

| Question | | Expected Answers | | | | Marks | Additional Guidance | |
|----------|-----|------------------|---|--------------|---------------|------------------|---|---|
| 1 | (a) | | Animal | Plant | Yeast | Bacterium | <p>Mark the first answer in each box. If the answer is correct and a further answer is given that is incorrect or contradicts the correct answer then = 0 marks</p> <p>Award 1 mark for each correct row</p> <p>ACCEPT tick / present & cross / not present / absent / none</p> <p>IGNORE ref to nucleoid</p> <p>CREDIT murein as alternative to peptidoglycan ACCEPT peptidoglycin DO NOT ACCEPT peptoglycan</p> <p>ACCEPT 'on RER' or 'in cytoplasm' for yes ACCEPT ref to size of ribosomes (large / 80S / 22nm in Eukaryotes, small / 70S / 18nm in bacteria)</p> | |
| | | | | | budding | | | ; |
| | | yes | yes | yes | no | | | ; |
| | | | cellulose | | peptidoglycan | | | ; |
| | | yes | yes | yes | yes | ; | 4 | |
| | (b) | (i) | <u>meristem</u> (atic) ; | | | | 1 | IGNORE position in plant such as 'root tip', cambium |
| | (b) | (ii) | nucleus / nucleolus / chromatin ; cytoplasm ; cross / end, (cell) walls ; | | | | 2 max | <p>Read through and award marks for correct features</p> <p>IGNORE ref to other individual organelles / vacuole</p> <p>IGNORE nucleous</p> <p>DO NOT CREDIT 'two nuclei in one cell'</p> <p>CREDIT end plates</p> <p>ACCEPT no end walls / no nucleus / no cytoplasm</p> <p>IGNORE walls between cells</p> |

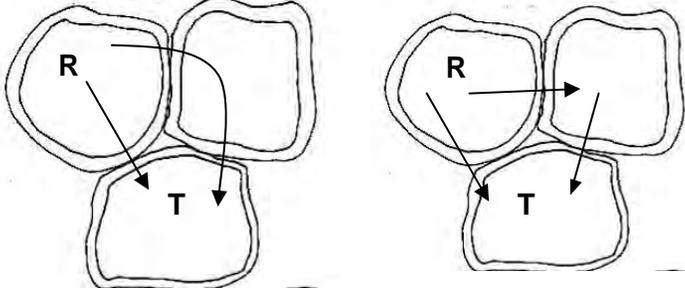
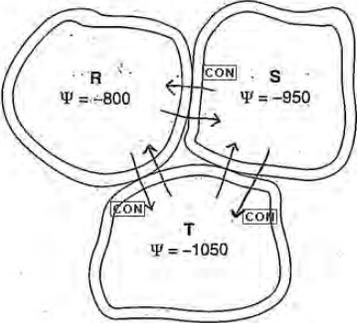
| Question | | Expected Answers | Marks | Additional Guidance |
|--------------|-----------|---|-----------|--|
| | (b) (iii) | <p>thicker ;</p> <p>lignified ;</p> <p>contain (bordered) pits ;</p> | 2 max | <p>IGNORE stronger</p> <p>CREDIT have lignin / contain lignin / reinforced with lignin / impregnated with lignin</p> <p>DO NOT CREDIT have lignin on the walls / lined by lignin / surrounded by lignin</p> <p>IGNORE ref to pattern of thickening</p> <p>IGNORE 'pore'</p> |
| | (c) | <p><u>sieve (tube) element</u> ;</p> <p><u>companion</u> (cell) ;</p> <p>parenchyma ;</p> | 2 max | <p>IGNORE 'sieve tube' 'sieve cell'</p> <p>ACCEPT fibres / sclereids / sclerenchyma</p> |
| Total | | | 11 | |

| Question | | | Answer | Marks | Guidance |
|----------|-----|------|---|-------|---|
| 2 | (a) | (i) | <p><i>division type 1</i> <u>mitosis</u></p> <p>and</p> <p><i>division type 2</i> <u>meiosis</u> ;</p> | 1 | <p>Mark the first answer on each prompt line. If the answer is correct and an additional answer is given that is incorrect or contradicts the correct answer then = 0 marks</p> <p>ACCEPT correct spelling only</p> <p>ACCEPT correct spelling only CREDIT meiosis I and II DO NOT CREDIT meiosis I / meiosis II alone</p> |
| 2 | (a) | (ii) | <p>A (DNA) replication ;</p> <p>B cytokinesis ;</p> | 2 | <p>Mark the first answer on each prompt line. If the answer is correct and an additional answer is given that is incorrect or contradicts the correct answer then = 0 marks</p> <p>IGNORE stages of cell division</p> <p>IGNORE cell division / stages of cell division</p> |

| Question | | Answer | Marks | Guidance |
|----------|-----|--|-------|--|
| 2 | (b) | <p>A1 independent assortment / random segregation , of (homologous) chromosomes / bivalents;</p> <p>A2 in , metaphase I / meiosis I ;</p> <p>A3 of chromatids in , metaphase II / meiosis II ;</p> <p>A4 (so) homologous chromosomes , have different alleles / come from different parents ;</p> <p>A5 produces large number of allele combinations ;</p> <p>C1 <u>crossing over</u> / (formation of) chiasma(ta) ;</p> <p>C2 in , prophase I / meiosis I ;</p> <p>C3 (so) <u>chromatids</u> will have new combination of <u>alleles</u> ;</p> <p>C4 amount of variation depends on distance between crossover points ;</p> <p>M1 mutation ;</p> <p>M2 changes the (DNA) nucleotide/ base, sequence ;</p> <p>M3 DNA checks (during duplication) did not recognise damage ;</p> <p>M4 <i>idea of</i> differences in (named) protein(s) ;</p> <p>N1 non-disjunction ;</p> <p>N2 homologous chromosomes do not separate (in metaphase I) ;</p> <p>N3 one , more / less , chromosome present ;</p> <p>F1 random, mating / fusion of gametes/ fertilisation ;</p> <p>F2 gametes are not genetically identical;</p> <p>F3 produces large number of (allele) combinations ;</p> | 8 max | <p>A1 ACCEPT Random assortment / independent segregation</p> <p>A2 /A3 DO NOT CREDIT metaphase /meiosis, I and II</p> <p>A2 /A3 ACCEPT correct anaphase stage linked to segregation</p> <p>A2 must be in context of independent assortment / random segregation</p> <p>A4/ A5 DO NOT CREDIT genes</p> <p>A4 ACCEPT pairs of chromosomes / maternal and paternal chromosomes, have different alleles/ come from different parents</p> <p>A5 ACCEPT different combinations of, chromatids /chromosomes, in gametes</p> <p>CREDIT figures e.g. for humans 2^{23} possible combinations</p> <p>C1 DO NOT CREDIT between sister chromatids</p> <p>C2 DO NOT CREDIT prophase / meiosis, I and II</p> <p>C2 must be in context of crossing over</p> <p>C3 ACCEPT shuffles / swaps/exchanges, <u>alleles</u> on <u>chromatids</u></p> <p>C4 e.g. more variation the further apart the crossovers occur</p> <p>M2 IGNORE 'pairs'</p> <p>M2 CREDIT deletion,/substitution/ addition, of, base / nucleotide</p> <p>M3 ACCEPT proof reading did not recognise damage</p> <p>M4 e.g. change in, amino acid sequence/primary structure</p> <p>N1 CREDIT inversion / translocation (chromosome mutation)</p> <p>N2 CREDIT description of inversion / translocation</p> <p>N3 CREDIT examples of chromosome changes e.g. Trisomy 21</p> <p>F2 ACCEPT gametes are genetically different</p> <p>F3 DO NOT CREDIT produce large number of gene combinations</p> |

| Question | | | Answer | Marks | Guidance |
|----------|--|--|--------------|-----------|--|
| | | | QWC ; | 1 | <p>Awarded for one change and consequence of that change</p> <p>Award if ONE of the following has been awarded</p> <p>mp A1 or A2 or A3 <u>and</u> mp A4 or A5</p> <p>OR</p> <p>mp C1 or C2 <u>and</u> mp C3 or C4</p> <p>OR</p> <p>mp M1 or M2 <u>and</u> mp M3 or M4</p> <p>OR</p> <p>mp N1 or N2 <u>and</u> mp N3</p> <p>OR</p> <p>mp F1 or F2 <u>and</u> mp F3</p> |
| | | | Total | 12 | |

| Question | | | Answer | Mark | Guidance |
|--------------|-----|-------|--|----------|--|
| 3 | (a) | (i) | mitosis ; | 1 | CREDIT correct spelling only ACCEPT binary fission |
| | | (ii) | in the grex / 3 ; | 1 | |
| | (b) | (i) | cell signalling ; | 1 | |
| | | (ii) | 1 attraction of <u>cell(s)</u> to folic acid from bacteria ; 2 attraction of <u>cells</u> to each other by cAMP ; 3 coordinated movement in grex ; 4 differentiation / described, of (grex / slime mould) <u>cells</u> in response to DIF ; | 2 max | NOTE must name the chemical involved for description (except mp 3 coordinated movement) ACCEPT attraction of cells to bacteria by folic acid IGNORE makes cells stick together |
| | | (iii) | contains , receptors / glycoproteins / glycolipids / glycocalyx ; for , folic acid / cAMP / DIF ; | 2 | DO NOT CREDIT <i>consists</i> of receptors |
| | (c) | | 17 (hours) ; | 1 | |
| Total | | | | 8 | |

| | | | |
|-------------------|---|---------------------|---|
| | <p>(iii) arrow from R to T ;</p> <p>arrow from R to S AND arrow from S to T OR arrow from R to S to T ;</p> | | <p>e.</p>  <p>If contradictory arrows to the above are drawn, apply CON for each arrow going from low Ψ to high Ψ.</p> <p>e.</p>  <p>2 gets 0</p> |
| <p>(b)</p> | <p>this is where cambium / meristem / xylem / phloem / vascular bundle, is found ;</p> <p>mitosis/cell division, occurs in cambium (to produce new cells for growth) ;</p> <p>new cells, differentiate / specialise, (into xylem and phloem) ;</p> <p>xylem supplies water for, (cell) elongation / (cell) growth ;</p> <p>phloem supplies, sugars / assimilates, for, energy / growth /respiration ;</p> | <p>max 2</p> | <p>CREDIT from a labelled diagram</p> <p>CREDIT description of position being close to the edge of trunk</p> <p>DO NOT CREDIT responses that suggest that cambium etc. are in or outside bark OR under cut surface</p> <p>ACCEPT cambium differentiates</p> <p>IGNORE nutrients</p> |

| | | | | |
|--|------------|--|------------------|---|
| | (c) | <p>tip / apex, of, shoot / root ;</p> <p>meristem ;</p> <p>bud ;</p> | <p>max 1</p> | <p>Mark the first answer on each prompt line. If the answer is correct and an additional answer is given that is incorrect or contradicts the correct answer then = 0 marks</p> <p>IGNORE root or shoot unqualified</p> <p>ACCEPT behind root tip</p> |
| | (d) | <p>allow <u>oxygen</u> to reach, cells / tissues (under bark) ;</p> <p>for (aerobic) respiration ;</p> <p>animals transport oxygen in, blood / circulation / transport system ;</p> <p>plants do not transport (much) oxygen in transport system ;</p> <p><i>idea that</i> (oxygen not supplied from leaves as) stomata only open in day / no leaves in winter ;</p> | <p>max 2</p> | <p>IGNORE refs to need for CO₂ / photosynthesis throughout</p> <p>ACCEPT correct formula O₂</p> <p>DO NOT CREDIT oxygen for photosynthesis</p> <p>ACCEPT gas(es) for oxygen</p> <p>ACCEPT gas(es) for oxygen</p> |
| | | <p>Total</p> | <p>10</p> | |

| Question | | Answer | Marks | Guidance |
|--------------|---------|--|-----------|---|
| 5 | (a) | metaphase I and metaphase II ; prophase I ; anaphase II ; telophase II ; anaphase I ; | 5 | Mark the first answer on each prompt line. If the answer is correct and an additional answer is given that is incorrect or contradicts the correct answer then = 0 marks |
| | (b) | to, halve chromosome number / reduce from 2n to n ; to separate homologous pairs (of chromosomes) and sister chromatids ; because, DNA (previously) replicated / chromosomes are two chromatids at start ; | 2 | IGNORE all references to mitosis CREDIT 'from diploid to haploid' ACCEPT 'from 46 to 23 chromosomes' IGNORE halve, genetic material / DNA ACCEPT genetic, material / information |
| | (c) (i) | sequence / order, of bases / nucleotides ; | 1 | CREDIT base pairs DO NOT CREDIT amino acid sequence |
| | (ii) | different, primary / secondary / tertiary, structure ; (protein) shorter due to, deletion / stop codon OR longer due to, insertion / duplication ; (protein) unchanged due to, silent mutation / non-coding DNA altered ; (function is) lost / worse / better ; | 3 | ACCEPT different <u>sequence</u> or <u>order</u> of amino acids ACCEPT different 3D folding or 3D shape for 'silent' CREDIT 'neutral' or a description of more than one triplet coding for one amino acid IGNORE different / altered function ACCEPT idea that change is harmful |
| Total | | | 11 | |