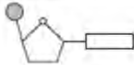


Question Number	Answer	Additional Guidance	Mark
1(a)(i)	C 		(1)

Question Number	Answer	Additional Guidance	Mark
1(a)(ii)	B G-A-U-U-C-A-C-G-U		(1)

Question Number	Answer	Additional Guidance	Mark
1(a)(iii)	C 3 1 4 2		(1)

Question Number	Answer	Additional Guidance	Mark
1(b)(i)	8		(1)

Question Number	Answer	Additional Guidance	Mark
1(b)(ii)	6		(1)

Question Number	Answer	Additional Guidance	Mark
1(c)	<ol style="list-style-type: none"> <li>DNA is {double stranded/ has a double helix} and RNA is {single stranded / does not have a double helix} ;</li> <li>DNA has {thymine / T} while RNA has { uracil / U } ;</li> <li>DNA has deoxyribose while RNA has ribose ;</li> <li>DNA is { larger / longer } than RNA / eq ;</li> </ol>	<p><b>1. ACC T</b> mixtures e.g. DNA double helix mRNA is single strand <b>IGNORE</b> hydrogen bonds</p> <p><b>2.NOT</b> thiamine, thymine</p>	(3)

Question Number	Answer	Additional Guidance	Mark
2(a)	idea that the (RNA) nucleotides attach to this strand OR idea of {nucleotide / base } sequence that directs the synthesis of {complementary sequence / mRNA / eq} ;	ACCEPT complementary to RNA nucleotides, codes for mRNA, {part of the DNA / antisense } strand that the mRNA is built along, NOT DNA nucleotides, plural strands	(1)

Question Number	Answer	Mark
2(b)(i)	D have a sugar-phosphate chain ;	(1)

Question Number	Answer	Mark
2(b)(ii)	C semi-conservative replication is possible ;	(1)

Question Number	Answer	Mark
2(b)(iii)	A 10% ;	(1)

Question Number	Answer	Additional Guidance	Mark												
2(c)	<table border="1" style="margin-left: 20px;"> <tr> <td>U</td><td>G</td><td>A</td><td>A</td><td>A</td><td>G</td><td>C</td><td>G</td><td>G</td><td>G</td><td>C</td><td>U</td> </tr> </table> <p>1. both uracils correct ; 2. the rest of the sequence correct ;</p>	U	G	A	A	A	G	C	G	G	G	C	U		(2)
U	G	A	A	A	G	C	G	G	G	C	U				

Question Number	Answer	Additional Guidance	Mark																		
2 (d)	<p>Any three from:</p> <table border="1" data-bbox="306 329 946 792"> <thead> <tr> <th></th> <th>replication</th> <th>transcription</th> </tr> </thead> <tbody> <tr> <td>1</td> <td>uses DNA nucleotides</td> <td>uses RNA nucleotides / eq ;</td> </tr> <tr> <td>2</td> <td>uses DNA polymerase / eq</td> <td>does not use DNA polymerase / uses RNA polymerase ;</td> </tr> <tr> <td>3</td> <td>reference to semi-conservative</td> <td>not semi-conservative / eq ;</td> </tr> <tr> <td>4</td> <td>(copies) both DNA strands / eq</td> <td>(copies) only {one strand / template / gene / eq} ;</td> </tr> <tr> <td>5</td> <td>makes DNA double helix / eq</td> <td>Makes single strand mRNA / eq ;</td> </tr> </tbody> </table>		replication	transcription	1	uses DNA nucleotides	uses RNA nucleotides / eq ;	2	uses DNA polymerase / eq	does not use DNA polymerase / uses RNA polymerase ;	3	reference to semi-conservative	not semi-conservative / eq ;	4	(copies) both DNA strands / eq	(copies) only {one strand / template / gene / eq} ;	5	makes DNA double helix / eq	Makes single strand mRNA / eq ;	<p>Must be clearly comparative for the mark  <b>IGNORE</b> destination of the molecules</p> <p>1. <b>ACCEPT</b> thymine / T, uracil / U comparison, deoxyribose and ribose, DNA and RNA bases  2. <b>ACCEPT</b> no ligase in transcription</p> <p>4. <b>ACCEPT</b> whole DNA molecule unzipped for replication with only part for transcription  <b>ACCEPT</b> all {DNA / genome} copied in replication only part in transcription  5. <b>NOT</b> just produces DNA and mRNA  <b>ACCEPT</b> two {new strands of DNA / DNA molecules} compared to one mRNA (each time)  <b>ACCEPT</b> if clear what is being produced elsewhere in the response</p>	<b>(3)</b>
	replication	transcription																			
1	uses DNA nucleotides	uses RNA nucleotides / eq ;																			
2	uses DNA polymerase / eq	does not use DNA polymerase / uses RNA polymerase ;																			
3	reference to semi-conservative	not semi-conservative / eq ;																			
4	(copies) both DNA strands / eq	(copies) only {one strand / template / gene / eq} ;																			
5	makes DNA double helix / eq	Makes single strand mRNA / eq ;																			