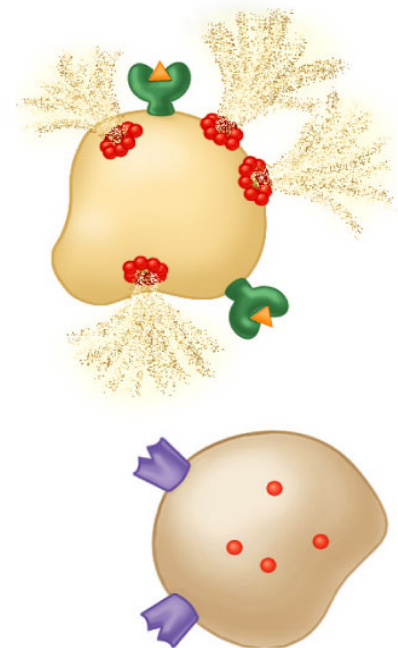
1 Cytotoxic T cell binds to infected cell



2 Perforin makes holes in infected cell's membrane and enzyme enters



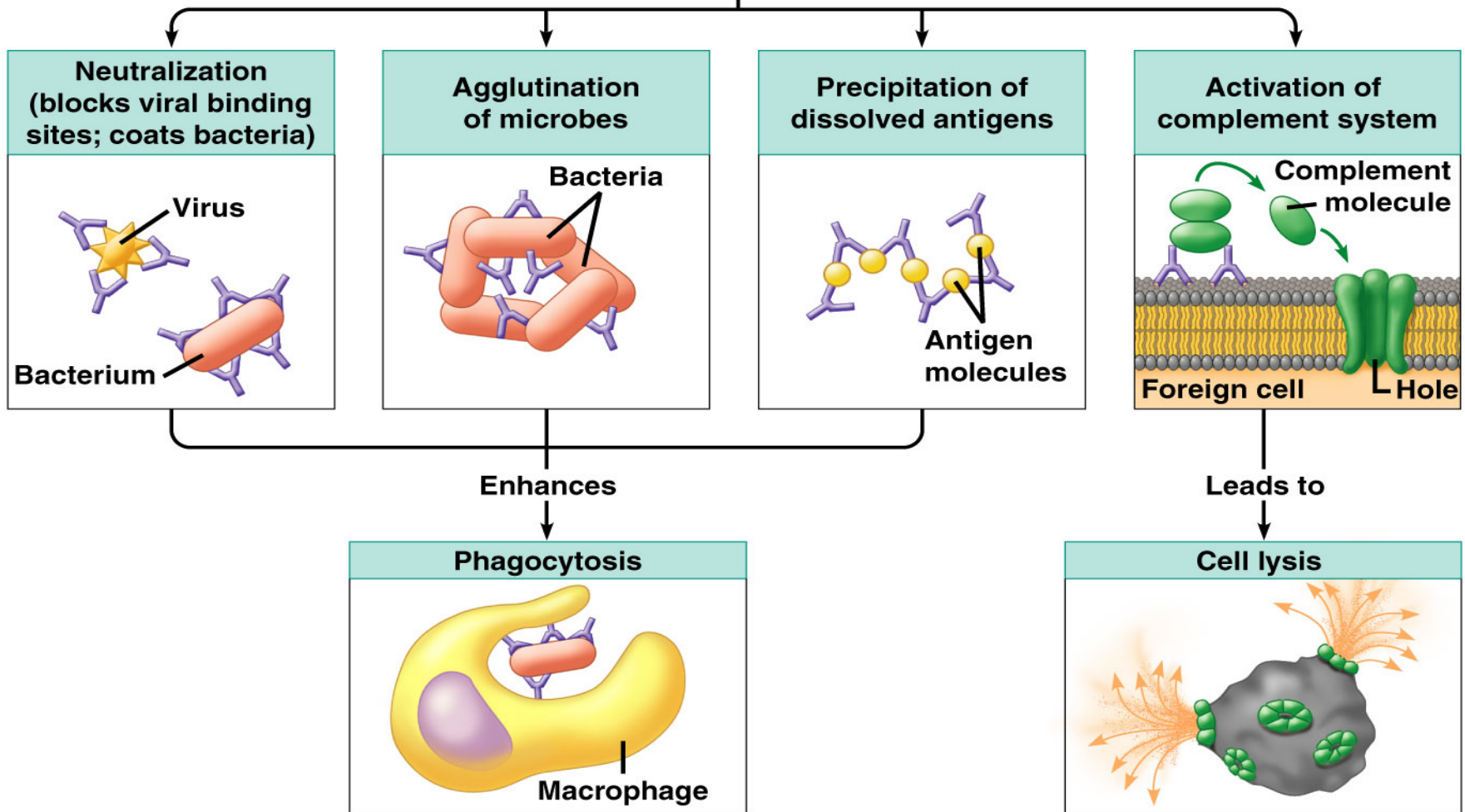
3 Infected cell is destroyed



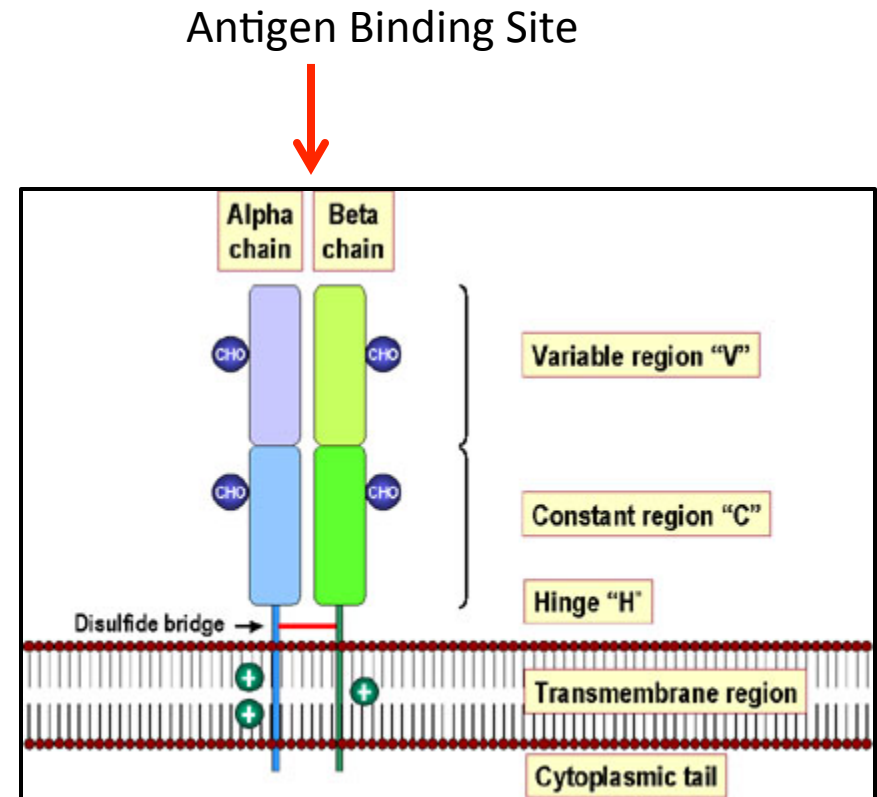
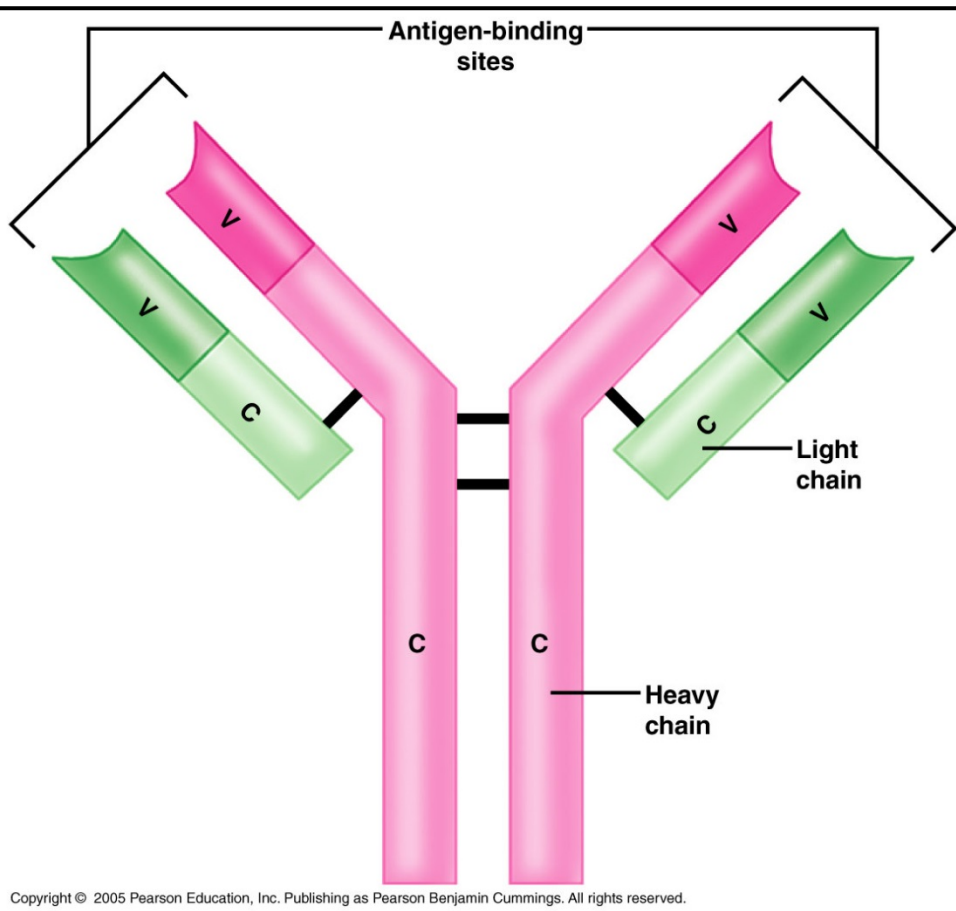
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Antibodies (secreted by B cells) work in several different ways

Binding of antibodies to antigens inactivates antigens by



"antigen" comes from **ANTI**-body **GEN**erating substances



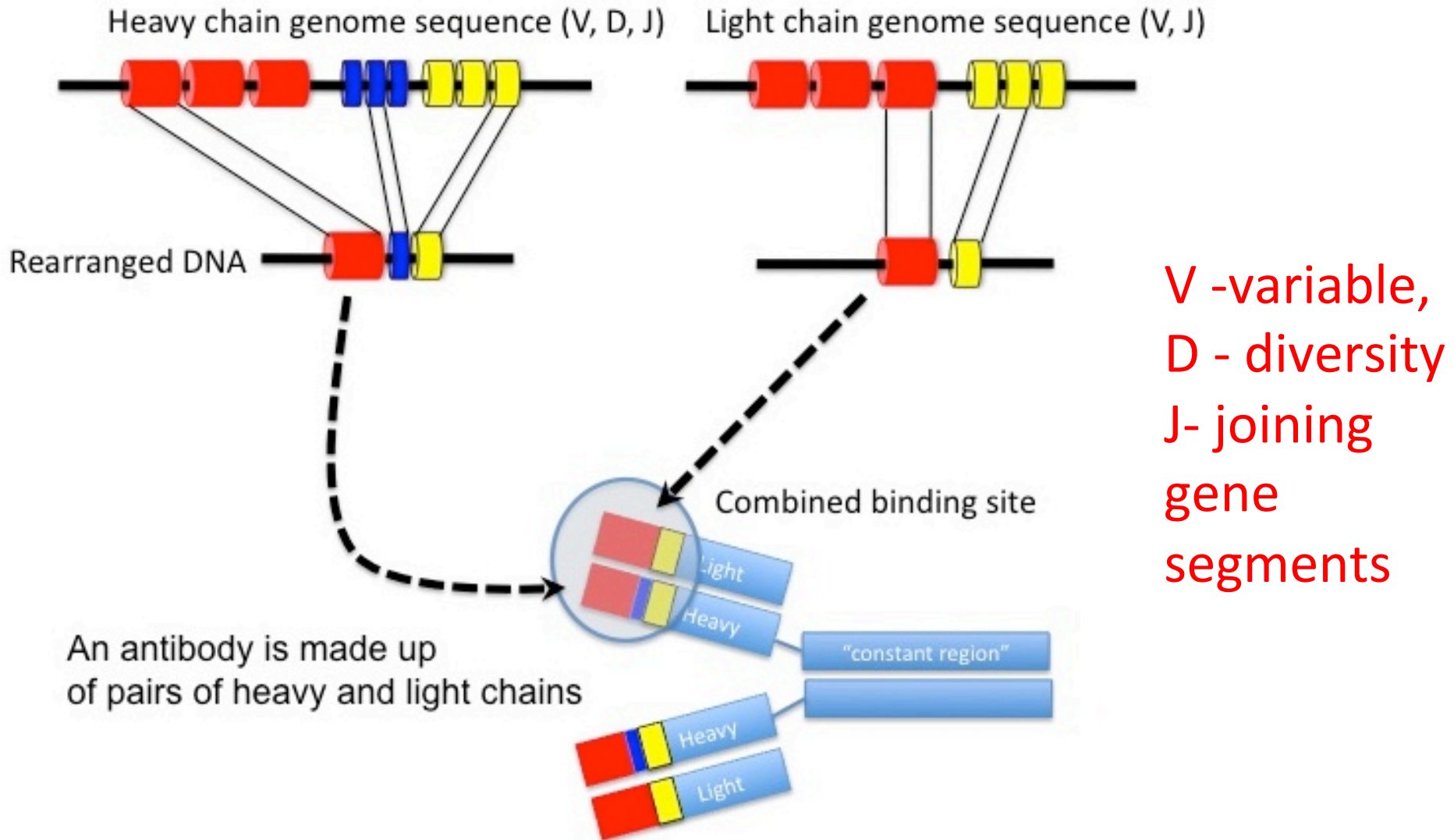
<http://www.austincc.edu/apreview/EmphasisItems/Inflammatoryresponse.html#ANTIB>

<http://pathmicro.med.sc.edu/bowers/mhc.htm>

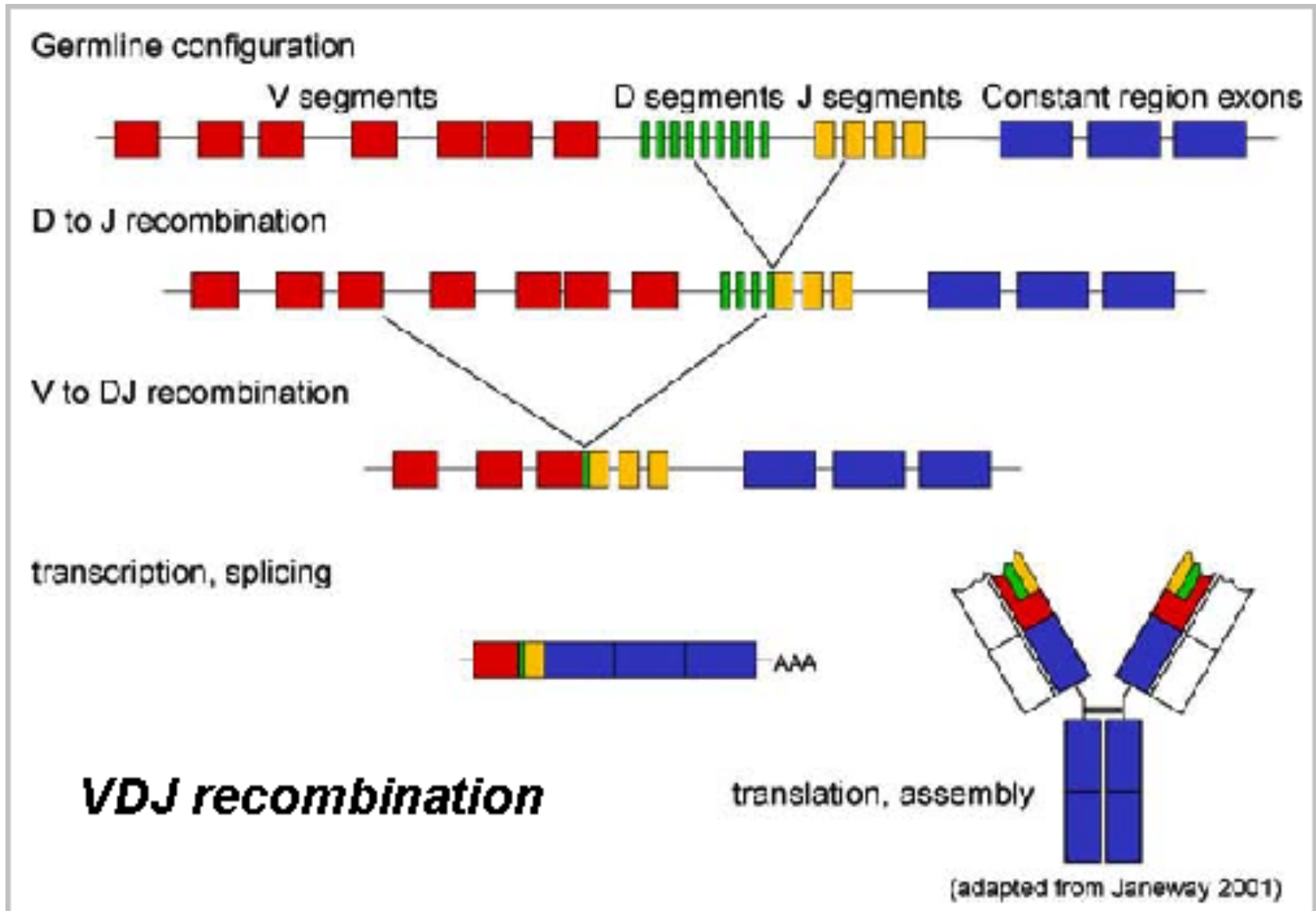
B-cell Immunoglobulin

T-cell Receptor

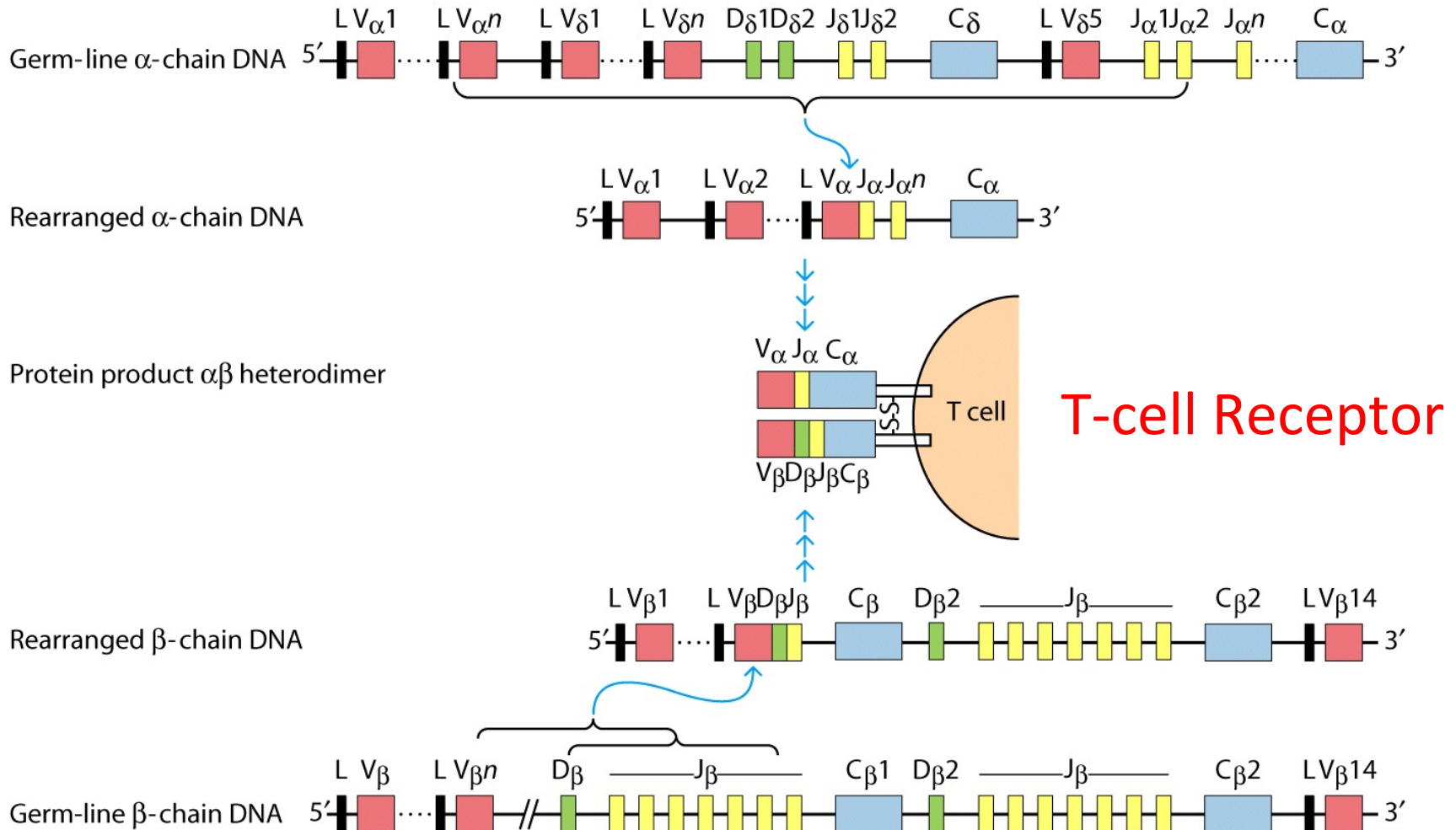
How Do the Variable Regions become Variable? Through Programmed NHEJ!!



How Do the Variable Regions become Variable? Through Programmed NHEJ!!



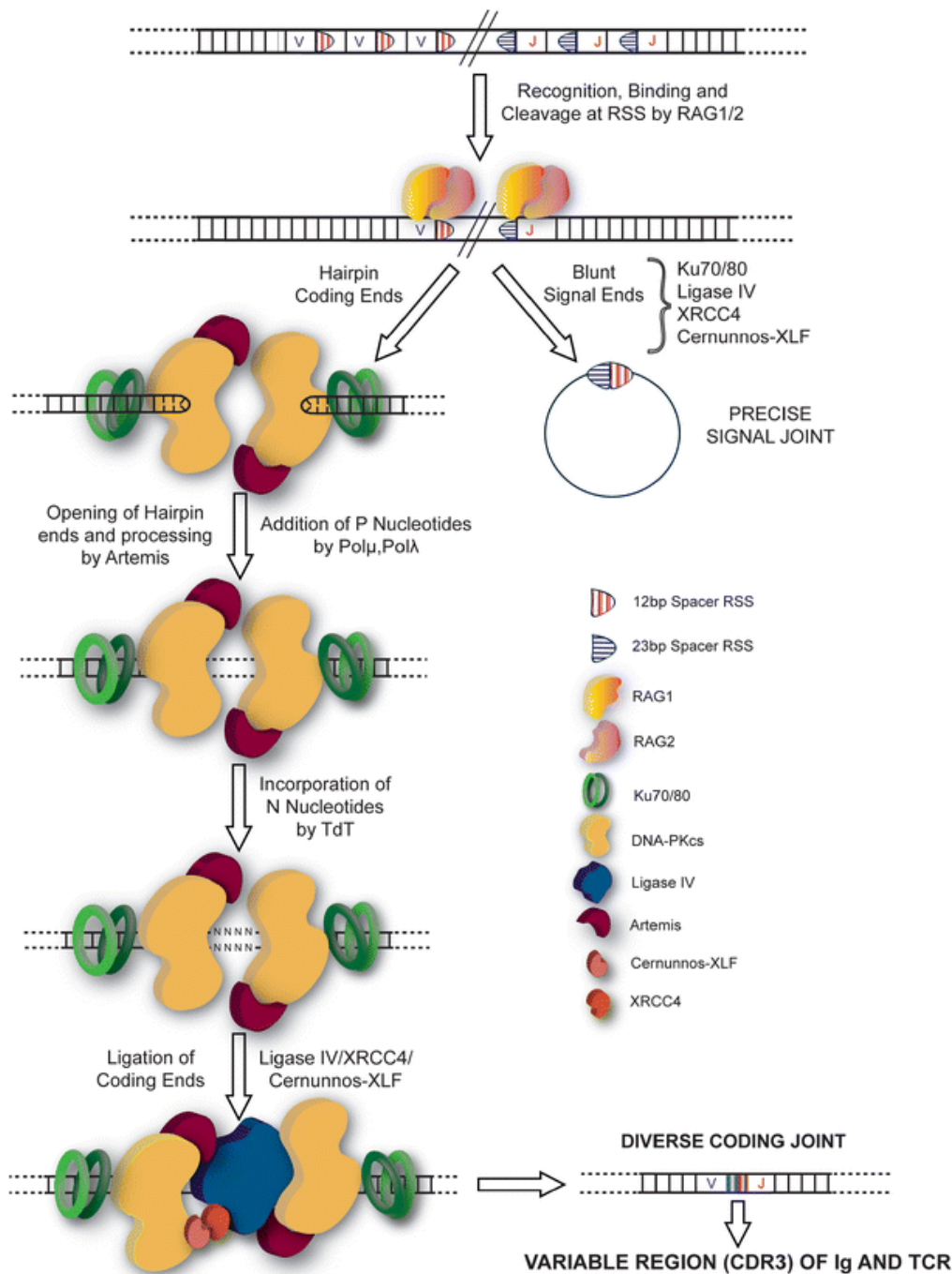
How Do the Variable Regions become Variable? Through Programmed NHEJ!!



V(D)J Gene Recombination

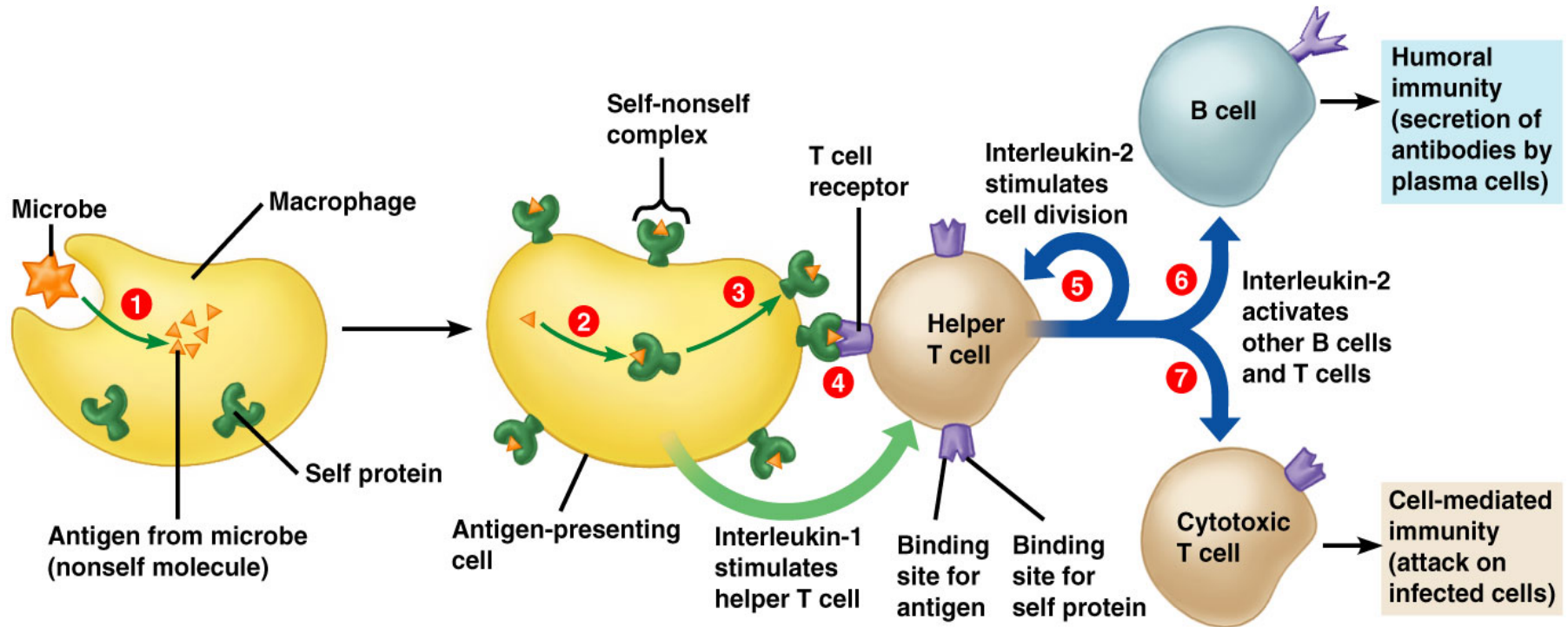
<http://www.youtube.com/watch?v=QTOBSFJWogE>

How Do the Variable Regions become Variable? Through NHEJ mediated DNA Recombination!



The rearrangement starts with the binding of products from recombination activating genes RAG1 and RAG2, whose expression is **unique to lymphoid progenitor cells**

The body contains **millions of different T-cells and B-cells**, each able to respond to one specific antigen.



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How Variable is Variable?

| Number of functional gene segments in human immunoglobulin loci | | | |
|--|--------------|-----------|-------------|
| Segment | light chains | | heavy chain |
| | κ | λ | H |
| Variable (V) | 40 | 30 | 65 |
| Diversity (D) | 0 | 0 | 27 |
| Joining (J) | 5 | 4 | 6 |

Over **15,000,000** combinations of variable, diversity and joining, V(D)J, gene segments are possible.

Imprecise recombination and mutation increase the variability into **billions of possible combinations**.

How Variable is Variable?

| | T cell receptor | |
|---------------------------------------|-----------------|---------|
| | α | β |
| Number of V gene segments | 54 | 67 |
| Number of diversity (D) gene segments | 0 | 2 |
| Number of joining (J) gene segments | 61 | 4 |

Millions of combinations of variable, diversity and joining gene segments are possible.

Imprecise recombination and mutation increase the variability into **billions of possible combinations**.

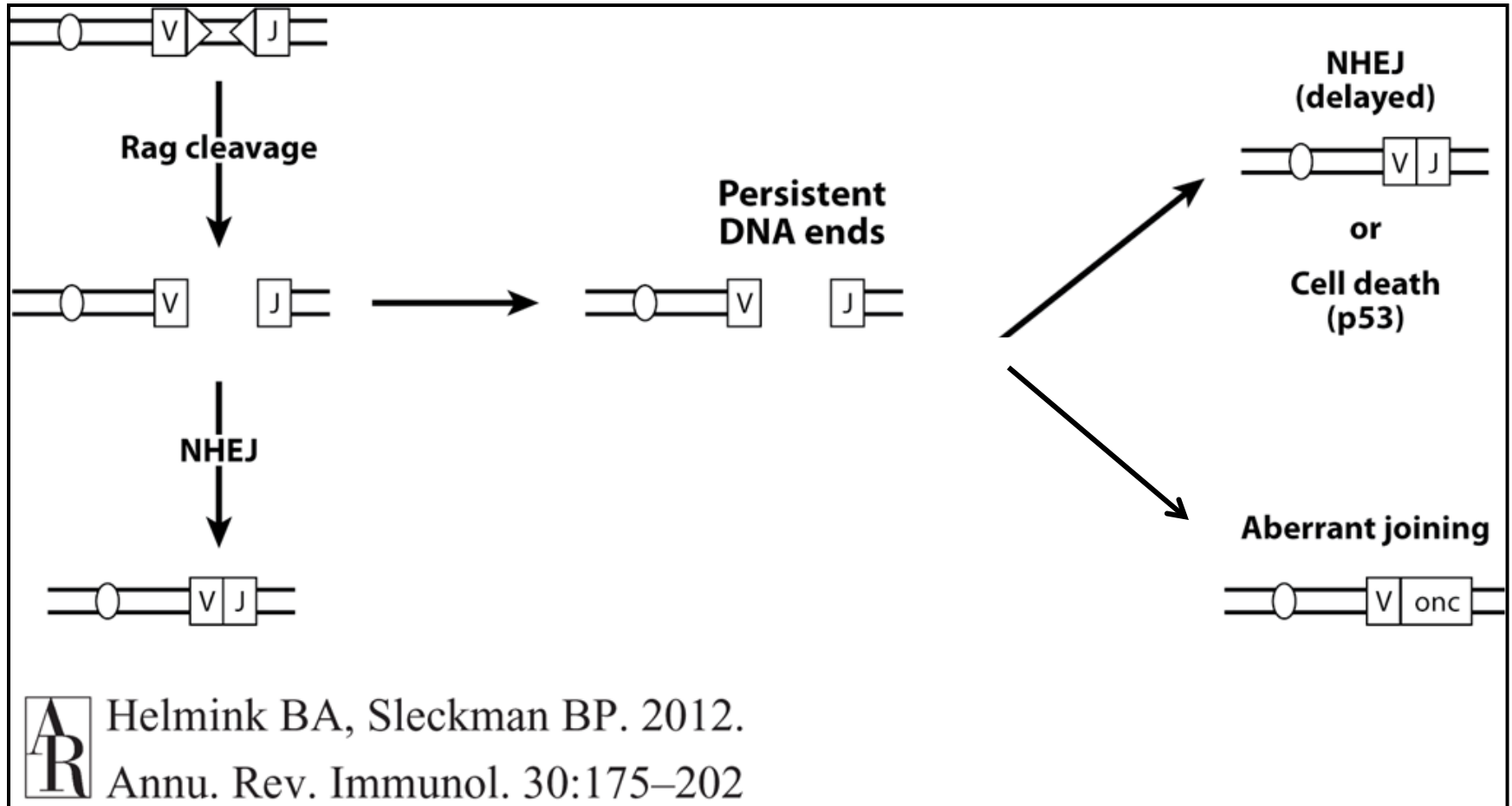
What happens if mice or people lose NHEJ capacity?

What happens if mice or people lose NHEJ capacity?

| NHEJ gene | Mouse knockout phenotype | Patient phenotype |
|--------------------------------------|---|--|
| <i>XRCC6</i> (encoding Ku70) | Viable, SCID, small size, radiosensitivity and thymoma ^{50,51} | None known |
| <i>XRCC5</i> (encoding Ku80) | Viable, SCID, small size, radiosensitivity, genomic instability and tumours, especially with p53 deletion ^{47,52-54} | None known |
| <i>PRKDC</i> (encoding DNA-PKcs) | Viable, SCID, some genomic instability and tumours with p53 (REFS 55-57) | Human hypomorph has SCID and radiosensitivity ⁵⁸ |
| <i>DCLRE1C</i> (encoding Artemis) | Viable, SCID, radiosensitivity and genomic instability ⁵⁹ | Null results in SCID and radiosensitivity; hypomorph shows reduction in lymphocytes, genomic instability and lymphoma ^{60,61} |
| <i>NHEJ1</i> (encoding XLF) | Mild lymphocytopaenia and radiosensitivity ⁶² | Cernunnos syndrome; immunodeficiency, developmental delay, microcephaly, reduced growth and genomic instability ⁶³ |
| <i>XRCC4</i> | Null is lethal with neuronal apoptosis; rescue with p53 results in SCID, radiosensitivity, early B lymphoma and genomic instability ^{49,64} | None known |
| <i>LIG4</i> | Knockout is lethal with neuronal apoptosis; rescue with p53 results in pro-B lymphoma and radiosensitivity; hypomorph is small, lymphopaenic and has reduced haematopoietic stem cell function ^{65,66} | LIG4 syndrome; immunodeficiency, reduced growth, developmental issues, microcephaly and malignancy ^{67,68} |

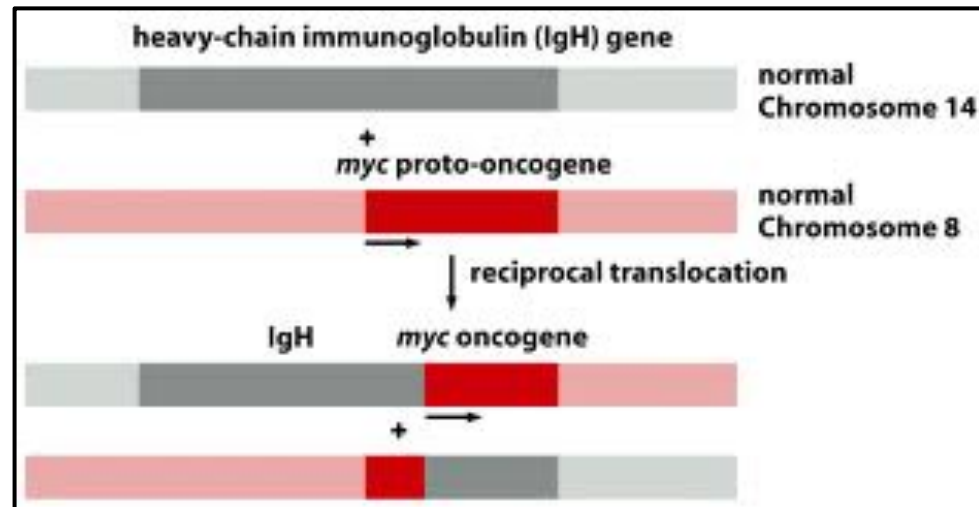
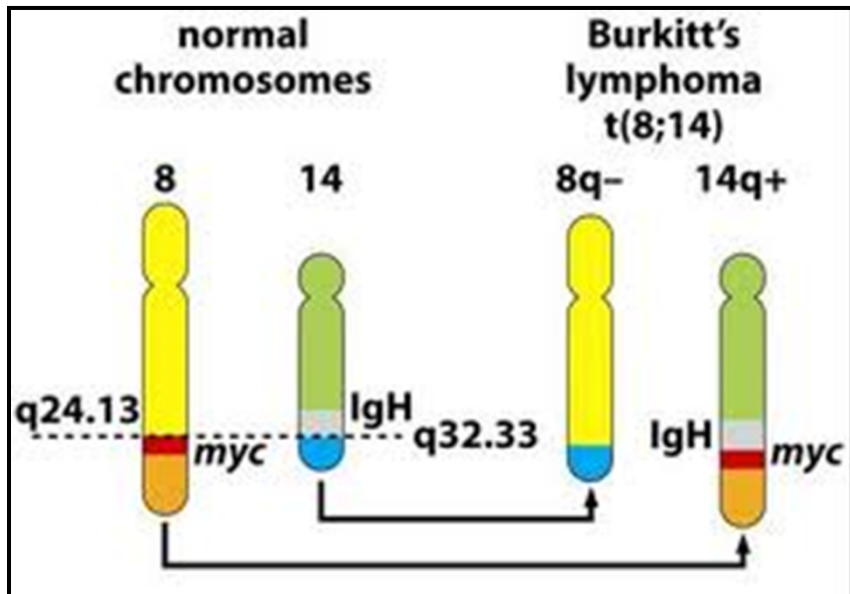
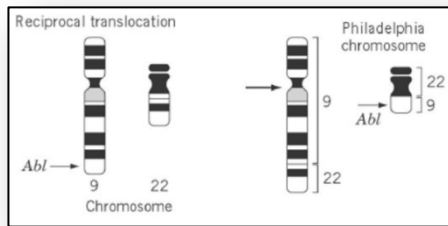
DCLRE1C, DNA cross-link repair 1C; DNA-PKcs, DNA-dependent protein kinase catalytic subunit; LIG4, DNA ligase 4; NHEJ, non-homologous end-joining; NHEJ1, NHEJ factor 1; PRKDC, protein kinase, DNA-activated, catalytic polypeptide; SCID, severe combined immunodeficiency; XLF, XRCC4-like factor; XRCC, X-ray repair cross-complementing protein.

Can V(D)J Recombination Go Wrong?

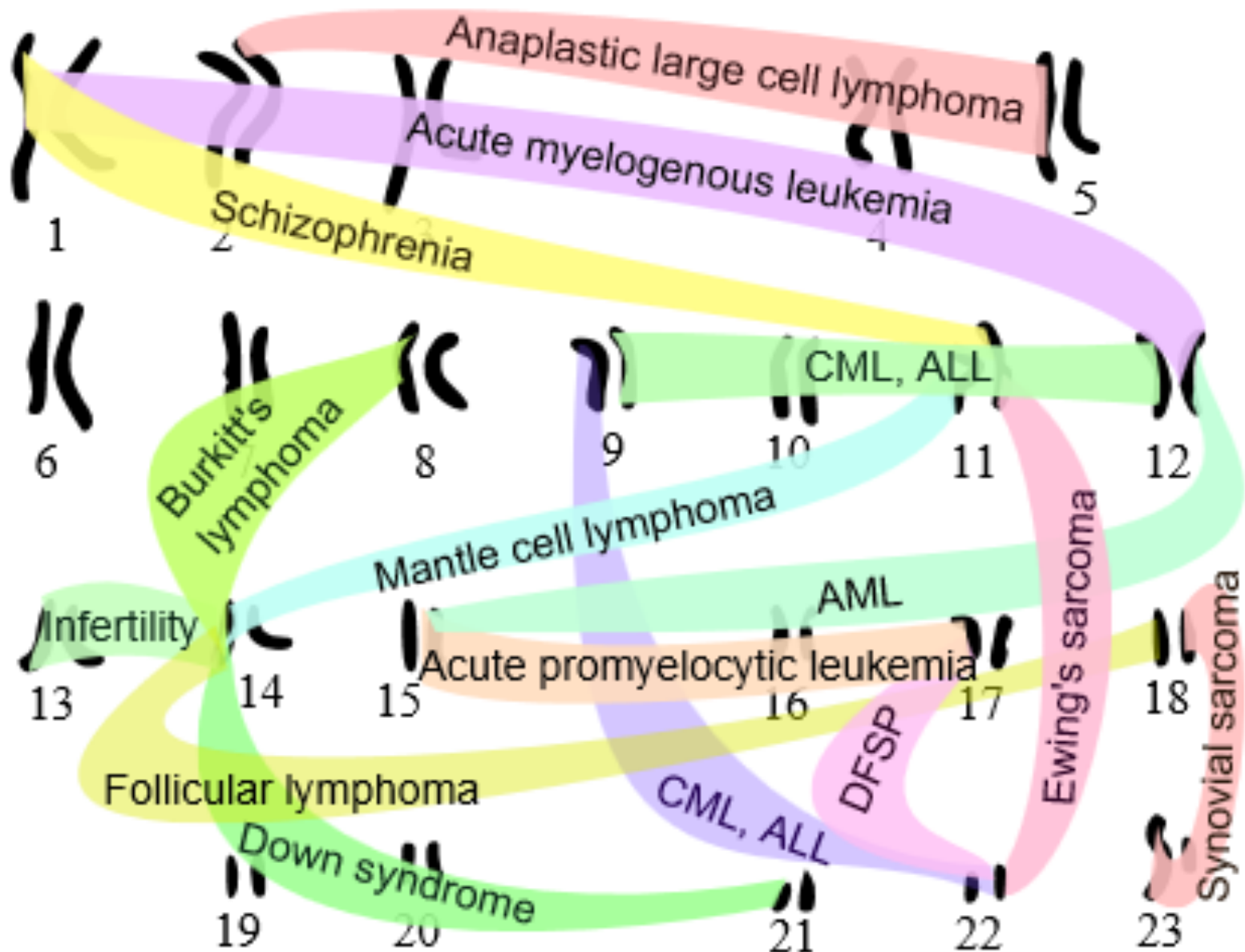


BURKITT'S LYMPHOMA

B-cell Lymphoma

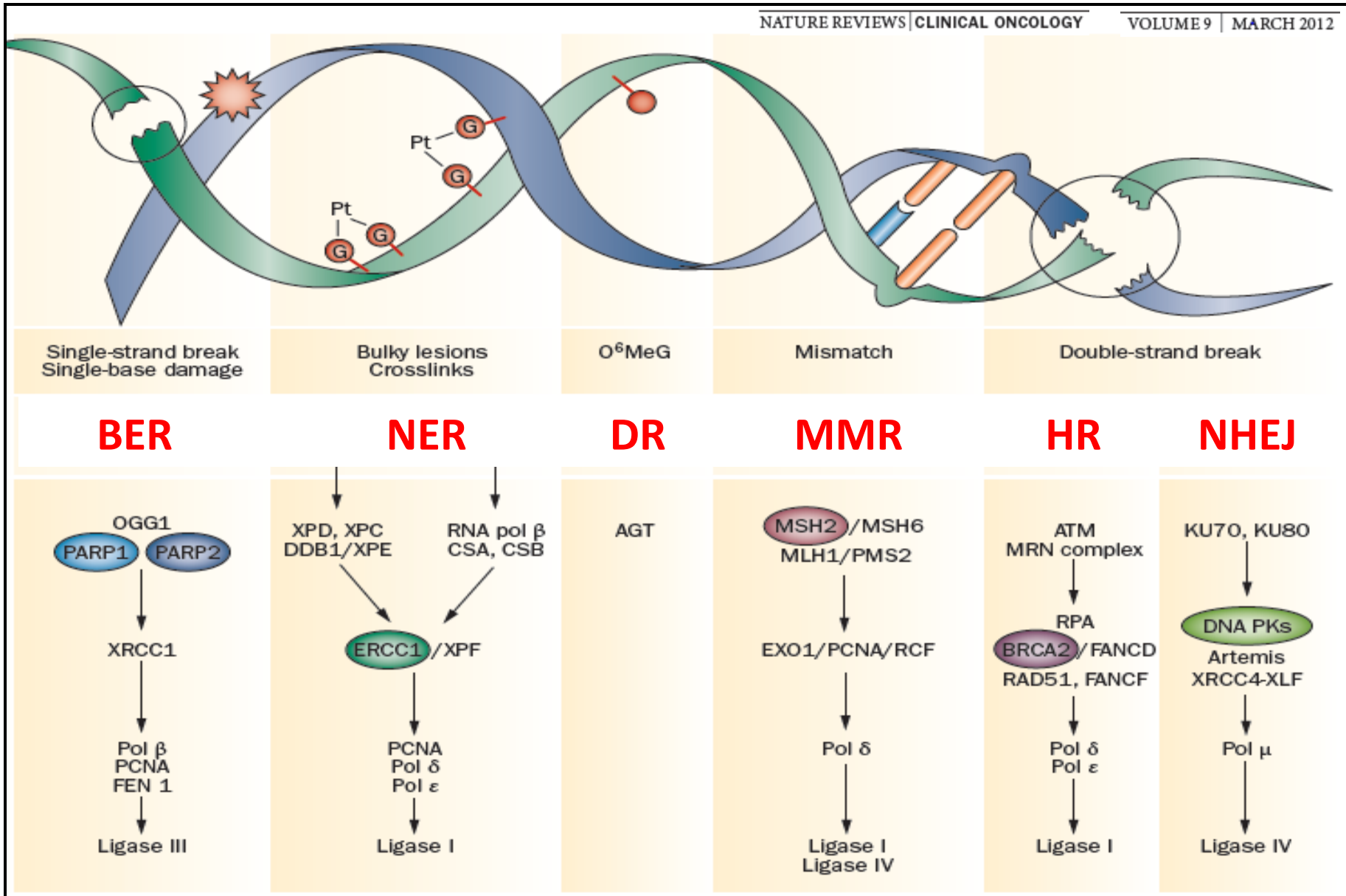


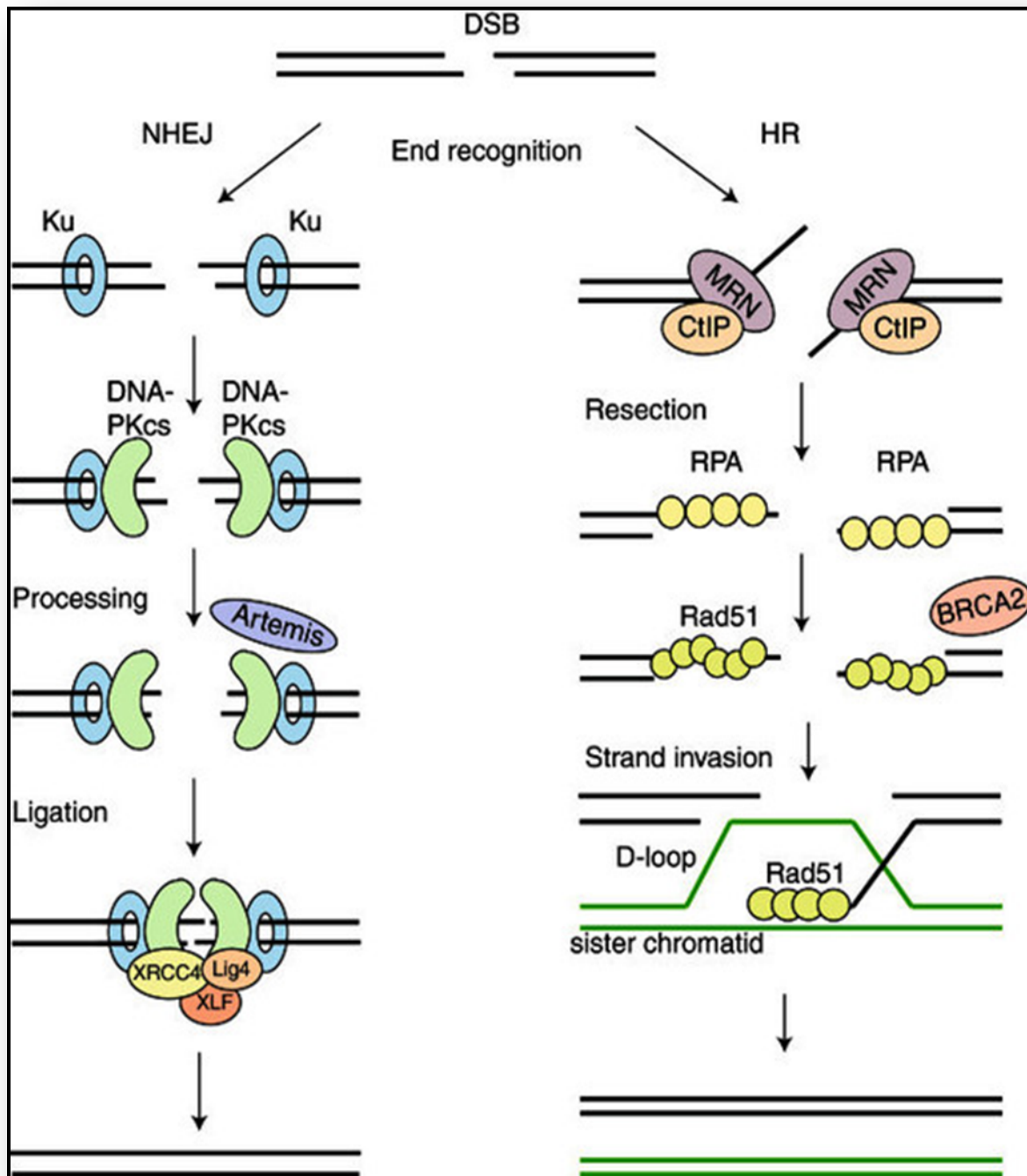
Diseases that involve Chromosome Translocations



Six Major DNA Repair Pathways

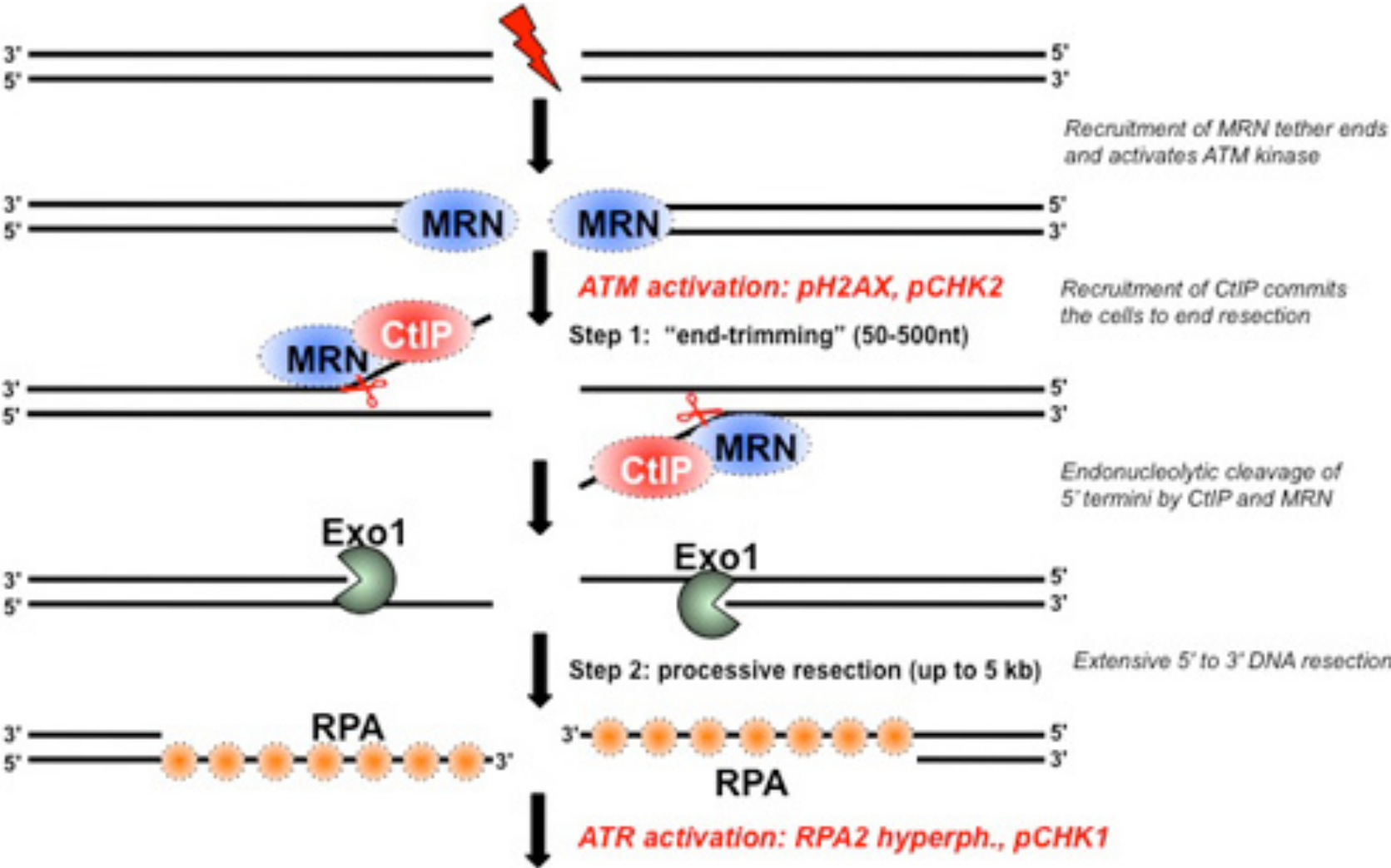
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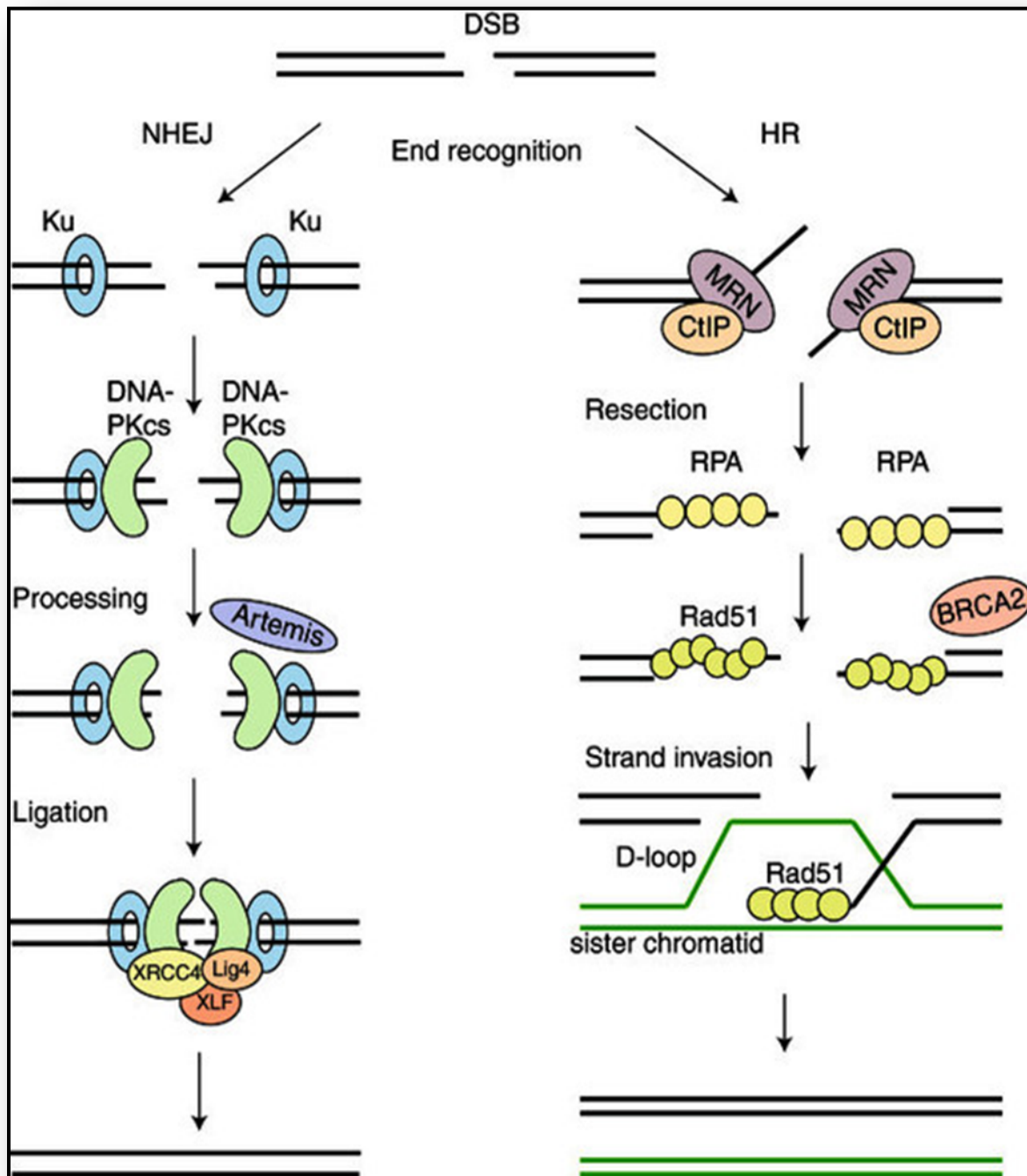


DNA Double
Strand
Break (DSB)
Repair –
What
Determines
Pathway
Choice?

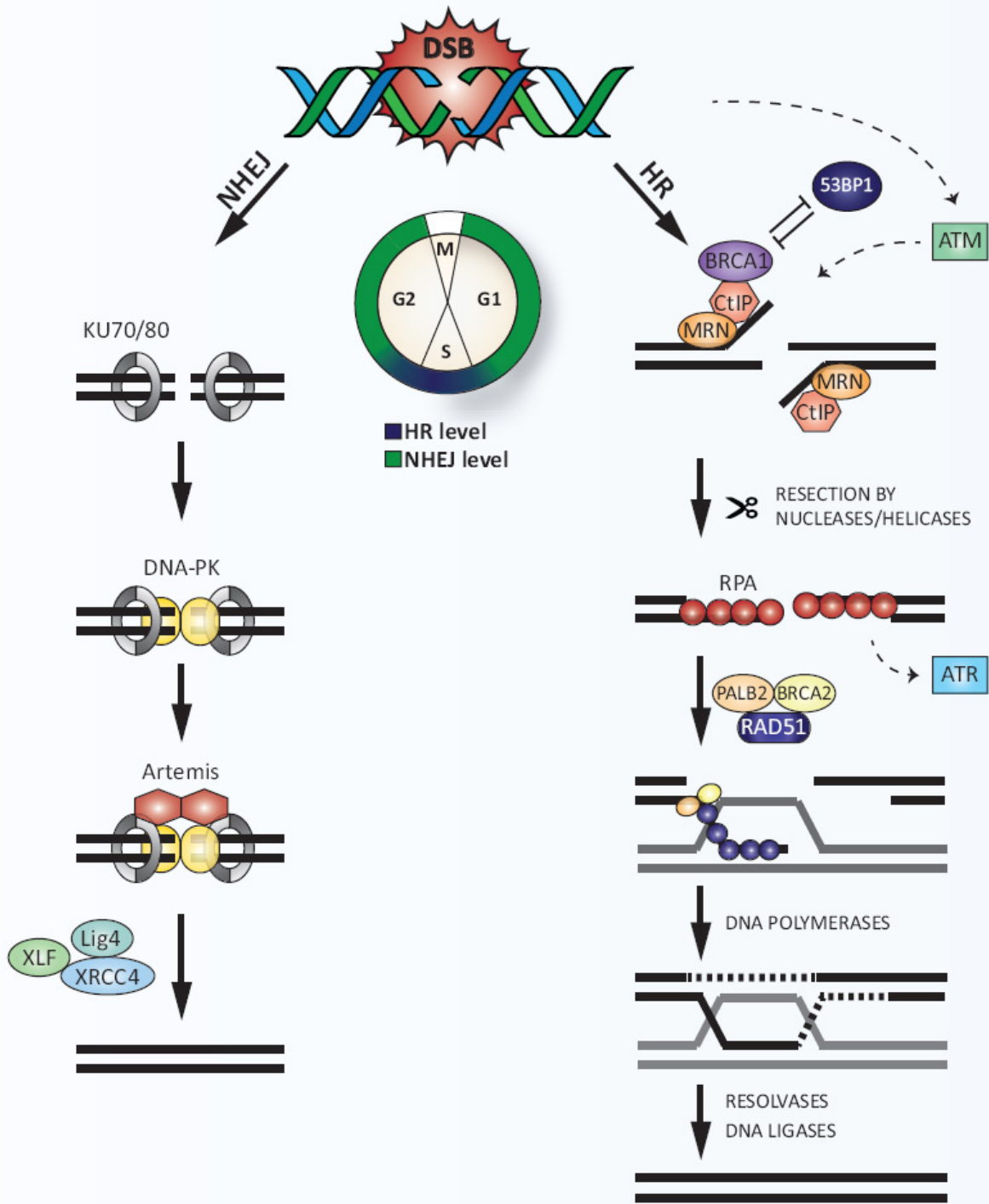
The MRN complex (Mre11/Rad50/Nbs1) competes with Ku for binding DSBs



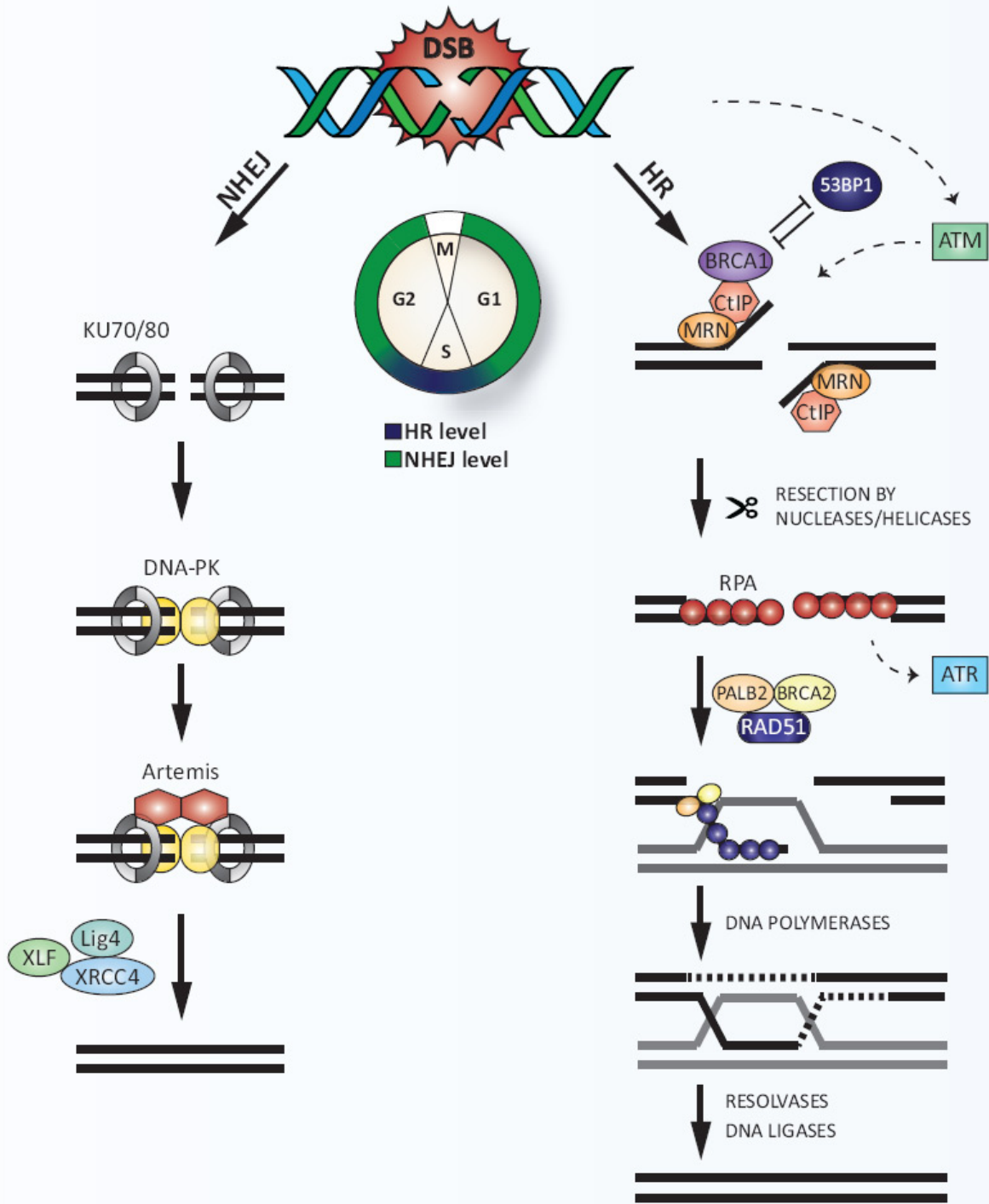
RAD51 nucleoprotein filaments → HR



DNA Double
Strand
Break (DSB)
Repair –
What
Determines
Pathway
Choice?

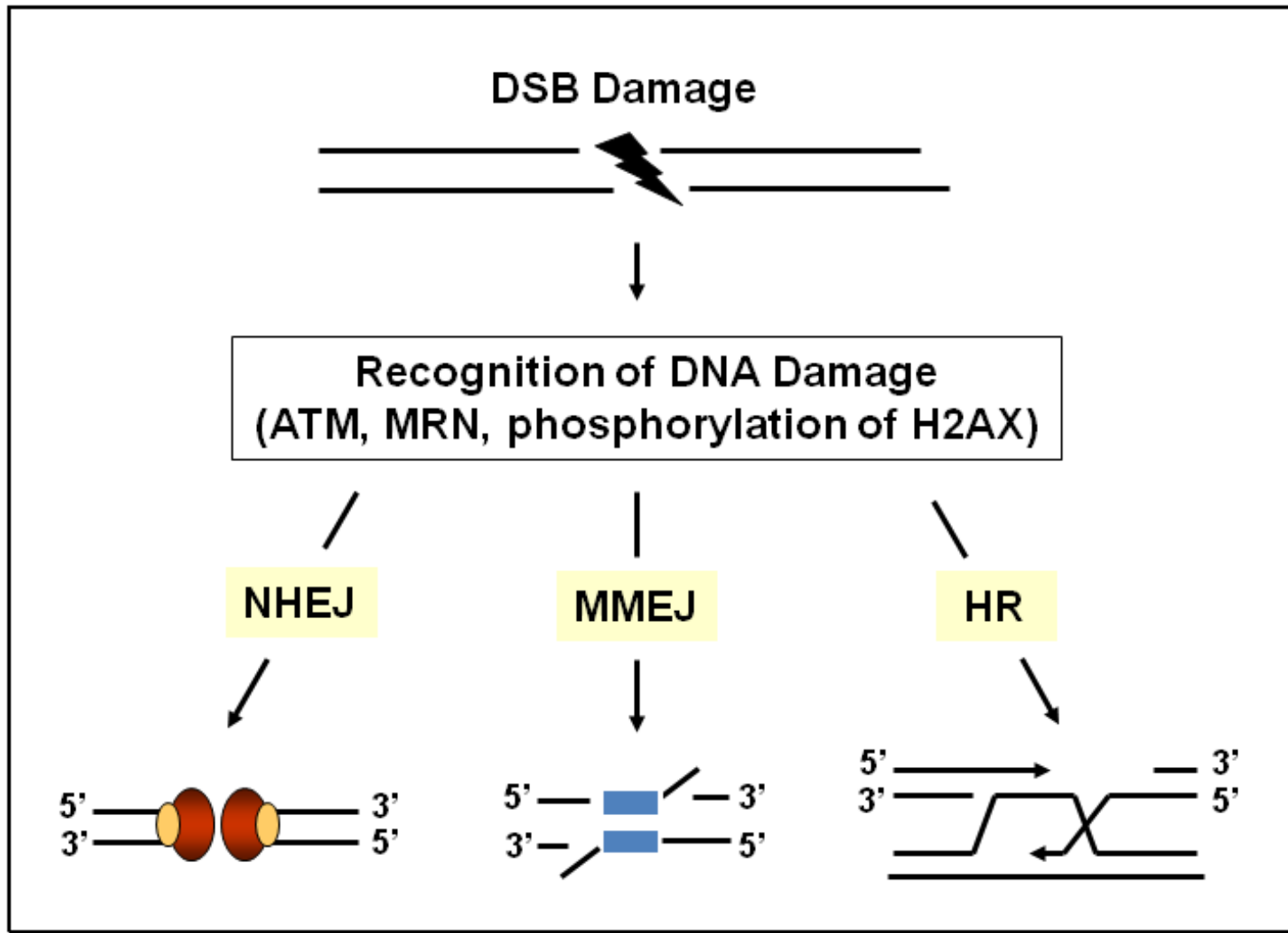


DNA Double
Strand
Break (DSB)
Repair –
What
Determines
Pathway
Choice?



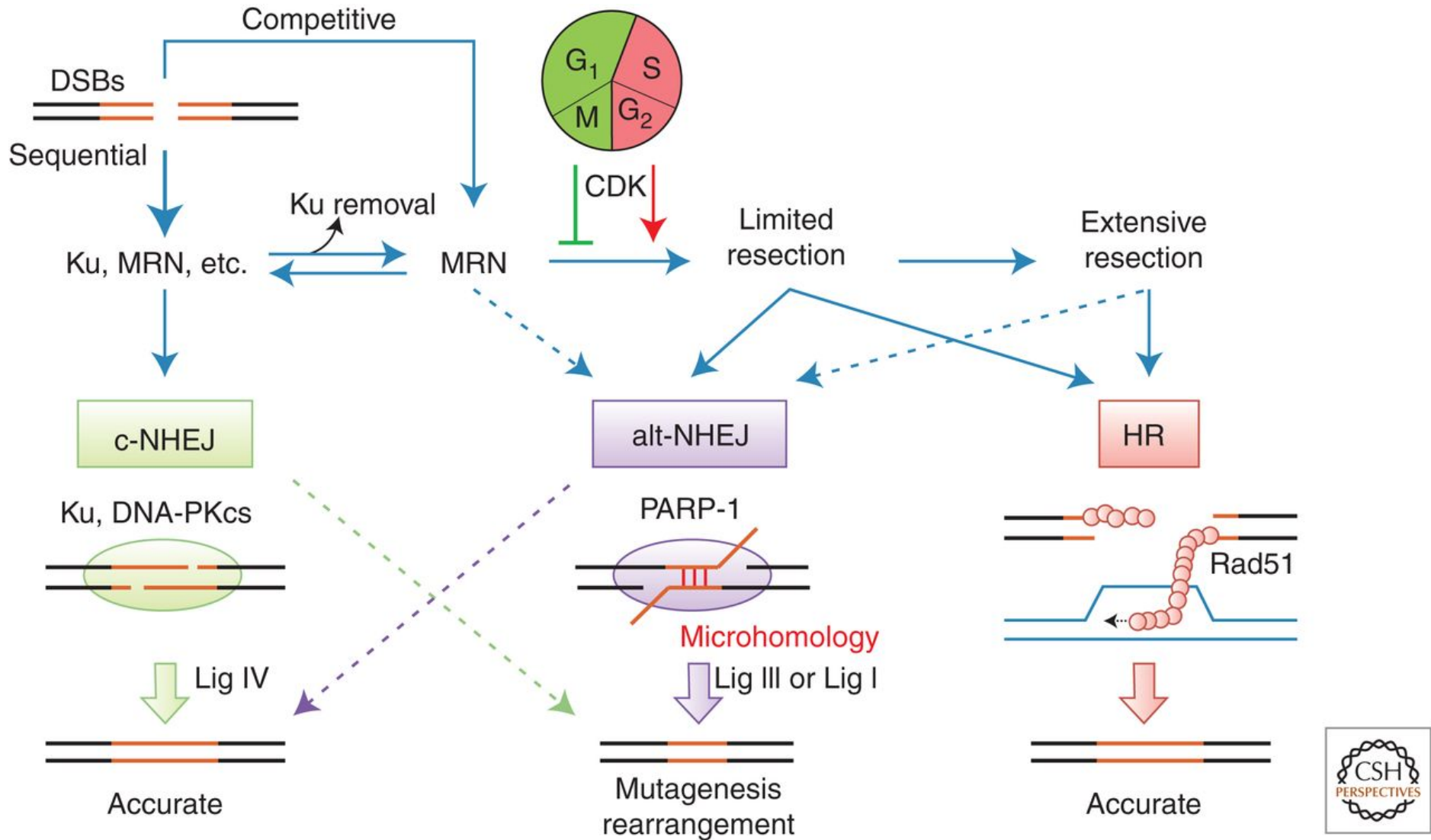
Are there really only two ways to repair DSBs?

Sub-Pathway for NHEJ



MMEJ = Microhomology Mediated End Joining

Disposition of DSBs between repair pathways.



MMEJ = alternate-NHEJ, alt-NHEJ

Non-Homologous End Joining

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

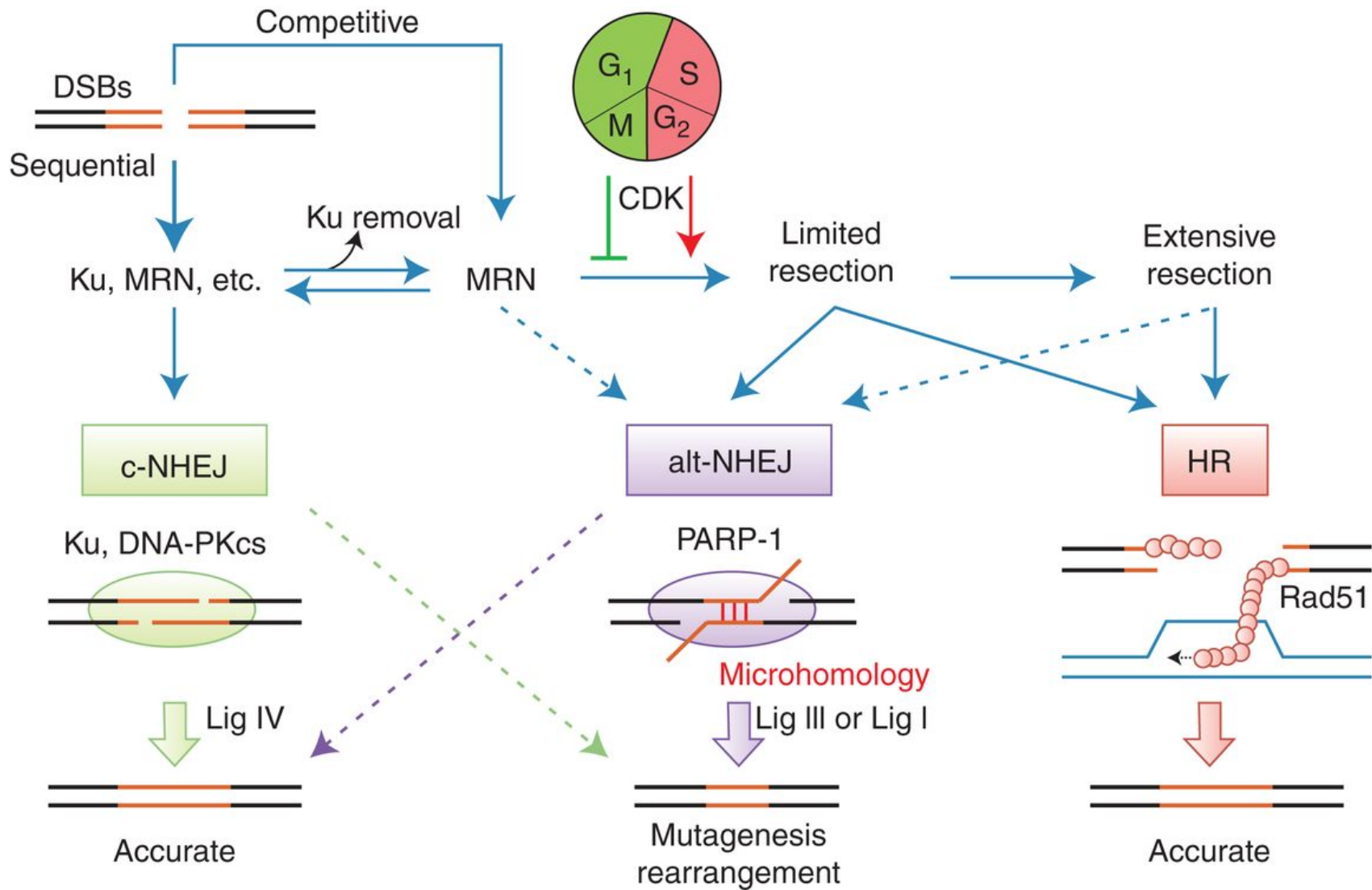
Double-Strand Break Repair via Single Strand Annealing - Alternate NHEJ - MMEJ

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

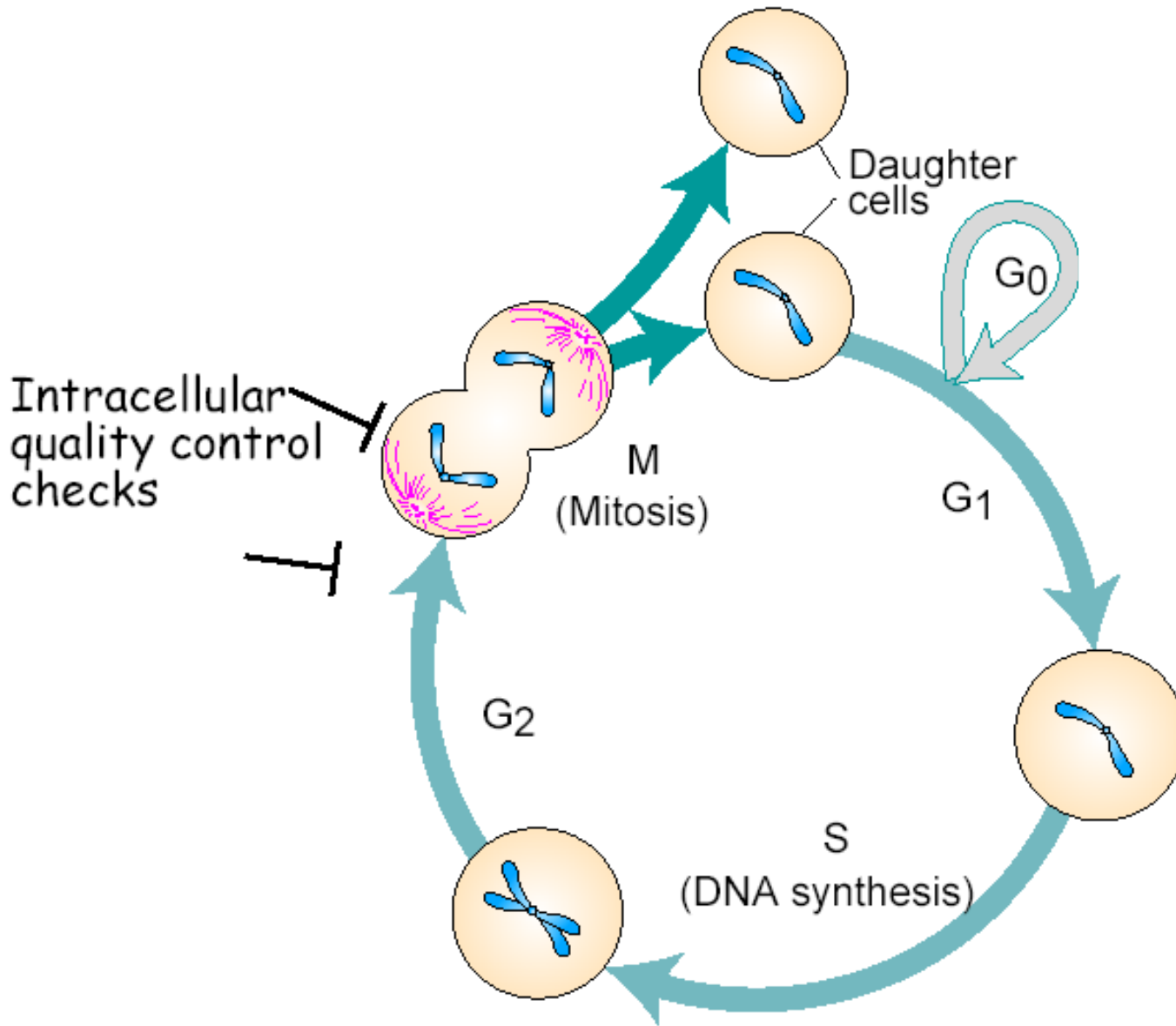
Synthesis-Dependent Strand Annealing (Homologous Recombination)

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

Disposition of DSBs between repair pathways.

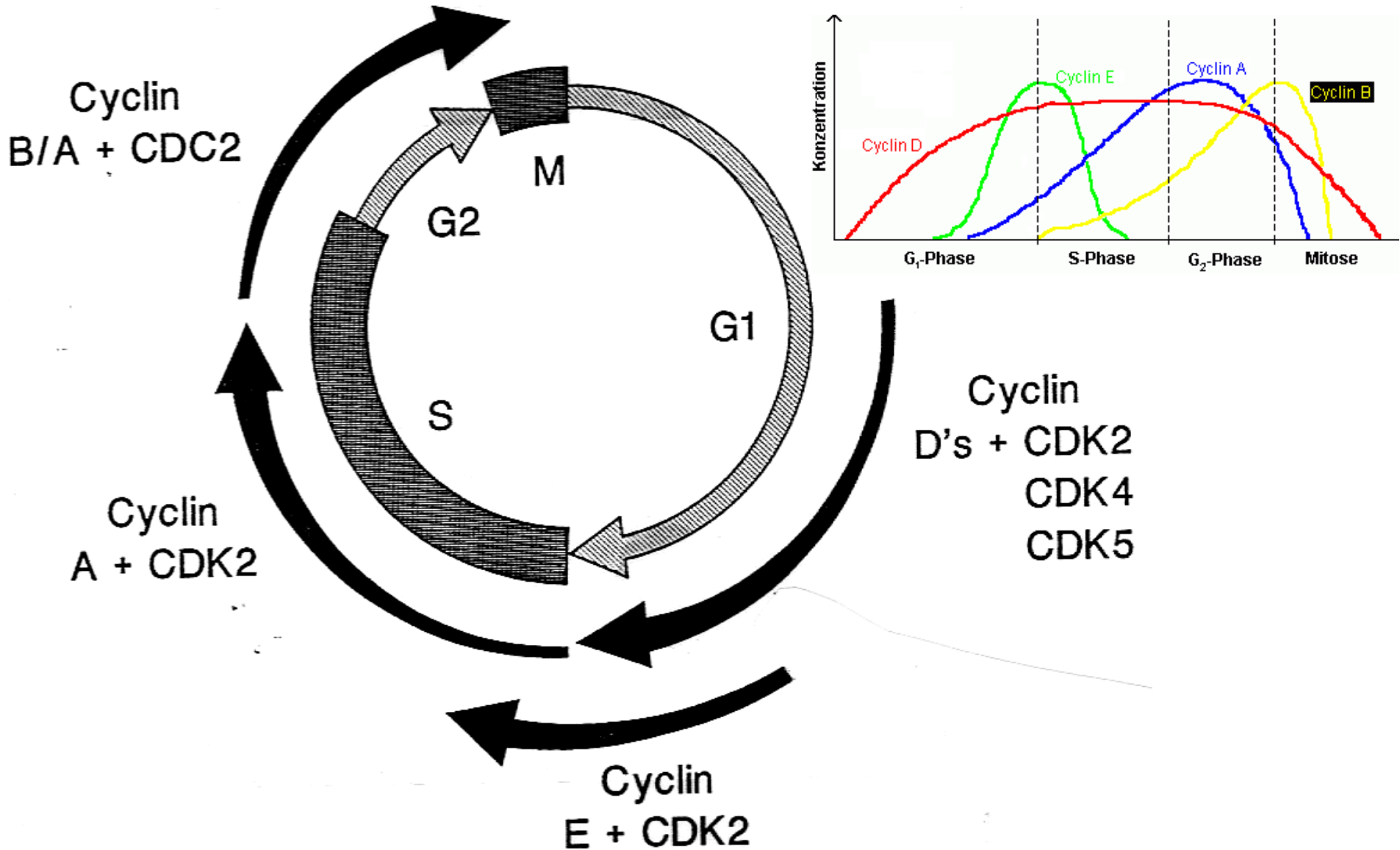


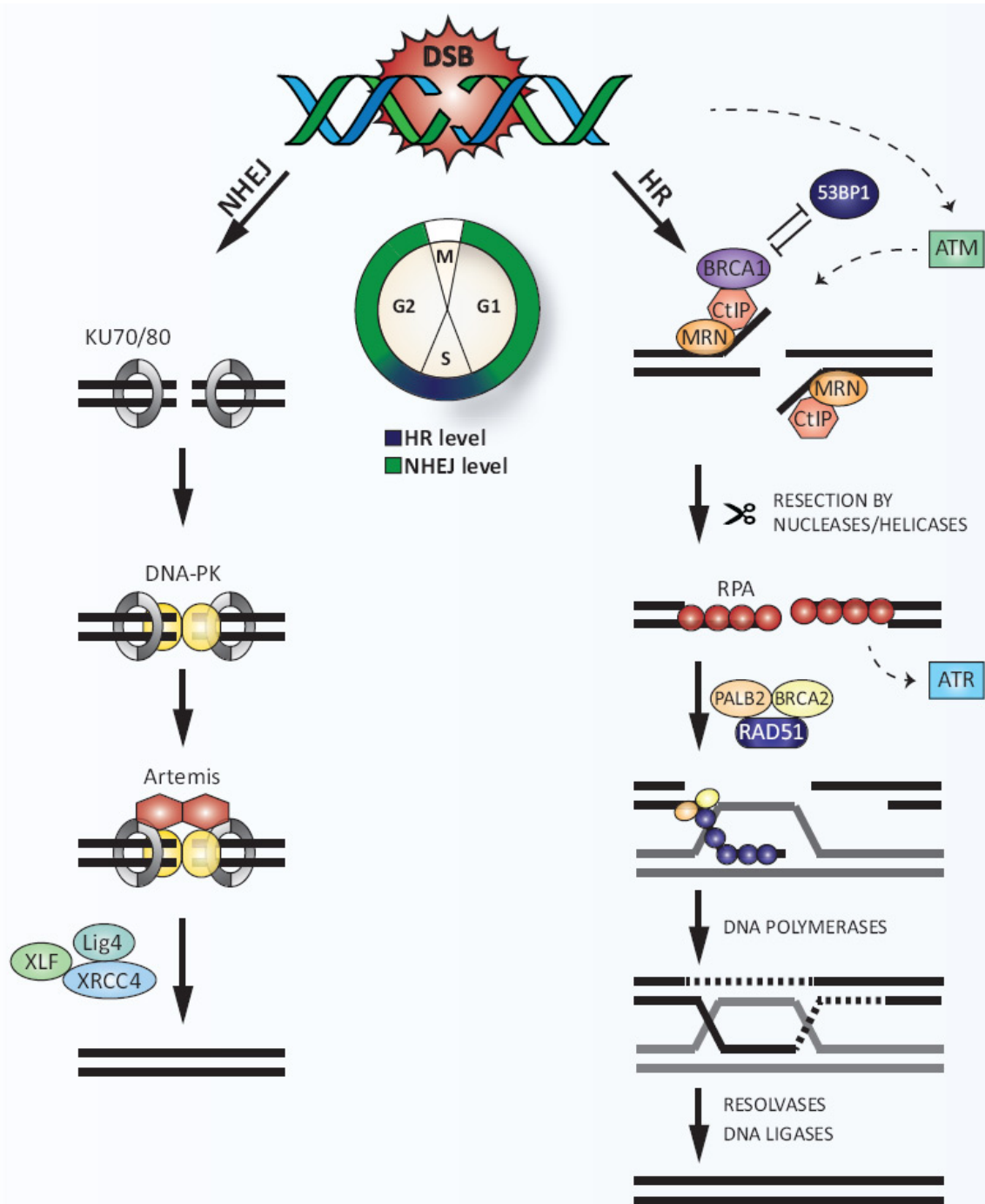
How does
the cell
"know"
where it is in
The Cell
Cycle



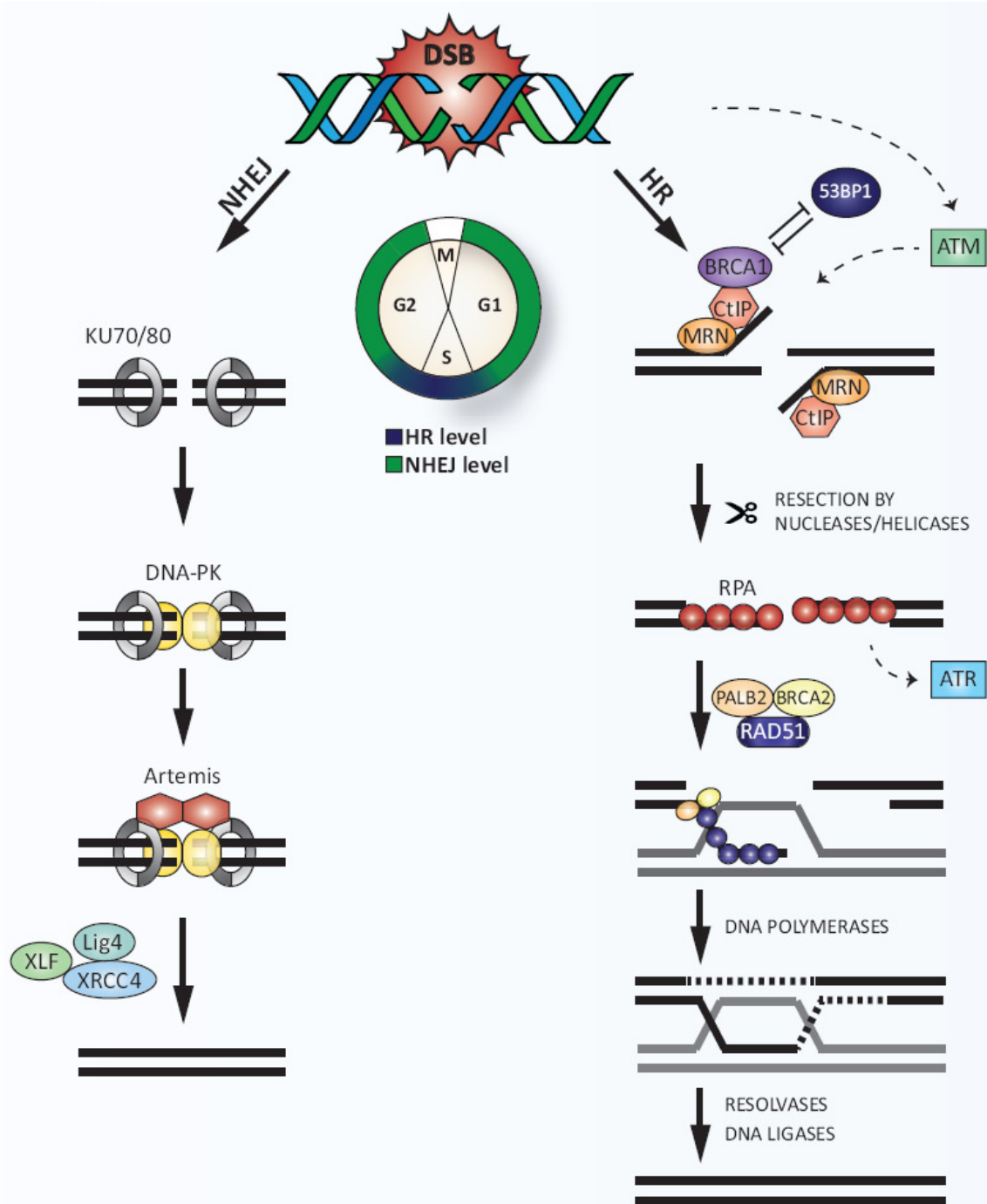
Duplication of chromosomes
DNA Replication

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

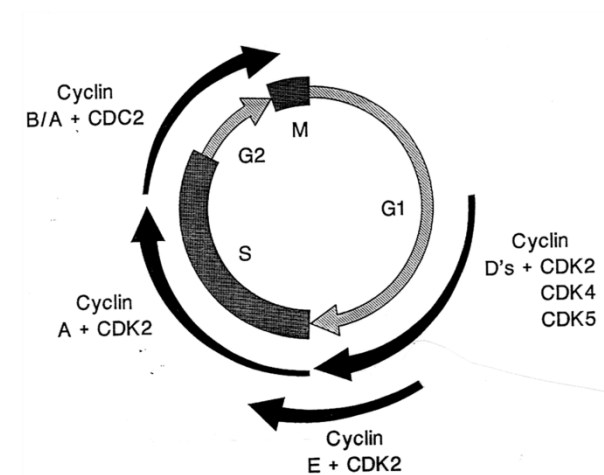


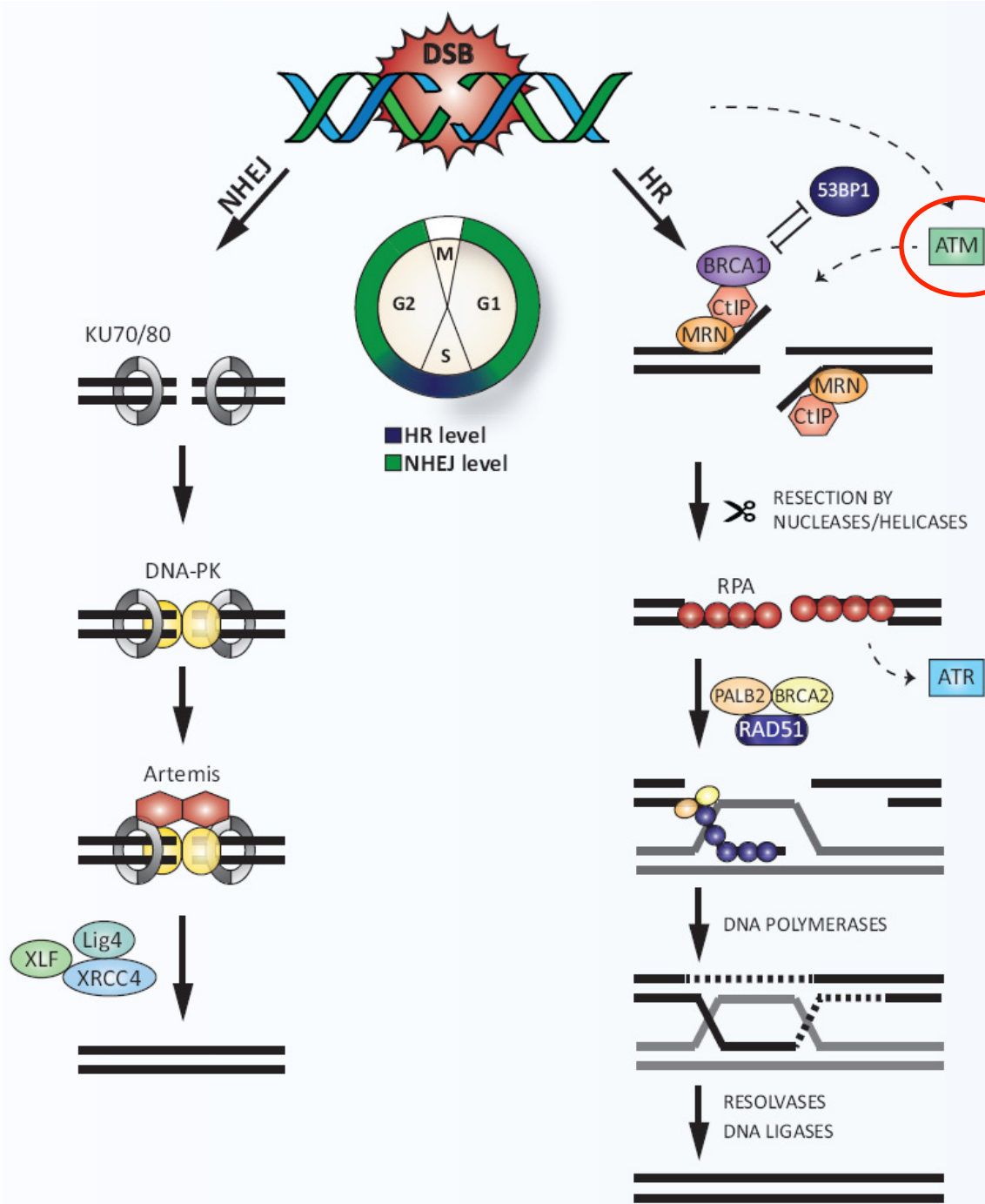


How does the cell decide which pathway to use?



CyclinA-
CDK2
targets the
CtIP/BRCA1
complex



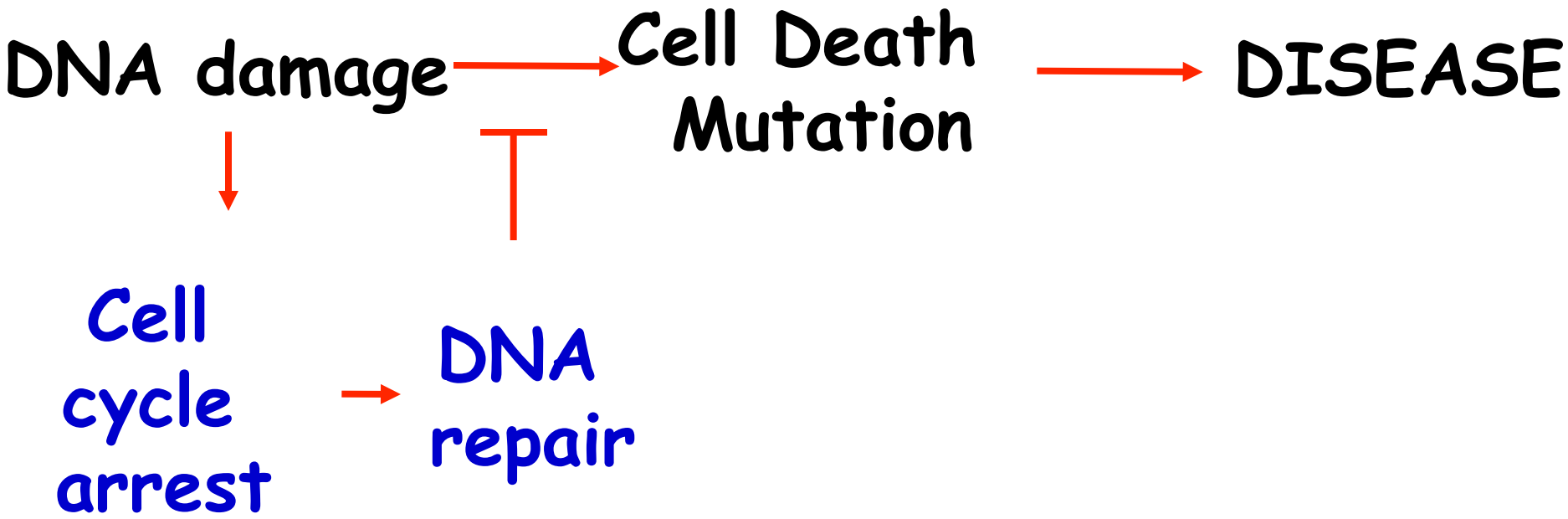


The **ATM** kinase also involved in activating the MRN complex and signaling for S-phase cell cycle arrest



DNA
repair





Ataxia Telangiectasia – Cancer Prone

Defective cell cycle
arrest in response to
DNA Damage affects
neurodegeneration
and cancer
susceptibility

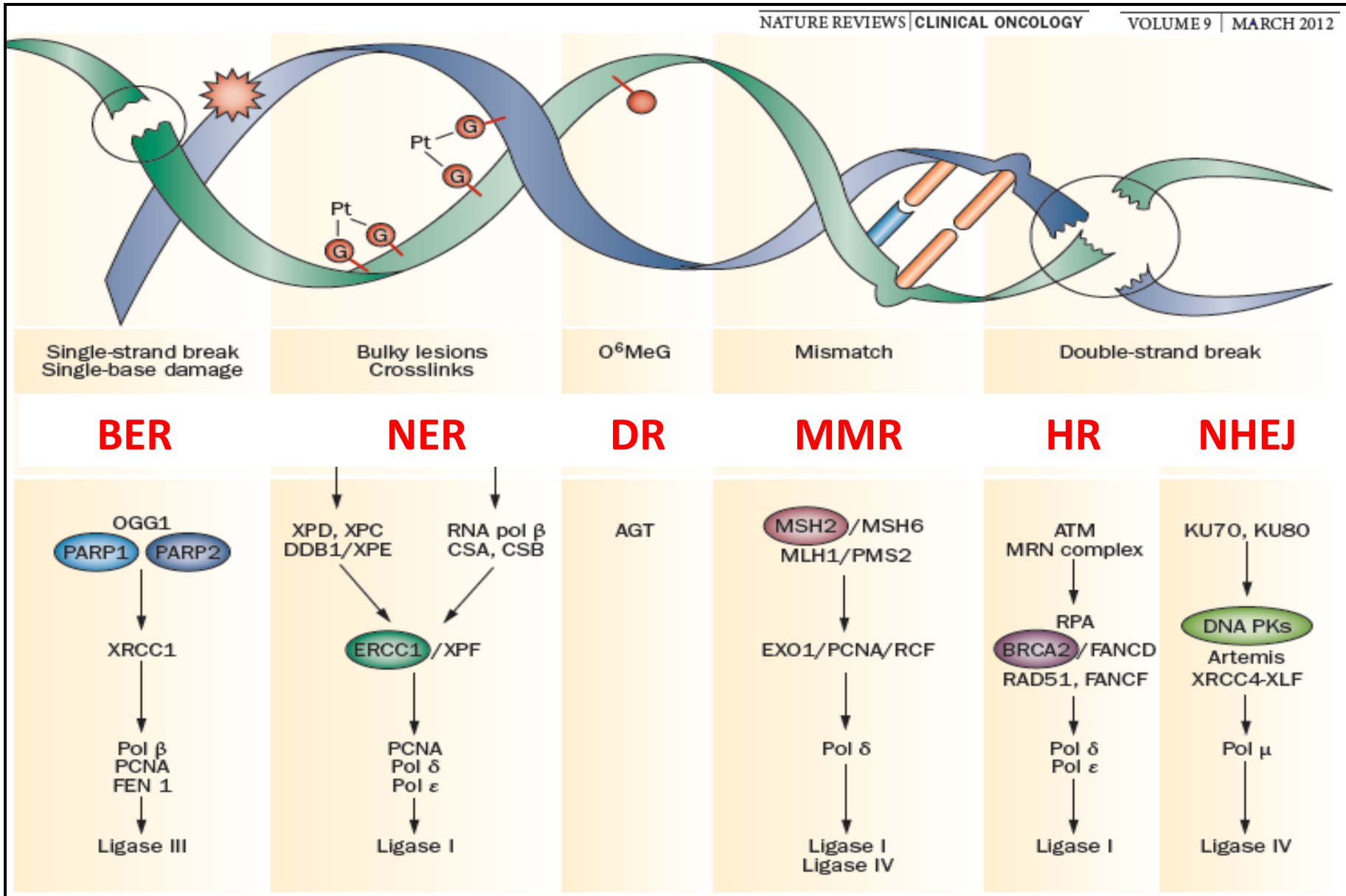


Ataxia Telangiectasia Mutated (ATM)

- 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

Six Major DNA Repair Pathways

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Key Experimental Methods for Module 2

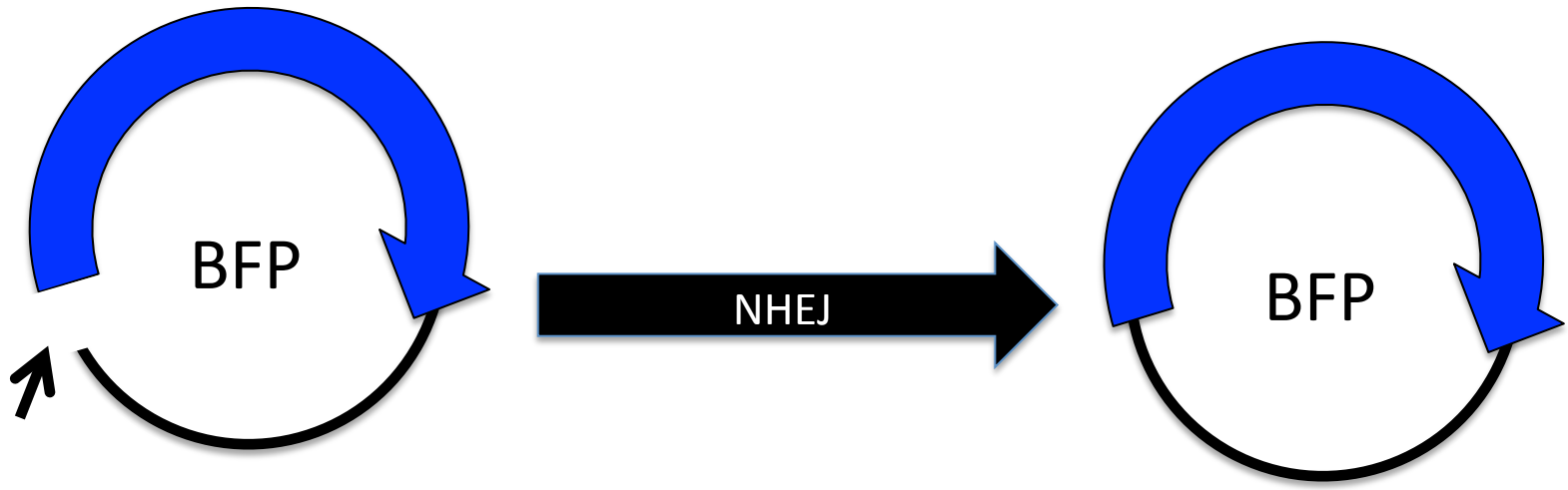
- Mammalian tissue cell culture
- Monitoring protein level by Western blot
- Generating plasmids with DNA damage
- Transfecting plasmids into mammalian cells
- Using fluorescent proteins as reporters of biological processes
- Flow cytometry to measure DNA repair
- Statistical analysis of biological data



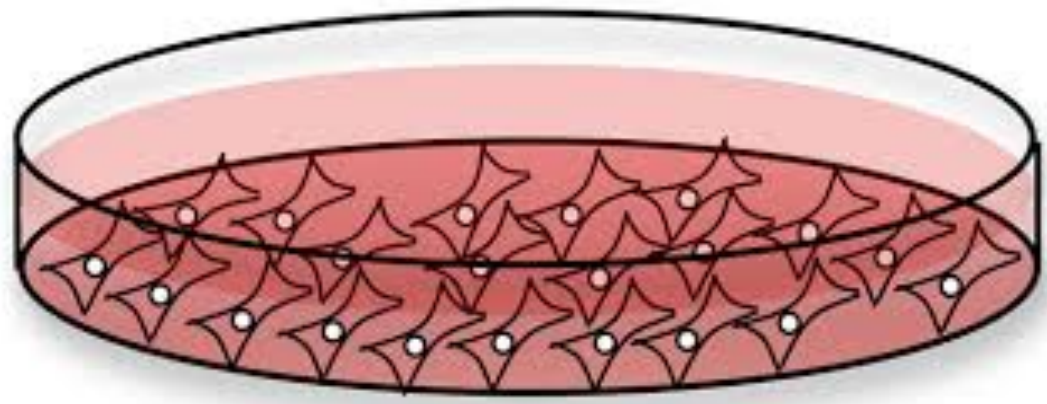
Basis for the fluorescent reporter assay:

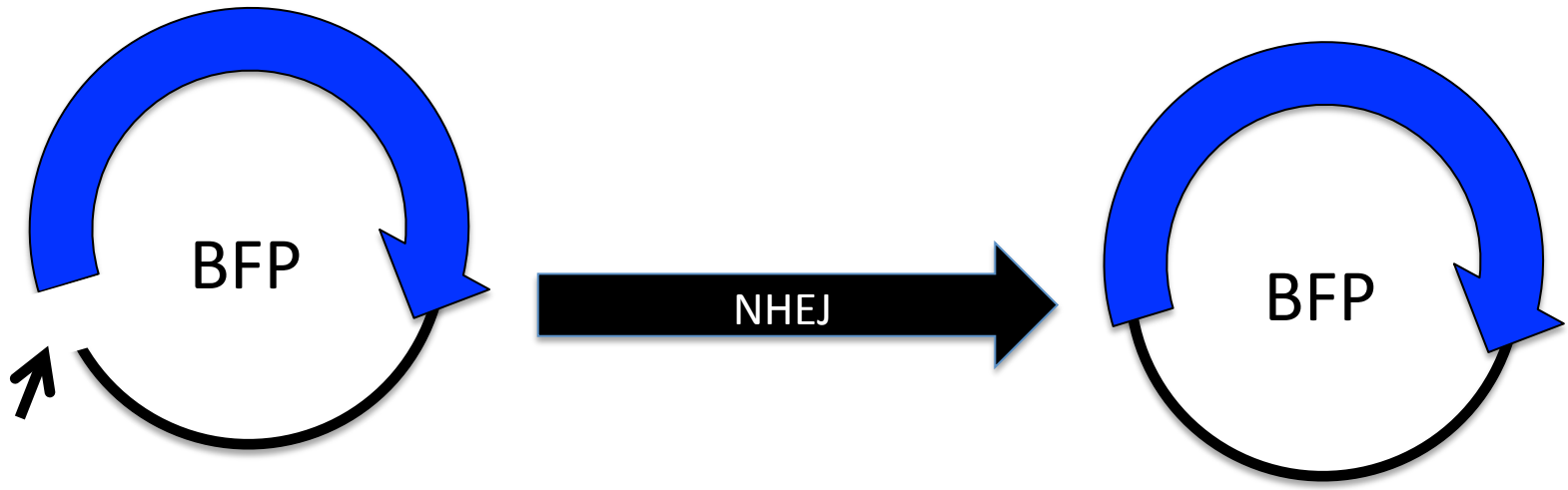


Following digest, the substrate contains a DSB in the 5' UTR that prevents fluorescent reporter expression

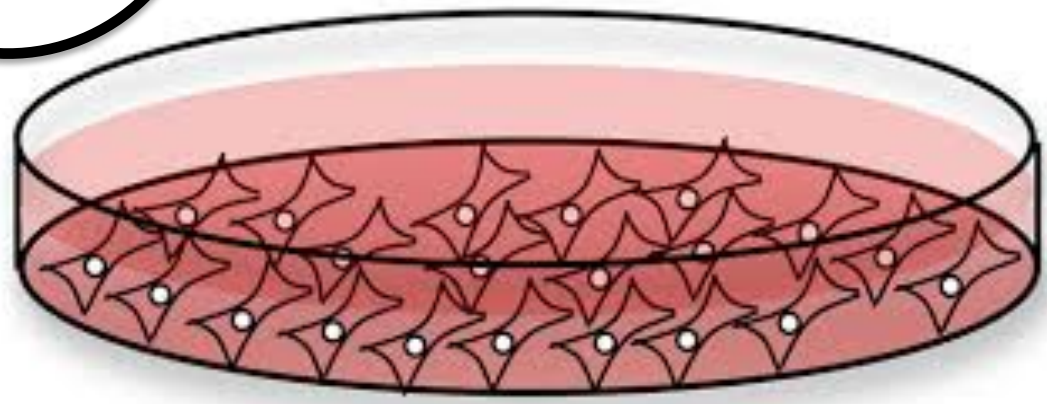
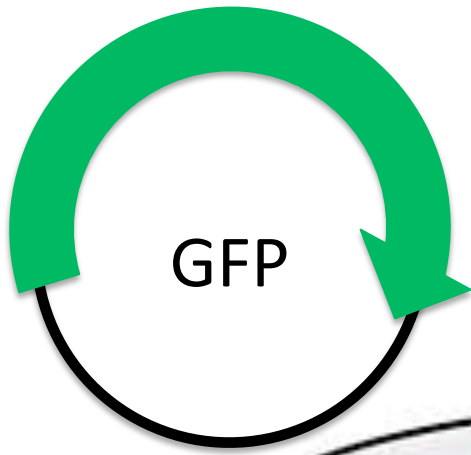


Substrate contains a DNA double strand break

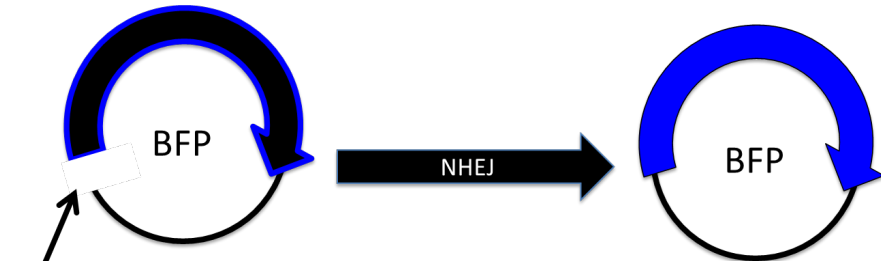
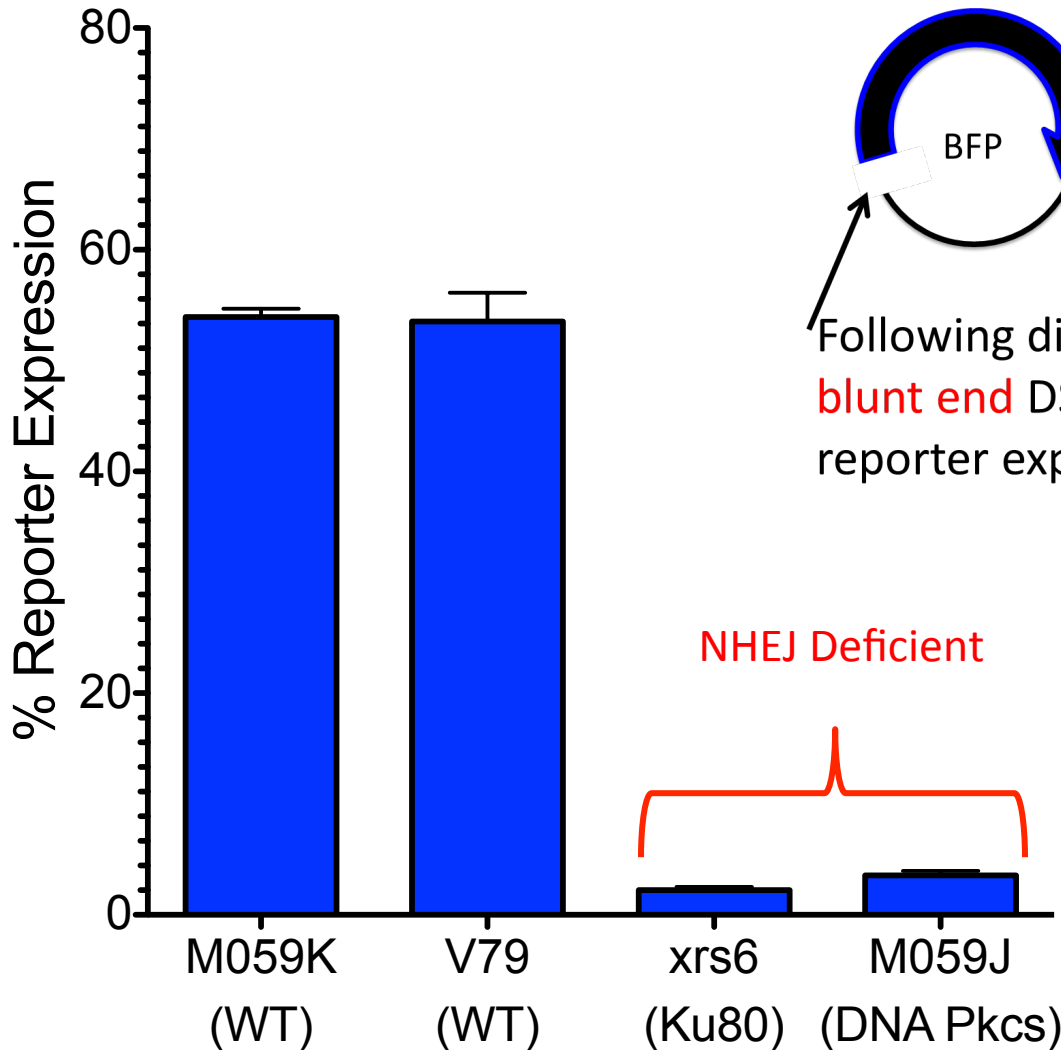




Substrate contains a DNA double strand break

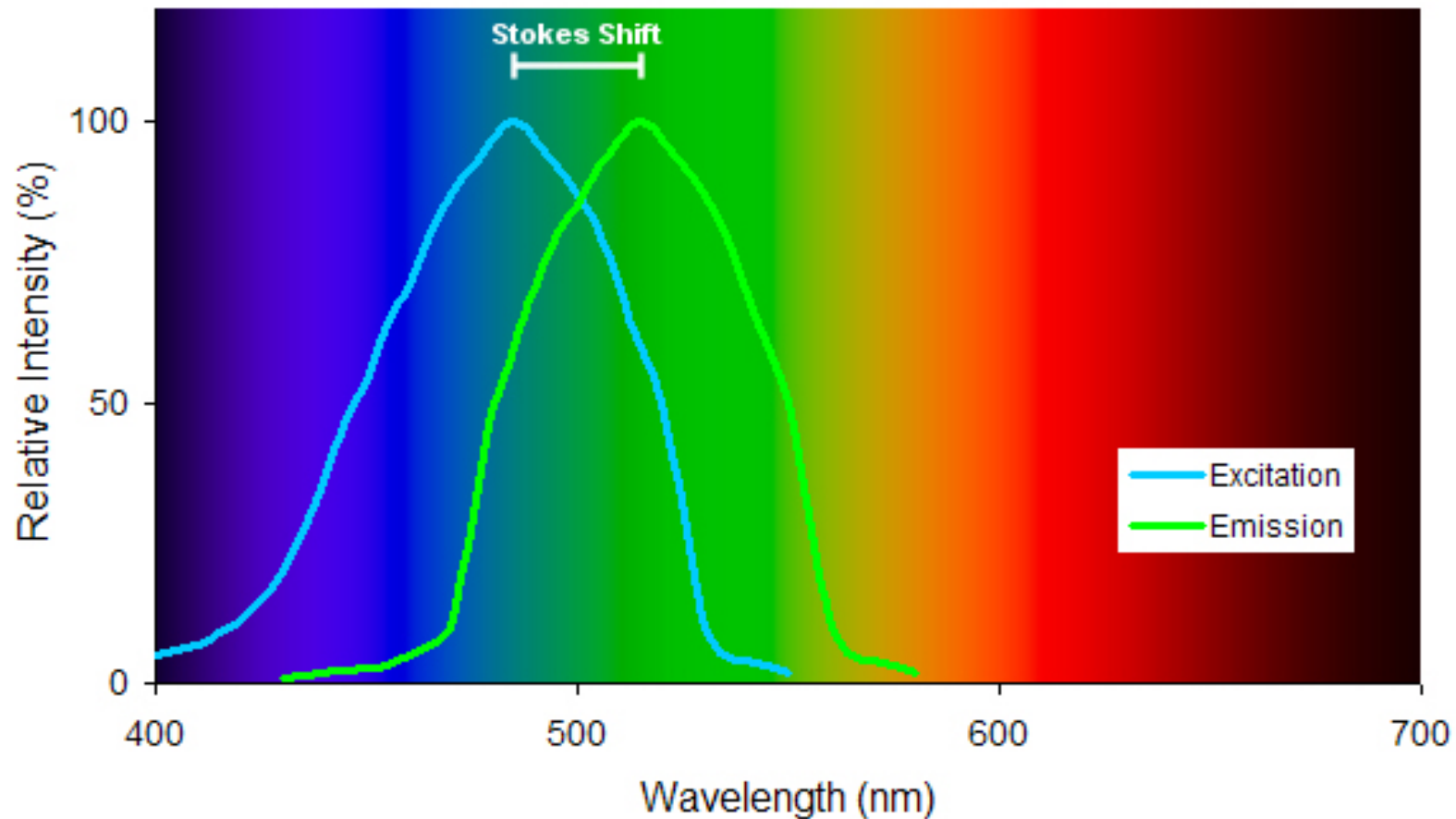
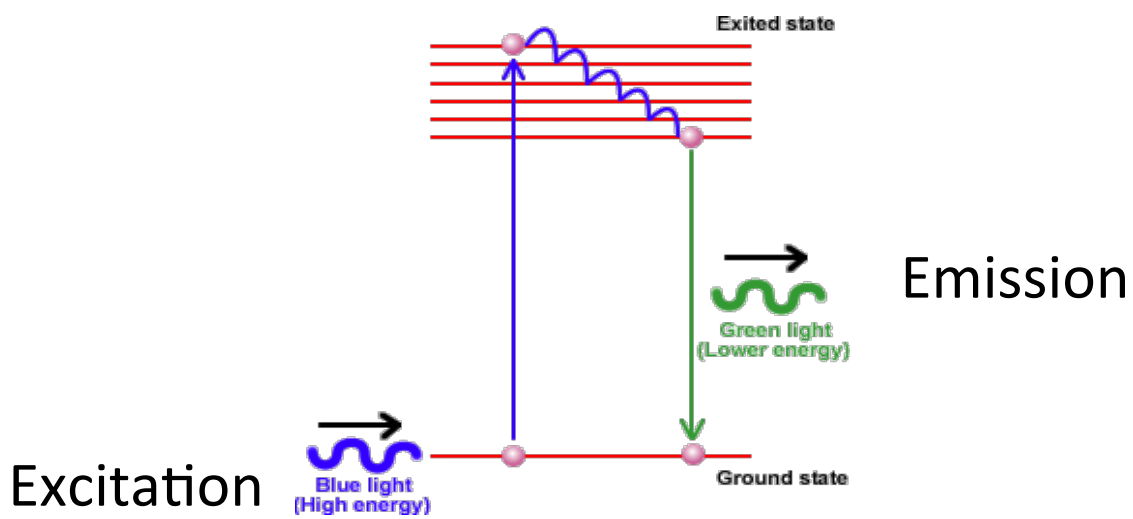


NHEJ HCR in WT and NHEJ defective cells at 18 hours post-transfection:

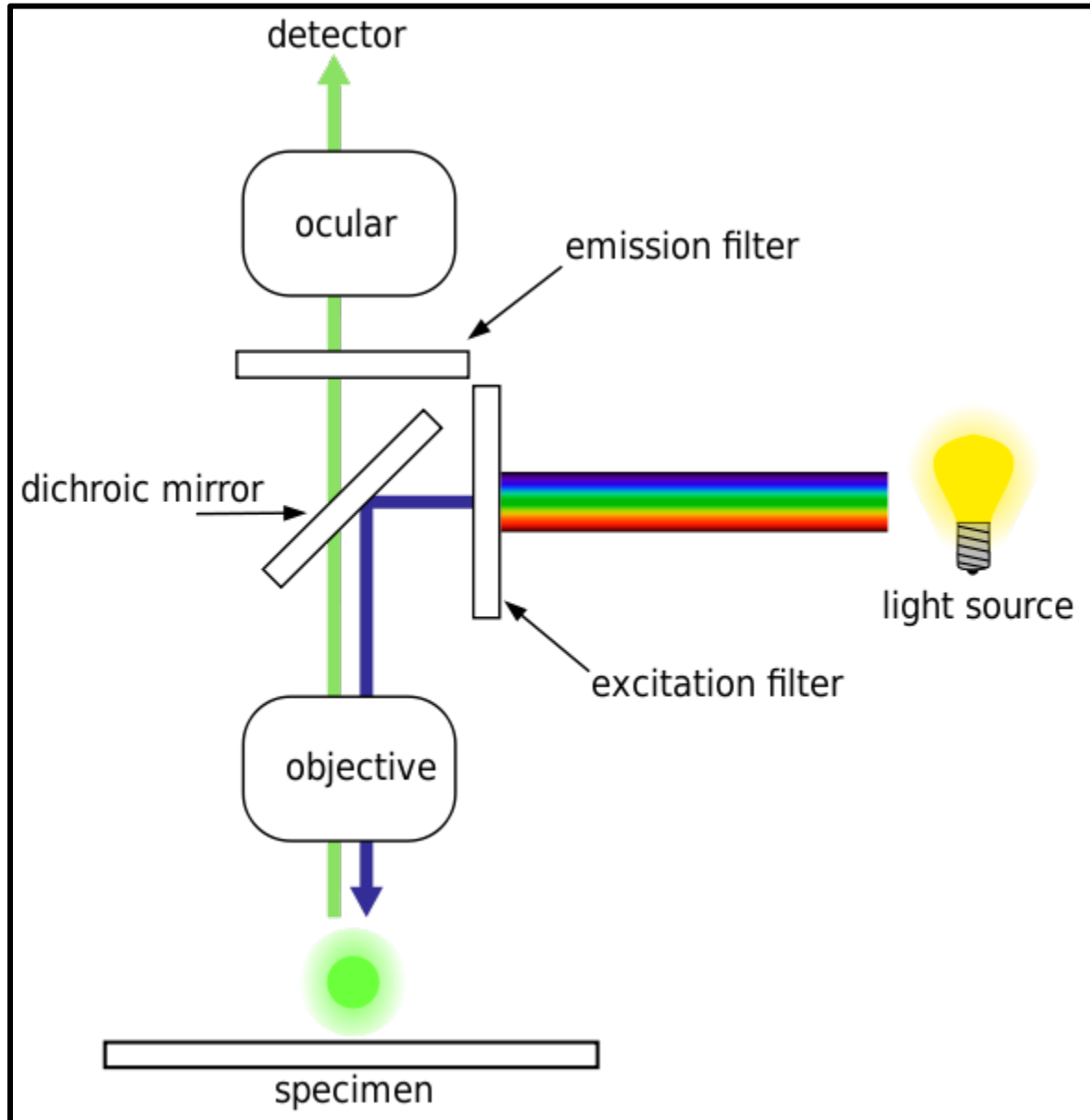


Following digest, the substrate contains a **blunt end** DSB that prevents fluorescent reporter expression

NHEJ Deficient

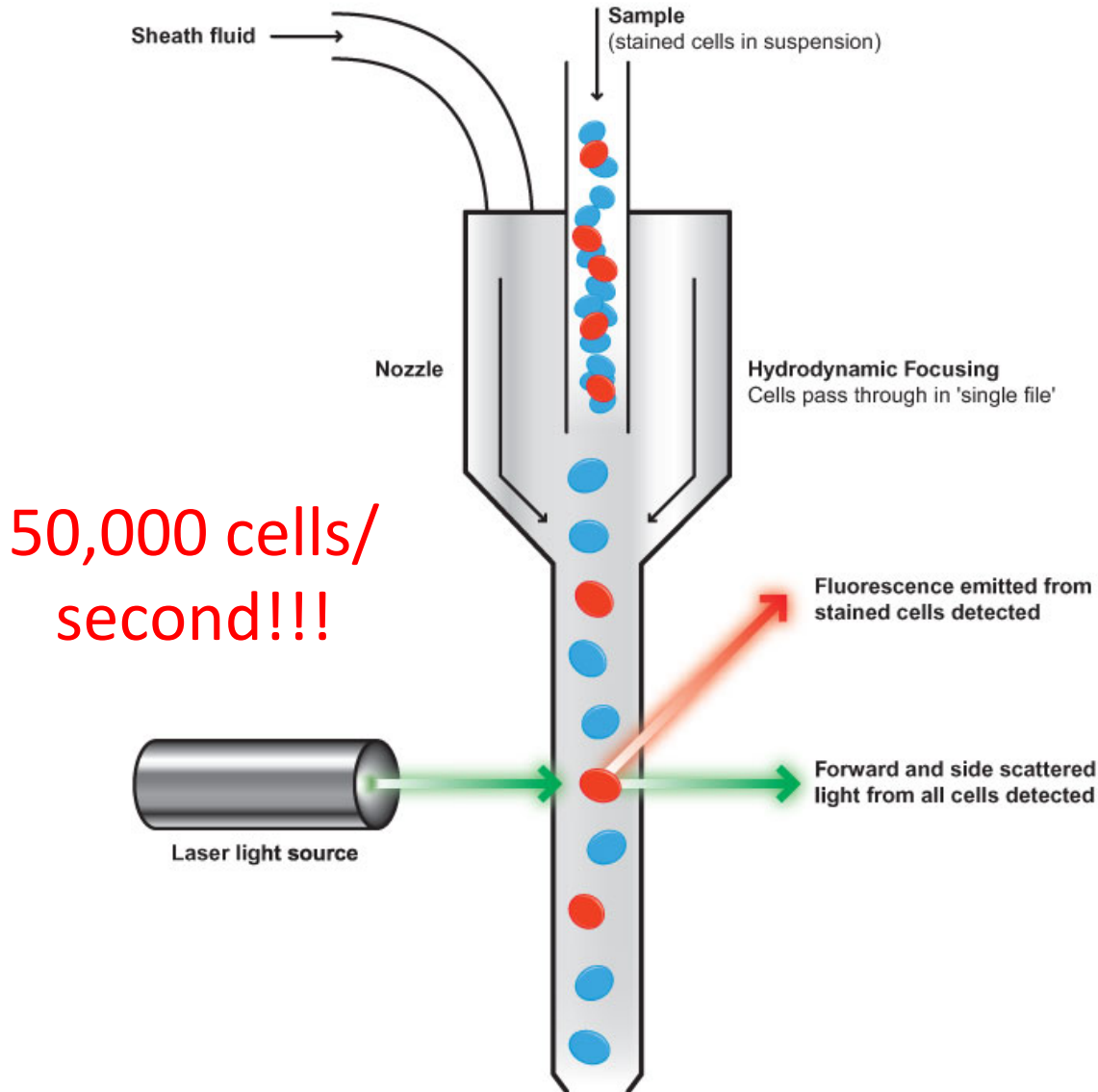


Fluorescence Detection



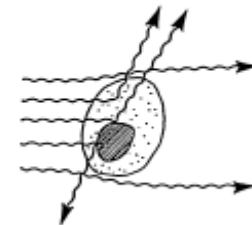
Flow Cytometry

Flow Cytometry



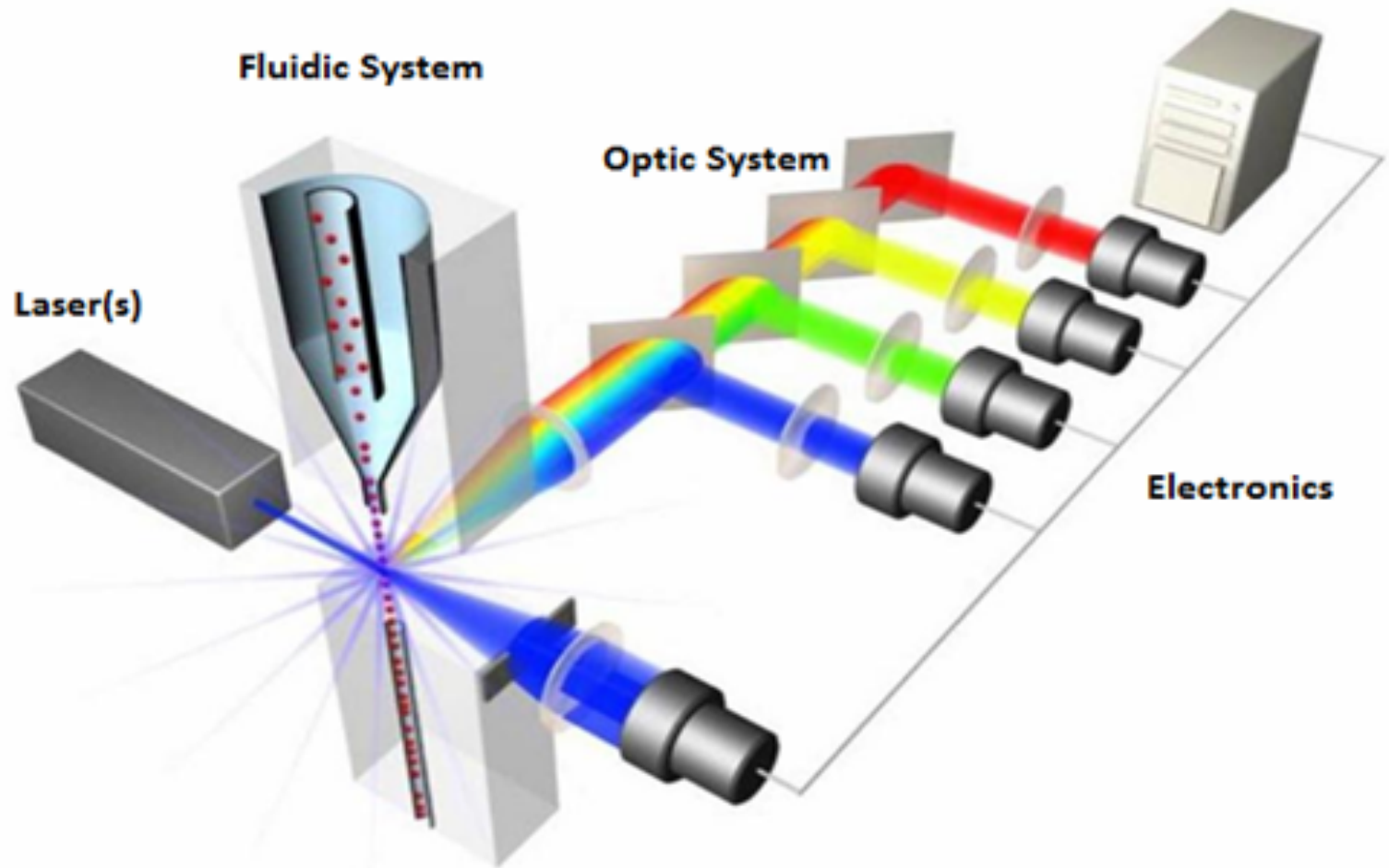
Reports on cell complexity

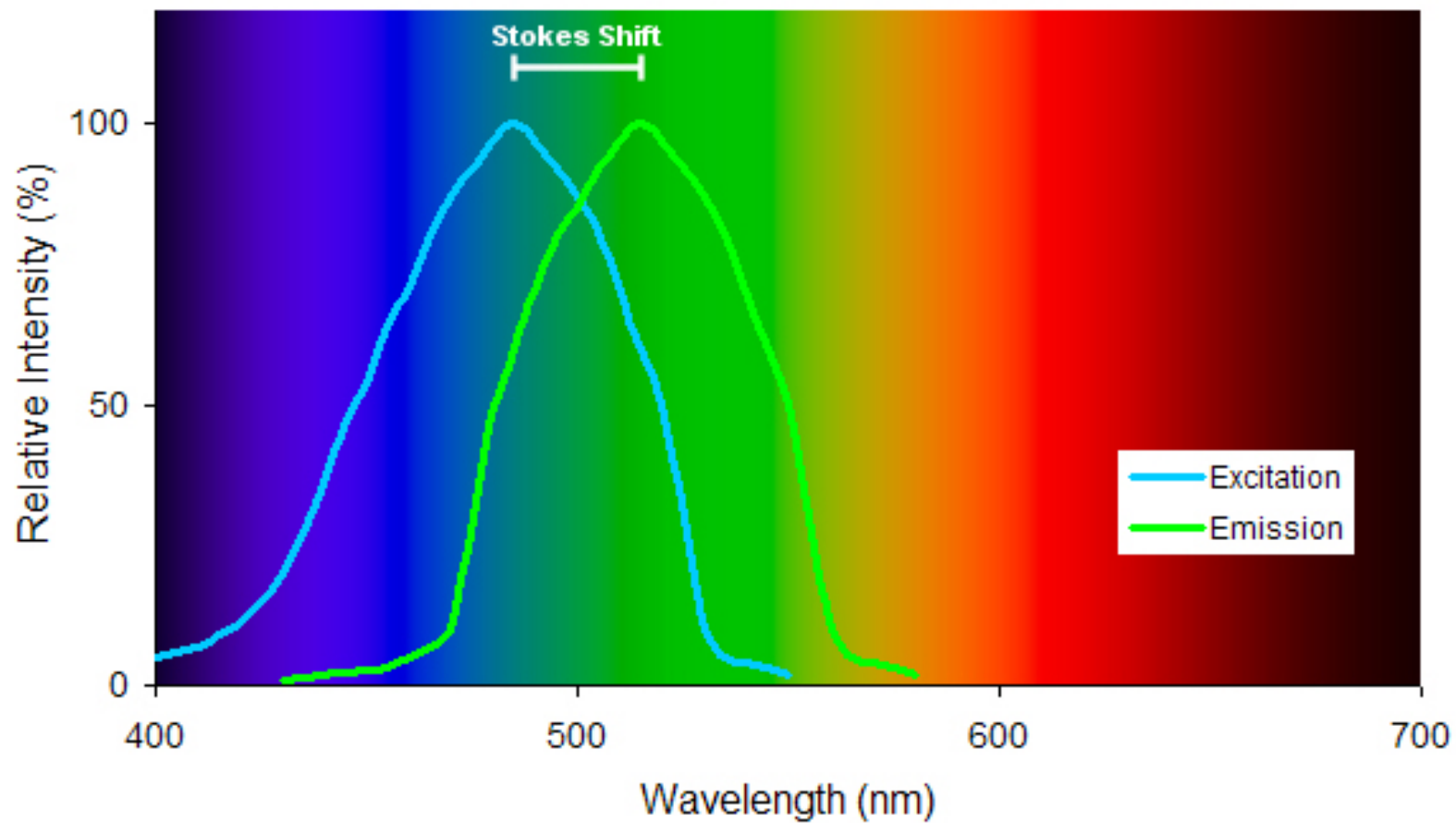
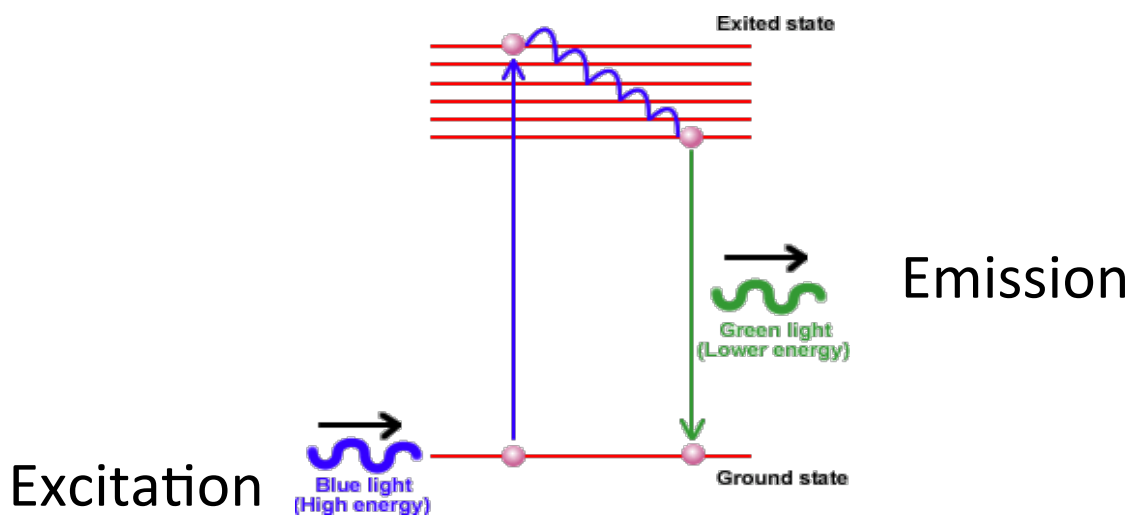
side scatter detector



forward scatter detector

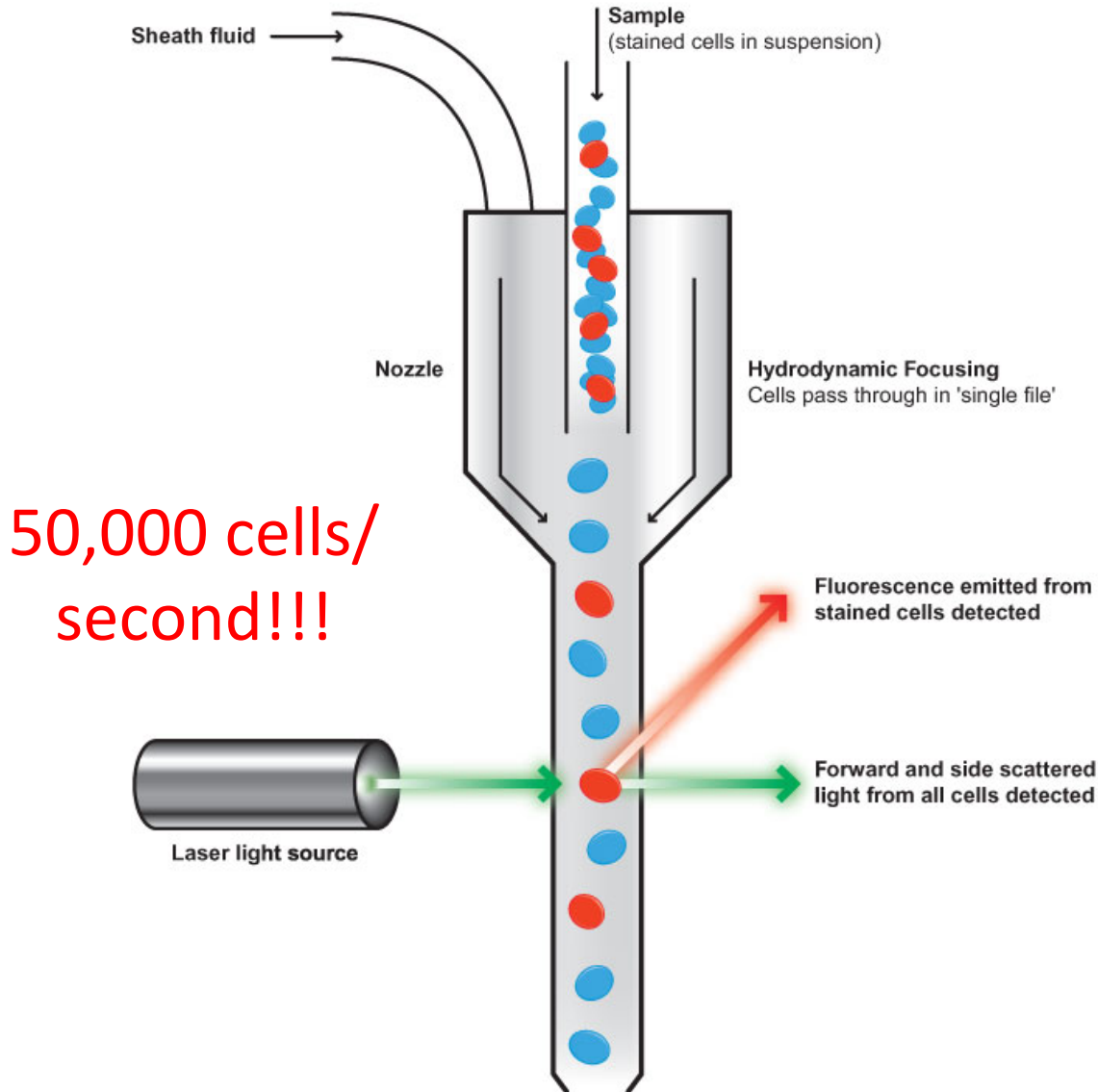
Reports on cell size





Flow Cytometry

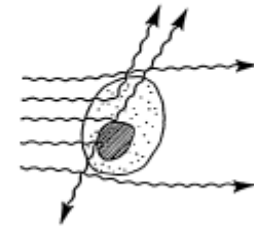
Flow Cytometry



**50,000 cells/
second!!!**

**Reports on cell
complexity**

side scatter detector



forward scatter detector

**Reports on
cell size**

Flow Cytometry

Flow cytometry analyzes cells one by one

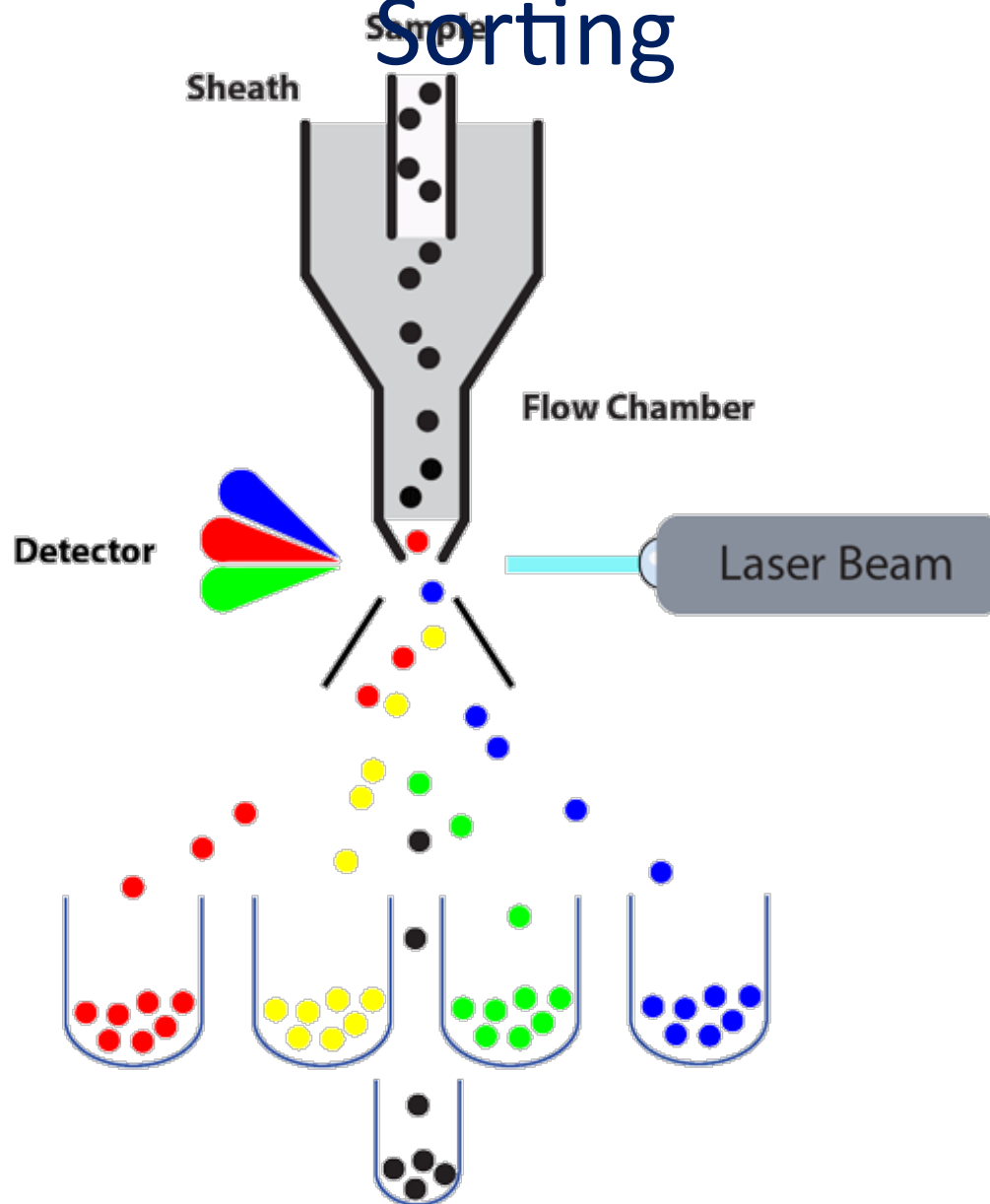
Fluorescence, diffracted and reflected light can be measured for each cell

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



20.109 Spring 2015 Module 2 – Lecture 5

System Engineering and Protein Foundations



Shannon Hughes

Noreen Lyell

Leslie McLain

Nova Pishesha (TA)

Leona Samson (Lectures)

Zachary Nagel (help with development) Alex Chaim

