

INTRODUCTION TO CELL BIOLOGY

<p>Biomolecules</p>	<ul style="list-style-type: none"> • Human cell contains > 20,000 unique <i>proteins</i> and 1000's of distinct <i>metabolites</i> (small molecules including lipids) • A key component of understanding how a protein works is realising where it is within the cell • Much is known about some genes/proteins, nothing about others 	
	<p>Gene ontology → a project that seeks to describe biology using a controlled vocabulary, and in a discrete way, such that a computer can understand what a protein does. This covers three domains:</p>	<p>i) Cellular Component: the parts of a cell with which the protein associates</p>
		<p>ii) Molecular Function: the elemental activities of a protein at the molecular level, such as binding or catalyst</p>
		<p>iii) Biological Process: operations or sets of molecular events with a defined beginning and end, pertinent to the functioning of integrated living units (cells, tissues, organs, organisms)</p>
<p>Protein location within the cell → important because:</p> <ul style="list-style-type: none"> • Eukaryotic cells are divided into various morphologically and functionally distinct compartments <ul style="list-style-type: none"> - These compartments are made up of 100's to 1000's of proteins which provide its structure and function • Proteins must be targeted to the appropriate compartment to ensure proper function • Knowing a proteins subcellular localisation is critical to understanding the function of individual proteins 		
<p>Detecting individual proteins in cells → cannot be observed with electron microscopy, so identification techniques have to be developed:</p> <ul style="list-style-type: none"> • Utilising the immune system <ul style="list-style-type: none"> - Body recognises foreign material and combats it (fight, kill, remove) - Develop <i>antigen-specific</i> antibodies to combat foreign material - By putting a human protein in another mammal, that mammal will develop antibodies <i>specific to that protein</i> - These antibodies can then be used to identify the location of that protein within the human cell <p>Indirect immunolabelling → the primary antibody developed in another mammal is directed against an immobilised antigen, antigen A. It binds, and then <i>marker-coupled</i> secondary antibodies directed at the non-human antibody bind, and mark the proteins location (as they are easily visible)</p>		

Immunofluorescence Microscopy	<ul style="list-style-type: none"> i) Cultivation: <i>culture</i> cells (mammalian cell culture is the process of growing animal cells in vitro in a flask or dish) ii) Fixation: in living cells, organelles move around. Thus, to stabilise the cells internal structures, the organelles are cross-linked together, making them static iii) Permeabilization: the lipid bilayer does not allow antibodies to pass through, and thus holes must be punched in it iv) Blocking: helps get specificity v) Primary antibody incubation: antigen-specific antibodies developed from a non-human mammal enter the cell and bind to their antigen vi) Secondary antibody incubation: these antibodies bind to the non-human antibody, and are marked with dyes that <i>fluoresce</i>, meaning they <i>absorb</i> light at one set of wavelengths and <i>emit</i> light at another set of wavelengths (process called <i>fluorescence</i>) vii) Nucleus staining: by staining the nucleus and other particular regions of the cell, a reference point is made, which can be used to identify fluorescent regions viii) Mounting: tissue is 'mounted' onto a 12mm x 1mm coverslip ix) Microscopy: tissue is observed
	Fluorescence Spectra <ul style="list-style-type: none"> • Fluorescent dyes absorb and emit light best at certain wavelengths <ul style="list-style-type: none"> - We can show this by plotting absorbance or emission versus wavelength on a graph - Since the light that gets absorbed by the dye excites the dye molecules to a more energetic state, it is called absorption <i>excitation</i>
	Stoke Shift → the difference between the peak emission and peak absorption wavelengths <ul style="list-style-type: none"> • Typically, excitation (absorption) light is many times brighter than the emission light <ul style="list-style-type: none"> - If we shone the excitation light onto the sample and looked for fluorescence, we might have a hard time seeing our emission - This is solved by using filters that allow the excitation light to get to the sample, but only the emitted light gets to our eyes/camera

The Airy Disk & Resolution

- When light from the various points of a specimen passes through the objective lens and is reconstituted as an image, the various points of the specimen appear in the image as small patterns (not points) known as *Airy patterns*
 - In Airy Disks, there is a maximum point of light intensity which is surrounded by rings of fluorescence
 - This scattering prevents us from seeing objects close together (i.e.- objects whose Airy Disks overlap)
- **Effective Resolution (d_0)** → the smallest distance between two objects that still allows for them to be seen as separate entities
 - For light microscopes, this is about $0.2\mu\text{m}$ (200nm)
 - E.g.- if two objects were less than 200nm apart, we could not resolve each one
- **Super Resolution Microscopy** → allows for greater resolutions because it can resolve more points of light
 - Resolution of 10-30 nm
- **Confocal Fluorescence Microscopy** → captures the light from one focal plane and removes any other sources of light
 - Allows the visualisation of thick specimens
 - Removes blur that conventional microscopy can't
- **Live Cell Imaging** → uses naturally occurring fluorescent proteins to observe living cells
 - Often able to be genetically coded into animals

MODULE 1- CELL CYCLE

<p>Cell → a membrane-bound structure containing biomolecules that acts as the structural, functional and biological unit of all organisms</p>	
<p>Central Dogma of Molecular Biology</p>	<p>Gene expression dictates cell identity and function:</p> <p style="text-align: center;">DNA → RNA → Protein</p> <p>Organisms genome encoded in DNA, which is <i>transcribed</i> into RNA by RNA polymerase. RNA is then translated into protein, which performs a function.</p>
<p>Gene structure and nomenclature</p>	<p>Promotor → transcription factors and polymerase bind here to begin transcribing the gene</p> <p>Exon → protein coding region that is translated into mRNA</p> <p>Introns → non-coding regions that are spliced out of mRNA (can have regulatory functions)</p> <p>Diploid organisms → have two copies of every gene. If both copies are identical, the organism is <i>homozygous</i> at that locus.</p> <p>Alleles → different versions of the same gene (differences in DNA sequences of the same gene). If the two copies of a gene are different, the organism is <i>heterozygous</i> at that locus.</p>
<p>DNA Transcription into mRNA</p>	<p>Capping → after transcription, the primary RNA transcript is capped at the 5' end with a special nucleotide; important for stability and translation</p> <p>Cleavage → primary RNA transcript is cleaved at the 3' end</p> <p>Polyadenylation → 3' end is polyadenylated by an enzyme called a polyadenylase; mRNA will have long tails of 'A's', important for stability and translation (binding site for proteins)</p> <p>RNA splicing → introns are removed by splicing factors that bind to acceptor and donor sites within introns. Results in mature mRNA</p>
<p>mRNA Translation into Protein</p>	<p>Single-stranded mRNA is translated into a protein within the ribosomes (after exiting the nucleus and travelling through the cytoplasm)</p> <p>Codons → exist within mRNA as nucleotide triplets; they specify what amino acid goes at their point in the sequence</p> <p>tRNA → have anticodons and amino acids attached; the anticodon is matched appropriately to the codon, such that the correct amino acid is retrieved</p>
<p>Proteins</p>	<p>Proteins → the workhorse of the cell:</p> <ul style="list-style-type: none"> • Structural • Sensors • Transporters • Enzymes • Transcription Factors • Cellular Communication • Signal Transduction

	<p>Amino Acids → amino acid sequences make up proteins, though there are only 20 amino acids</p> <p>Protein (amino acid) sequences → determine protein <i>structure</i></p> <p>Protein structure → determines protein <i>function</i></p> <p>Prion → infectious protein with <i>normal</i> DNA sequence; infectious nature allows it to convert normal proteins into prion form</p> <p>Goes against Central Dogma because while the prion is a mutated protein, its DNA is normal (i.e.- protein structure/function is not determined by DNA sequence, but by something else)</p>
Approaches for Studying Cells	Cell Biology → through direct observation (microscopy); labelling cell structures of interest and observing them
	Biochemistry → isolating and describing proteins
	Genetics → looking at mutant genes, their mutant proteins, and the effects on the cell or animal
	Genomics and Proteomics → looking at all genes (or proteins) at the same time
	Developmental Biology → studying differential gene expression and the signals that lead to the mature organism
The Cell Cycle	Interphase → everything except Mitosis
	G1 Phase (Gap 1) → recovery from mitosis, growth
	S Phase (Synthesis) → DNA is duplicated
	G2 Phase (Gap 2) → pre-mitosis checkpoints
	M Phase (Mitosis) → chromosome segregation and cell division
	G0 Phase (Gap 0) → temporary or permanent exit from the cell cycle
Chromosome Structure and Movements	Chromosome → the structural unit of genetic material consisting of genetic material consisting of double stranded DNA and proteins
	Chromatid → one copy of a duplicated chromosome (still a chromosome)
	Sister Chromatids → identical copies of a chromosome joined by a centromere
	Homologous Chromosomes → chromosome pair that includes one from each parent (maternal and paternal). Different alleles.
	Cohesins → proteins that hold the sister chromatids together
	Centromere → repetitive DNA sequence that serves as a landing pad for mitotic machinery
	Kinetochores → protein complex that binds to the centromere, linking the centromere to microtubules
	<p>Chromosome Segregation in Mitosis →</p> <ul style="list-style-type: none"> • Duplicated chromosomes line up independently of one another • Each pair of sister chromatids (identical) separate • Each daughter cell gets <i>all</i> of the genetic information

Stages of Mitosis	Interphase → chromosome duplication and cohesion; centrosome duplication, one for each daughter cell (both occur in S Phase)	
	Prophase → breakdown of interphase microtubule and its replacement by two mitotic asters (centrosome + emerging microtubules); mitotic aster separation; chromosome condensation for movement	
	Prometaphase → nuclear envelope breakdown; condensed chromosomes captured, bi-oriented and brought to spindle equator by microtubules	
	Metaphase → chromosomes aligned <i>independently</i> at the metaphase plate	
	Anaphase → Anaphase Promoting Complex (APC/C) activated, and cohesins degraded	Anaphase A → chromosome movement to poles
		Anaphase B → spindle pole separation
	Telophase → nuclear envelope reassembly; assembly of contractile ring	
	Cytokinesis → reformation of interphase microtubule array; contractile ring forms cleavage furrow	
Mitotic Machinery	Centrosomes (Spindle Poles) → microtubule organising centres	Centrioles → contained within the centrosomes and are composed of bundles of microtubules (source of microtubules)
	Microtubules → capture and move chromosomes; anchor to the plasma membrane	
	Molecular Motors → drive chromosome movement	
Mitosis vs Meiosis	Meiosis → cell division that only occurs in the germline cells; the goal of meiosis is to produce four unique gametes that are appropriate for sexual reproduction (i.e.- shuffle alleles and reduce to 1n)	
	Mitosis → cell division that occurs in the somatic cells; the goal of mitosis is to produce two identical daughter cells (i.e.- keep all alleles the same)	
	n → number of chromosomes of each type	
	Differences Between Mitosis and Meiosis → <ul style="list-style-type: none"> • Mitosis <ul style="list-style-type: none"> - In a mitotic cell, the number of chromosomes goes from 2n to 4n after S-phase - Most cells are somatic, and have either 2n or 4n • Meiosis <ul style="list-style-type: none"> - A reductive process that produces gametes viable for sexual reproduction - Reduces the number of chromosomes down to 1n - In sexual reproduction, new combinations of existing alleles create new phenotypes (this drives evolution) 	

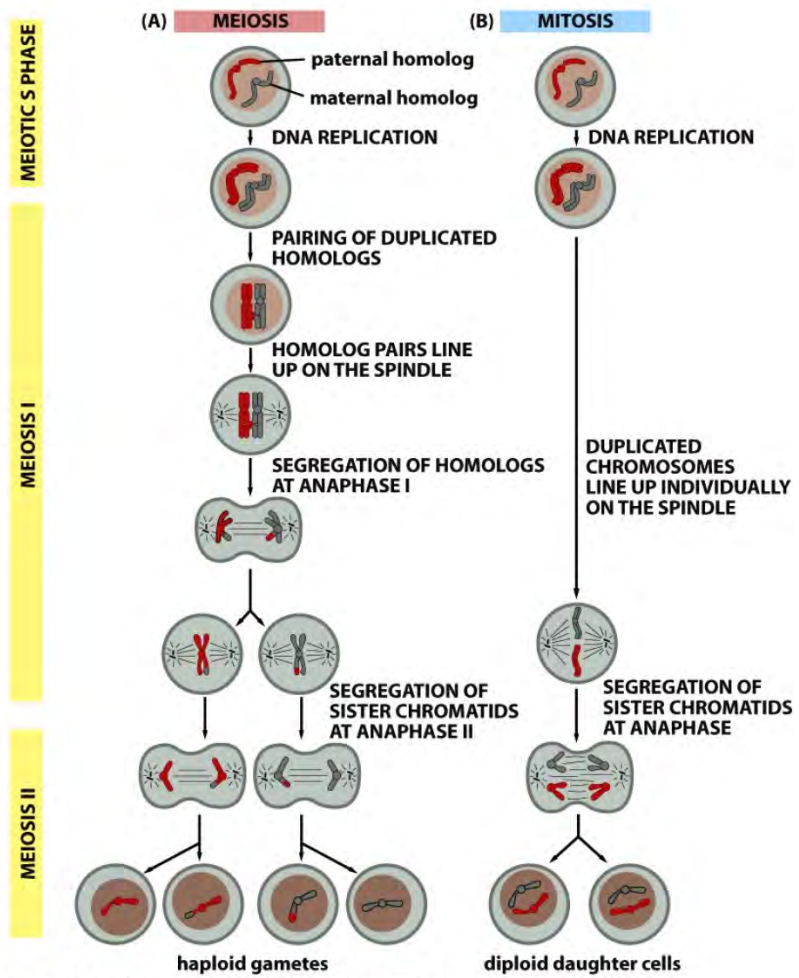


Figure 17-47 Molecular Biology of the Cell 5/e (© Garland Science 2008)

Stages of Meiosis

Meiosis I →

- i) In meiotic S-phase, DNA replication occurs
- ii) Homologous chromosomes pair and line up on the mitotic spindle; *recombination* occurs between homologous chromosomes
- iii) *Segregation* of homologous chromosomes at Anaphase I; sister chromatids remain intact; each daughter cell gets a different set of chromosomes; each daughter cell is $2n$, but contains either a maternal or paternal version of the chromosome

Meiosis II →

- i) Chromosomes are lined up *independently* on the metaphase plate
- ii) Segregation of sister chromatids in Anaphase II, forming four unique haploid gametes ($1n$)

**The Cell Cycle:
Mechanisms of
Regulation**

Phosphorylation → the enzymatic process of adding phosphate groups to target substrates to activate or inactivate them; temporary and reversible

- Kinase
 - An enzyme which adds phosphate groups to their targets
 - Phosphate group taken from ATP (making it ADP)

- Phosphatase
 - An enzyme that removes a phosphate from its target
 - Requires a water molecule to remove phosphate group

- Phosphate groups are attached to the side chains of specific amino acids
 - Only Ser, Thr, and Tyr can be phosphorylated, because they have hydroxyl as their side chain
 - Specific kinases phosphorylate specific residues on specific proteins

Phosphorylation: Kinases → heterodimeric protein kinases drive the cell cycle

- Cyclin dependent kinases (CDKs) are a catalytic subunit present throughout the cell
 - The cyclin-regulatory subunit is cyclical, and expressed at specific cell cycle stages
 - Cyclin recognises the substrate and determines the CDK's specificity
 - CDK cannot exert kinase activity without being bound to cyclin

- Distinct CDK's regulate different cell cycle transitions
 - G0: CDK's are inactive
 - CDK's are essential for progressing through the cell cycle

- A kinase assay can be used to test the activity of a CDK
 - Pull down cyclin/CDK complex using antibodies
 - Add substrate (histone, H1, protein) and radioactive ATP
 - Quantify the amount of labelled phosphate transferred to substrate on an SDS PAGE gel

Ubiquitination → a mechanism which degrades a protein; permanent and irreversible

- Ubiquitin-protein ligases attach ubiquitin to a target protein
 - Repeats multiple times, resulting in polyubiquitination
 - Proteasome recognises *polyubiquitination*, and destroys the protein

	<p>Ubiquitin-Protein Ligases →</p> <ul style="list-style-type: none"> • SCF Complex <ul style="list-style-type: none"> - Involved in the G1-S phase transition • Anaphase Promoting Complex or Cyclosome (APC/C) <ul style="list-style-type: none"> - Involved in metaphase-anaphase and anaphase-telophase transitions
<p>The G1/S Phase Transition</p>	<p>G1 Cyclin/CDK Complexes → promote S-phase entry:</p> <ul style="list-style-type: none"> • G1 Cyclin-CDKs <ul style="list-style-type: none"> - Phosphorylate transcription factors - Transcription factors drive the expression of genes that code for tools of DNA replication - Transcribed genes include enzymes to make deoxynucleotides, DNA polymerases, replication proteins and S-phase cyclins • SCF Ubiquitin-Protein Ligases <ul style="list-style-type: none"> - The boundary between G1 and S phase is defined by an inhibitor of S-phase cyclin/CDKs (Sic1) - During G1, S-phase cyclins are created and bind to their CDK's, but the complexes action is prevented by these inhibitors - However, the inhibitor serves as a substrate for the binding of G1/S cyclin-CDK's, which phosphorylates the inhibitor, making it a substrate for the ubiquitinating SCF Ubiquitin-Protein Ligase - Once the inhibitor has been ubiquitinated, the S-phase cyclin-CDK's become active - The cell is then abruptly pushed into S-phase • S-phase cyclin-CDKs promote DNA replication <ul style="list-style-type: none"> - Phosphorylates and activates numerous proteins that go onto replicate the DNA - The onset of DNA replication means that S-phase has begun - S-phase cyclin/CDKs also prepare the cell for mitosis in a similar way that G1/S CDKs play in G1 • The G1/S transition is abrupt because the S-phase cyclin-CDK inhibitor is a poor substrate <ul style="list-style-type: none"> - Therefore, requires high levels of G1/S kinase to become phosphorylated (G1/S kinase peaks mid G1 phase) - Needs to be phosphorylated on multiple sites - Makes it one of the last substrates to get phosphorylated in G1

<p>Identifying Cell Cycle Genes</p>	<p>Experimental Process →</p> <ul style="list-style-type: none"> • Yeast mutants identified crucial players in the G2/M transition, including activators and inhibitors of the mitotic cyclin/CDK • Screen for temperature-sensitive mutants <ul style="list-style-type: none"> - Mutagenize, and grow up cells at permissive temperature - Then shift them to restrictive temperature - Characterise lines that fail to grow after the temperature shift • Cell growth and cell division are uncoupled in <i>S. Pombe</i> <ul style="list-style-type: none"> - Mitosis-defective mutants thus form long rod-shaped cells - Mutants that enter mitosis prematurely show a phenotype of very small cells <p>Identified Genes →</p> <ul style="list-style-type: none"> • Cdc2 is a cyclin dependent kinase (CDK) <ul style="list-style-type: none"> - cdc2, when lost, gives a long phenotype - cdc2, when dominant, gives a small phenotype • Cdc13 is ac cyclin that forms heterodimers with cdc2 <ul style="list-style-type: none"> - Cdc13 mutants also give a long phenotype - Cdc13 = mitotic cyclin • Cdc25 drives mitosis (is a phosphatase) <ul style="list-style-type: none"> - Deficit of Cdc25 results in a long cell phenotype (increased G2) - Excess of Cdc25 results in a small cells phenotype (decreased G2) • Wee1 inhibits mitosis (is a kinase) <ul style="list-style-type: none"> - Deficit of Wee1 results in a small cell phenotype (decreased G2) - Excess of Wee1 results in a long cell phenotype (increased G2)
<p>Entry into Mitosis</p>	<p>Entry into Mitosis → controlled by a cascade of kinase and phosphatase activity</p> <ul style="list-style-type: none"> • Mitotic cyclin and CDK subunits are assembled <ul style="list-style-type: none"> - Wee1 phosphorylates Tyrosine Y15 of the CDK subunit, inactivating it - CAK phosphorylates Thymine T161, activating it (however, the inhibition of Y15 means the CDK remains inactive overall) - Cdc25, a phosphatase, reverses the phosphorylation of Y15, creating an active mitotic kinase and allowing mitosis to begin • If Wee1 is lost, there is less regulation, and the cell will enter mitosis early, making it shorter <ul style="list-style-type: none"> - If Cdc25 is lost, the cell can never enter mitosis, making it long