

### **Chapter 1 Terminology**

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1. Acetylation	Addition of a chemical group derived from acetic acid. Acetyl group are added covalently to some proteins, such as histones, as a post-translational	14. Dicer	Protein involved in the processing of miRNAs and siRNAs
		15. <b>Differentiation</b>	Process by which a cell undergoes a change to an overtly specialized cell type.
2. Amino acid	modification.  Organic molecule containing both an amino group and a carboxyl group. Those that serve as building blocks of proteins are alpha amino acids, having both the amino acid and carboxyl groups linked to the same carbon atom.	16. DNA methylation	Addition of methyl groups to DNA.  Extensive methylation of the cytosine base in CG sequences is used in vertebrates to keep genes in an inactive state.
		17. Drosha	Protein involved in the processing of microRNAs
3. Antibody	Protein produced by B cells in response to a foreign molecule or invader microorganism. Binds tightly to the foreign molecule or cell, inactivating it or marking it for destruction. Can be used as a specific reagent to detect the protein it	18. Enhancer	Regulatory DNA sequence which can increase the rate of transcription of a structural gene that can be many thousands of base pairs away.
		19. <b>Enzyme</b>	Protein that catalyses a specific chemical reaction
4. Antigen	recognizes.  A molecule that can induce and immune response or that can bind to an antibody or T-cell receptor.	20. Erythrocyte (Red blood cell)	Small hemoglobin-containing blood cell of vertebrates that transports oxygen to, and carbon dioxide from, tissues. No longer synthesizes globin or other proteins.
5. Antisense RNA	RNA complementary to the RNA transcript of a gene. Can hybridize to the specific RNA and block its function.	21. Gain-of- function mutation	Mutation that increases the activity of a gene, or makes it active in inappropriate circumstances. Usually dominant.
6. Autoradiography	Method in which the position of a radioactive probe is detected by exposure to X-ray film. Used in conjuction with blotting techniques.	22. <b>Genome</b>	The totality of genetic information belonging to a cell or an organism; in particular, the DNA that carries this information.
7. Base pair	Two nucleotides in an RNA or DNA molecule that are held together by	23. Genomic DNA	DNA constituting the genome of a cell or an organism. Often used in contrast to cDNA.
8. B cells ( B lymphocytes)	hydrogen bonds.  Type of lymphocyte that makes antibodies	24. Germ cell	A cell in the germ line of an organism, which includes the haploid gametes and their specific diploid precursor cells. Germ cells contribute to the formation of a new generation of organisms and are distinct from somatic cells, which form the body and leave no descendants.
9. Chromatin	Complex of DNA, histones and non- histone proteins found in the nucleus of a eukaryotic cell. The material of which chromosomes are made.		
10. Chromatography	Broad class of biochemical techniques in which a mixture of substances is separated by charge, size, hydrophobicity, non covalent binding affinities, or some other property by allowing the mixture to partition between a moving phase and a stationary phase.	25. <b>Germ line</b>	The cell lineage that consists of the haploid gametes and their specific diploid precursor cells.
		26. Growth factors	Extracellular signal protein that can stimulate a cell to grow. They often have other functions as well, including stimulating cells to survive or proliferate.
ıı. Complementary	Of nucleic acid sequences: capable of forming a perfect base-paired duplex	27. Heavy chains (H chains)	The larger of the polypeptide chains in an immunoglobin or other protein molecule.
12. Complementary DNA (cDNA)	with each other.  DNA copy of mRNA	28. Histone deacetylase 1 (HDAC)	Enzymes able to remove acetyl groups from histones.
13. Cytoplasm	Contents of a cell that are contained within its plasma membrane but in the case of eukaryotic cells outside the nucleus.	29. Homologous	Genes, proteins or body structures that are similar as a result of a shared evolutionary origin.

30. <b>Hormone</b>	Signal molecule secreted into bloodstream, which can then carry it to distant target cells	44. Nucleic acid	RNA or DNA; a macromolecule consisting of a chain of nucleotides joined together by phosphodiester bonds.
31. <b>HP 1</b>	Heterochromatin protein 1. Involved in producing the tightly packed structure of heterochromatin.	45. <b>Nucleus</b>	Prominent membrane-bounded organelle in eukaryotic cells, containing DNA organized into chromosomes.
32. Immune system	System of lymphocytes and other cells in	46. Peptide	Short polymer of amino acids.
	the body that provides defense against infection.	47. <b>Phenotype</b>	The observable character (including both physical appearance and behaviour) of a
33. Immunoglobulin (Ig)	An antibody molecule able to recognize specific proteins.	48. <b>piRNAs</b>	cell or organism  Piwi-interacting RNAs. Short singel-
34. <b>Insulin</b>	Polypeptide hormone that is secreted by Beta cells in the pancreas to help regulate glucose metabolism in animals.		stranded RNAs that regulate gene expression. Defined on the basis that they interact with piwi proteins and do not require the Dicer protein for their mutation.
35. Light chains (L chain)	One of the smaller polypeptides of a multi-subunit protein such as myosin or		
36. Long noncoding RNAs	immunoglobin.  (IncRNAs) RNA of 200 bases or more which cannot encode a protein but can	49. Polyadenylation	Addition of a long sequence of A nucleotides (the poly A tail) to the 3'end of a nascent mRNA molecule.
37. <b>MALDI</b>	regulate gene expression.  Matrix-assisted laser desorption-ionization method for determining the molecular	50. Polymerase	Enzyme that catalyzes polymerization reactions such as the synthesis of DNA and RNA.
	weight of a peptide.	51. Polymerase	Technique for amplifying specific regions
38. Mass spectrometry	Technique for identifying compounds on the basis of their precise mass-to-charge ratio. Powerful tool for identifying proteins and sequencing polypeptides.	chain reaction	of DNA by the use sequence-specific primers and multiple cycles of DNA synthesis, each cycle being followed by a brief heat treatment to separate complementary strands.
acid sequence of a prof eukaryotes by processii molecule made by RNA complementary copy o translated into a proteir	RNA molecule that specifies the amino acid sequence of a protein. Produced in eukaryotes by processing of an RNA molecule made by RNA polymerase as a	52. Polytene chromosomes	Giant chromosomes in which the DNA has undergone repeated replication and the many copies have stayed together.
	complementary copy of DNA. It is translated into a protein in a process catalyzed by ribosomes.	53. Post- transcriptional control	Any control of gene expression that is exerted at a stage after transcription.
40. Micro RNAs	Short eukaryotic RNAs, produced by the processing of specialized RNA transcripts coded in the genome, that regulate gene expression through complemetary base pairing with mRNA. Depending on the extent of the base pairing, miRNAs can lead either to destruction of the mRNA or	54. <b>Primers</b>	Oligonucleotide that pairs with a template DNA or RNA strand and promotes the synthesis of a new complementary strand by polymerase.
		55. <b>Promoter</b>	Region of DNA close to the transcription start site which directs transcription.
41. Northern	to a block in its translation.  Technique in which RNA fragments separated by electrophoresis are immobilized on a paper sheet, and a specific RNA is detected by hybridization with a labeled nucleic acid probe.  Method for measuring the transcription rate of a particular gene by adding radioactive precursor to isolated nuclei	56. Protease	Enzyme that degrades proteins by hydrolyzing some of the peptide bonds between amino acids.
blotting		57. Proteomics	Study of all the proteins, including all the covalently modified forms of each, produced by a cel, tissue or organism. Proteomics often investigates changes in this larger set of proteins caused by changes in the environment or by extracellularsignals.
42. Nuclear run-on assay			
43. <b>Nuclear</b>	and measuring its incorporation into the RNA transcript.	58. Pseudogene	Nucleotide sequence of DNA that has accumulated multiple mutations that have
transplantation	Transfer of a nucleus from one cell to another by microinjection.		rendered it inactive and nonfunctional.

59. Pulse labeling	Process for determining the synthesis rate of RNA (or any other molecule) by adding radioactive precursor to cells for a brief period and measuring its incorporation into the RNA.
60. Quiescent	Nondividing cells
61. Recombinant DNA	Any DNA molecule formed by joining DNA segments from different sources.
62. Regulatory RNA	RNA which cannot encode a protein but which regulates gene expression.
63. Reticulocytes	Precursor to the mature red blood cellwhich lacks the nucleus but continues to synthesize globin from long-lived mRNA.
64. Reverse transcriptase	Enzyme first discovered in retroviruses that makes a double-stranded DNA copy from a single stranded RNA template molecule.
65. <b>Ribosome</b>	Particle composed of RNAs and ribosomal proteins that catalyzes the synthesis of protein using information provided by mRNA.
66. RNA polymerase	Enzyme that catalyzes the synthesis of an RNA molecule on a DNA template from ribonucleoside triphosphate precursors. RNA polymerase I transcribes the gene encoding the 28S, 18S and 5.8S ribosomal RNA's RNA polymerase II transcribes the proteincoding genes RNA polymerase III transcribes the genes encoding transfer RNAs, 5S ribosomal RNA RNA polymerase IV and V are found ONLY in plants and are involved in transcriptional repression by siRNAs
67. <b>Robe</b>	Defined fragment of DNA or RNA, radioactively or chemically labeled, used to locate specific nucleic acid sequences by hybridization.
68. <b>siRNAs</b>	Small interfering RNAs. Short double- stranded RNAs that inhibit gene expression by directing destruction of complementary mRNAs. Production of siRNAs is triggered by double-stranded RNA.
69. Somatic cell	Any cell of plant or animal other than the germ cell.
70. Southern Blotting	Technique in which DNA fragments separated by electrophoresis on immobilized on a paper sheet. Specific fragments are then detected with a labeled nucleic acid probe.
71. Stop codons	Codons in the mRNA which produces termination of its translation rather than the insertion of another amino acid.
72. <b>Totipotent</b>	Describe a cell that is able to give rise to all the different cell types in an organism.

73. Transcriptomics	Study of all the mRNA's of a cell, tissue or organism.
74. <b>Transposons</b>	DNA element that can move from one position in the genome to another.
75. <b>Tubulin</b>	The protein subunit of microtubules
76. URNAs	Family of small uridine-rich RNAs. Several such s U1, U2, U4, U5 and U6 are involved in RNA splicing. Also known as snRNA.
77. <b>Variable region</b>	Region of an immunoglobin light or heavy chain that differs from molecule to molecule and forms the antigen binding site.
78. Western blotting/ immunoblotting	Techniques by which proteins are separatedny electrophoresis and immobilized on a paper sheet and then analyzed, usually by means of a labeled antibody.
79. <b>Yeast</b>	Common name for several families of unicellular fungi. Includes species used for brewing and bread-making, as well as pathogenic species. Among the simplest of Eukaryotes.

# Quizlet

### **Chapter 3 terminology**

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5-azacytidine	Analog of the C nucleotide in DNA which cannot be methylated. Used to probe the consequences of the loss of DNA methylation on C residues.	13	13. Cyclic AMP  14. DNase I hypersensitive sites	(cAMP) Nucleotide that is generated from ATP by adenylyl cyclase in response to various extracellular signals. It acts as small intracellular signaling molecule, mainly by activating cAMP-dependent protein kinase.
2. Active demethylation	Process in which DNA is actively demethylated by TET enzymes			It is hydrolyzed to AMP by a phosphodiesterase.
3. <b>Air</b>	Long noncoding RNA which regulates the genomic imprinting of several genes	14		Sites within chromatin which are highly sensitive to digestion with DNasel. Frequently located at target sites for
4. Argonaute	Family of proteins found in the RISC and RITS complexes which play a key role in the repression of gene		Epigenome	transcription factor binding.  Features such as DNA methylation or histone modification which are
5. Barr body	expression by miRNAs and siRNAs Inactive X chromosome whose very			superimposed onto the genomic DNA sequence
	condensed chromatin structure makes it visible by microscopy.	16	16. Estrogen response	DNA sequence which binds the estrogen receptor and mediates the activation of
<ul><li>Basal transcription factors</li></ul>	Any of the proteins whose assembly at the promoter is required for the		element (ERE)	genes in response to estrogen
	binding and activation of RNA polymerase and the initiation of transcription.	d activation of RNA 17. <b>GAGA factor</b> e and the initiation of	GAGA factor	Transcription factor involved in chromatin remodeling/ gene activation. A member of the trithorax family
7. Bromodomain	Protein domain found in many proteins which can convert chromatin to a more open structure	18	Genetic code	Set of rules specifying the correspondence between nucleotide triplets (codons) in DNA or RNA and amino acids in proteins
8. <b>CBP</b>	CREB-binding protein. Transcriptional co-activator for a variety of transcriptional activators. Originally identified on the basis of its binding to	19	Genomic imprinting	Phenomenon in which a gene is either expressed or not expressed in the offspring depending on which parent it is inherited from.
Chromatin	the CREB transcription factor.	20. Glycosylation	. Glycosylation	Modification of histones and other proteins by addition of sugar residues
immunoprecipitation		21	GRE (glucocorticoid response element)	DNA sequence which binds the glucocorticoid receptor and mediates the activation of genes in response to glucocorticoid
10. Chromodomain	Protein domain found in many proteins which can convert chromatin to a more tightly packed structure.	22	Heat-shock factor	(HSF) Transcription factor which is activated by heat or other stresses when activated. It binds to the heat shock element in the genes encoding heat shock proteins and
11. Citrullination	Modification of arginine residues in histones and other proteins to produce the amino acid citrulline which is not coded for in the genetic code	23	23. Helix-turn- helix	DNA binding structural motif present in
				many gene regulatory proteins, consisting of two alpha-helices held at a fixed angle and connected by a short chain of amino acids, constituting the turn. Distinct from the helix-loop-helix motif.
12. CpG island	Region of DNA with a greater than average density of CG sequences; these regions generally remain unmethylated			
		24	4. Homologous chromosomes	The maternal and paternal copies of a particular chromosome in a diploid cell

25. <b>Hybridization</b>	In molecular biology, the process whereby two complementary nucleic acid strands form a base-paired duplex DNA-DNA, DNA-RNA, RNA-RNA molecule. Forms the basis of a powerful experimental technique for detecting specific nucleotide sequences.	41. Steroid	Hydrophobic lipid molecule with a characteristic four-ringed structure; derived from cholesterol. Many important hormones such as glucocorticoid, estrogen and testosterone are steroids that activate intracellular nuclear receptors
26. Inner cell mass	Cluster of undifferentiated cells in the early mammalian embryo from which the whole of the adult body is derived	42. <b>SV40</b>	Simian virus 40. Small DNA virus which infects monkeys and has been widely used in studies of eukaryotic gene regulation
27. Maintenance methylase	Enzyme which can recognize and methylate CG sites in DNA which are methylated only on one strand of DNA. It then fully methylates them.	43. <b>TATA Box</b>	AT-rich sequence in the promoter region of many eukaryotic genes that bind the general transcription factor TBP and hence specifies the position at which transcription is initiated.
28. <b>MeCP2</b>	Regulatory protein which binds specifically to methylated DNA		
29. <b>MeF2</b>	Myocyte enhancer factor two Family of transcription factors	44. <b>TBP</b>	Transcription factor which plays a key role in transcription by all three RNA polymerases. Binds to the TATA box
30. <b>OGT enzyme</b>	Enzyme able to glycosylate histones and other proteins by addition of O-linked N-acetylglucosamine	45. <b>TET</b>	Ten eleven translocase enzymes which are able to produce the active
31. Oligonucleotides	Short DNA sequence		demethylation of methylated C- residues in DNA
32. Oogenesis	Formation and maturation of oocytes in the ovary	46. Transdifferentiation	Process in which one differentiated cell type can differentiate, proliferate, and then differentiate into a different cell type.
33. <b>p300</b>	Transcriptional co-activator for a variety of transcriptional activators. Closely		
	related to CBP. Named on the basis of its molecular weight of 300 kDa	47. <b>TSIX</b>	Long noncoding RNA which is transcribed from the active X-
34. Pioneer transcription factors	Regulatory protein which can bind to tightly packed chromatin and open its structure so that other regulatory		chromosome utilizing the opposite DNA strand to that used to probe the XIST RNA
35. Polycomb complex	multiprotein complex which produces a closed chromatin structure, incompatible	48. X Inactivation	Inactivation of one copy of the X chromosome in the somatic cells of female mammals
36. Regulatory gene	processes such as gene transcription or RNA splicing.	49. X-inactivation center	(XIC) Site in an X chromosome at which inactivation is initiated and spreads outwards
		50. <b>XIST</b>	Long noncoding RNA whose transcription initiates in the X-inactivation center and which is critical
37. Restriction fragment	Fragment of DNA generated by the action of restriction enzyme(s)		
38. <b>RISC</b>	RNA-induced-silencing-complex. Protein complex which plays a key role in post- transcriptional repression by miRNAs and siRNAs		for the process of X-inactivation
39. <b>RITS</b>	RNA-induced-transcriptional-silencing- complex. Protein complex which play a key role in transcriptional repression by siRNAs		
40. Small regulatory RNAs	Small RNAs between 20 and 31 bases in length which regulate gene expression		

## **NORTHERN BLOTTING**

Method

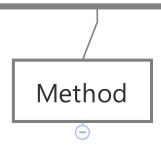
1. Isolate total mRNA from cell types A and B



3. Transfer mRNA to nylon membrane

4. Label a gene-specific probe to detect mRNA by hybridisation

## **SOUTHERN BLOTTING**



1. Isolate DNA from various tissue types

2. Digest DNA with restriction enzyme(s) gel electrophoresis to obtain thousands of fragments

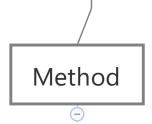
3. Separate DNA fragments from samples by agarose gel electrophoresis

4. Analyse all samples for comparative purposes

5. Transfer to membrane

6. Hybridize with gene-specific probe

### **WESTERN BLOTTING**



- 1. Obtain proteins from various tissue types
  - 2. Denature proteins with heat & SDS
    - 3. Separate proteins by SDS-PAGE
    - 4. Transfer proteins to membrane
    - 5. Submerge membrane in a solution containing the monospecific antigen against protein-X

6. Visualise binding