

الصفحة	الامتحان الوطني الموحد للبكالوريا المسالك الدولية الدورة العادية 2020 - عناصر الإجابة -		الجمهورية المغربية وزارة التربية الوطنية والتكوين المهني والتعليم العالي والبحث العلمي المركز الوطني للتقويم والامتحانات
1			
4	SSSSSSSSSSSSSSSSSSSSSSSSSSSSSS		NR 32E

3	مدة الإنجاز	علوم الحياة والأرض	المادة
7	المعامل	شعبة العلوم والتجريبية مسلك علوم الحياة والأرض (خيار إنجليزية)	الشعبة أو المسلك

Key and marking scale

Questions	Response elements	Scores
Section I : Knowledge Retrieval (5 pts)		
I	a- definition (accept any correct definitions) Genetic engineering: set of techniques used to modify genetic cell or organism in order to express new traits	0.5
	b- two example of application of genetic engineering: -in agricultural field: producing plants resistant to pests	0.25
	-in medical field: producing human insulin.....	0.25
II	(1, b) ; (2, c) ; (3, b) ; (4, c)	0.5x4
III	1- true 2- false 3- false 4- false	0.25x4
IV	(1, b) ; (2, a) ; (3, d) ; (4, c)	0.25x4
Section II : Scientific reasoning and communication in graphic and written modes (15 pts)		
Exercise 1 (5 pts)		
1	Necessary conditions of muscle contraction: Presence Ca ⁺⁺ ions.....	0.25
	ATP hydrolysis.....	0.25
	Justification (any correct justifications is accepted) By inhibiting ATP hydrolysis of ATP (medium 2), the myofibrils don't contract...	0.5
	By inhibiting the action of Ca ⁺⁺ ions (medium 3) the myofibrils don't contract....	0.5
2	Ca ⁺⁺ ions in sarcoplasmic reticulum → relaxed fibers	0.25
	Ca ⁺⁺ ions in sarcoplasm → contracted fibers	0.25
3	Explanation of the evolution of tension of muscle fiber During contraction phase (phase A)	0.25x6
	- fixation Ca ⁺⁺ ions on the troponin and tropomyosin displacement → exposing the binding sites to the myosin heads in actin and forming the actin-myosin complex → release the ADP and Pi → the myosin heads swiveling and sliding of myofilaments (contraction of the muscle fiber) → the actin-myosin complex is dissociated by ATP → ATP hydrolysis and myosin heads returning to their original status and resume the cycle of contraction.	
	During relaxation phase (phase C):	0.25x2
	In the absence of Ca ⁺⁺ ions, the actin-myosin complex is not formed → relaxation of muscle fiber even though the ATP is present.	

الصفحة 2 4	NR 32E	<p style="text-align: center;">الامتحان الوطني الموحد للبكالوريا - الدورة العادية 2020 - عناصر الإجابة - مادة: علوم الحياة والأرض - شعبة العلوم التجريبية مسلك علوم الحياة والأرض (خيار إنجليزية)</p>																					
4		<p>Explanation of the Rigor Mortis</p> <p>Exhaustion and non-renewal of the ATP after death → the actin-myosin complex is not dissociated (document 3) → stop of muscle contraction cycle in contraction phase (document 3) → maintaining strong muscle tension (Phase B of document 2) causing rigor mortis.</p>	0.25x4																				
Exercise 2 (6.5 pts)																							
1		<p>Protein-trait relationship:</p> <ul style="list-style-type: none"> - With normal Endoglin, the complex “membrane receptor-growth factor” is functional so a normal angiogenesis → healthy person - With abnormal Endoglin, the complex “membrane receptor-growth factor” is dysfunctional so an abnormal angiogenesis → person with ROW disease..... - So any modification in the protein Endoglin causes a modification in the trait (healthy or sick person)..... 	<p>0.25</p> <p>0.25</p> <p>0.25</p>																				
2		<p>mRNA sequences corresponding to:</p> <ul style="list-style-type: none"> - the normal allele fragment : <li style="padding-left: 40px;">CCC-CAC- GUG- GAC-AGC-AUG-GAC-CGC - the abnormal allele fragment : <li style="padding-left: 40px;">CCC-CAC- AUG- GAC-AGC-AUG-GAC-CGC <p>Amino acids sequences corresponding to :</p> <ul style="list-style-type: none"> - the normal allele fragment : <li style="padding-left: 40px;">Pro-His-Val-Ac.asp-Ser-Met- Ac.asp -Arg - the abnormal allele fragment: <li style="padding-left: 40px;">Pro-His-Met- Ac.asp -Ser-Met- Ac.asp -Arg <p>Mutation by substitution of first nucleotide G for A at the level of third triplet of un transcribed strand (DNA) → incorporation the Met instead of Val in amino acid sequence → synthesis abnormal Endoglin protein → abnormal angiogenesis (the ROW disease appear).</p>	<p>0.25</p> <p>0.25</p> <p>0.25</p> <p>0.25</p> <p>0.5 pt</p>																				
3		<p>a. The responsible allele for disease is dominant and the studied gene is carried by an autosome:</p> <p>The daughter III₁ is healthy phenotype while her parents II₅ and II₆ are sick phenotype → parents are heterozygous → responsible allele for disease is dominant. (Accept also the answer: any person affected must descending from affected person).....</p> <ul style="list-style-type: none"> -The disease is present in both sexes → the responsible allele is not carried by chromosome Y. -The daughter III₁ is healthy, her father II₅ is sick and responsible allele for disease is dominant → the girl III₁ will inherit of her father the responsible allele for diseases so the daughter should be affected → the responsible allele for disease is not carried by chromosome X (accept any correct answers)..... →the responsible allele for disease is not carried by chromosome X or chromosome Y so the responsible allele for disease is carried by autosome..... <p>b. the probability for that couple II₈ et II₉ to give birth to healthy child:</p> <table style="width: 100%; border-collapse: collapse;"> <tr> <td style="width: 30%;">Parents :</td> <td style="width: 20%;">II₈ ♂</td> <td style="width: 10%; text-align: center;">x</td> <td style="width: 20%;">II₉ ♀</td> <td style="width: 10%;"></td> </tr> <tr> <td>Phenotypes :</td> <td>[r]</td> <td></td> <td>[R]</td> <td></td> </tr> <tr> <td>Genotypes :</td> <td>r//r</td> <td></td> <td>R//r</td> <td></td> </tr> <tr> <td>Gametes :</td> <td>r/ 1</td> <td></td> <td>R/ ½ r/ ½</td> <td></td> </tr> </table>	Parents :	II ₈ ♂	x	II ₉ ♀		Phenotypes :	[r]		[R]		Genotypes :	r//r		R//r		Gametes :	r/ 1		R/ ½ r/ ½		<p>0.5</p> <p>0.25</p> <p>0.25</p> <p>0.25</p> <p>0.25x2</p>
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3		Punnet square: <table border="1" style="margin: 10px auto; border-collapse: collapse;"> <tr> <td style="text-align: center;">Gametes ♂</td> <td style="text-align: center;">r</td> <td style="text-align: center;">1</td> </tr> <tr> <td style="text-align: center;">♀</td> <td style="text-align: center;">R</td> <td style="text-align: center;">(R/r) [R]</td> </tr> <tr> <td style="text-align: center;"></td> <td style="text-align: center;">1/2</td> <td style="text-align: center;">1/2</td> </tr> <tr> <td style="text-align: center;">r</td> <td style="text-align: center;">(r/r) [r]</td> <td style="text-align: center;">1/2</td> </tr> <tr> <td style="text-align: center;"></td> <td style="text-align: center;">1/2</td> <td style="text-align: center;">1/2</td> </tr> </table>	Gametes ♂	r	1	♀	R	(R/r) [R]		1/2	1/2	r	(r/r) [r]	1/2		1/2	1/2	0.25
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r	(r/r) [r]	1/2																
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4		The probability for that couple II ₈ et II ₉ to give birth to healthy child is 1/2 a. The frequency of the normal allele and abnormal allele we have : $f([R]) = p^2 + 2pq = 1/5000$ we know $p^2 + 2pq + q^2 = 1$ So $q^2 = 1 - 1/5000 = 0.9998$ - Normal allele frequency is: $f(r) = q = \mathbf{0.9998}$ - Abnormal allele frequency is: $f(R) = p = 1 - q = \mathbf{0.0002}$ b. The frequencies of different genotypes in studied population. $f(r/r) = q^2 \approx 0.9998$ $f(R/r) = 2pq \approx 0.0003$ $f(R/R) = p^2 \approx 0$	0.25 0.5 0.5 0.25 0.25 0.25															
Exercise 3 (5 pts)																		
1		Deduction and justification: -We study transmission of a hereditary trait for each cross → monohybrid cross... -The descending of two crosses are homogenous → the parents are for pure lineage according to Mendel's first law -The descendants of the first cross have erect ears → responsible allele for erect ears form is dominant (D) and responsible allele for floppy ears is recessive (d)... -The descendants of the first cross have light muzzle → responsible allele for light muzzle is dominant (S) and responsible allele for dark muzzle is recessive (s).....	0.25 0.25 0.25 0.25															
2		The test cross gives two parental phenotypes with a percentage 83% upper to percentage of recombined phenotype 17% (Mendel's third law is not verified) → The two studied genes are linked..... Deduction : the parental genotype The genotype of sheep with dominant phenotype : <table style="display: inline-table; vertical-align: middle;"><tr><td style="border: 1px solid black; padding: 2px;">D</td><td style="border: 1px solid black; padding: 2px;">S</td></tr><tr><td style="border: 1px solid black; padding: 2px;">d</td><td style="border: 1px solid black; padding: 2px;">s</td></tr></table> The genotype of double recessives sheep : <table style="display: inline-table; vertical-align: middle;"><tr><td style="border: 1px solid black; padding: 2px;">d</td><td style="border: 1px solid black; padding: 2px;">s</td></tr><tr><td style="border: 1px solid black; padding: 2px;">d</td><td style="border: 1px solid black; padding: 2px;">s</td></tr></table>	D	S	d	s	d	s	d	s	0.5 0.25 0.25							
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3		<p>Chromosomal interpretation of the third cross with Punnet square:</p> <p>Phenotypes : ♀ [D,S] x [d, s] ♂</p> <p>Genotypes : $\frac{D}{d} \frac{S}{s}$ x $\frac{d}{d} \frac{s}{s}$</p> <p>Gametes : 45% $\frac{D}{d} \frac{S}{s}$ 100% $\frac{d}{d} \frac{s}{s}$</p> <p style="text-align: center;">38% $\frac{d}{d} \frac{s}{s}$</p> <p style="text-align: center;">9% $\frac{D}{d} \frac{s}{s}$</p> <p style="text-align: center;">8% $\frac{d}{d} \frac{S}{s}$</p> <p>Punnet square :</p> <table border="1" style="width: 100%; border-collapse: collapse; text-align: center;"> <tr> <td style="width: 15%;">$\gamma^{\text{♀}}$</td> <td style="width: 15%;">$\frac{D}{d} \frac{S}{s}$</td> <td style="width: 15%;">$\frac{d}{d} \frac{s}{s}$</td> <td style="width: 15%;">$\frac{D}{d} \frac{s}{s}$</td> <td style="width: 15%;">$\frac{d}{d} \frac{S}{s}$</td> </tr> <tr> <td style="width: 15%;">$\gamma^{\text{♂}}$</td> <td style="width: 15%;">45%</td> <td style="width: 15%;">38%</td> <td style="width: 15%;">9%</td> <td style="width: 15%;">8%</td> </tr> <tr> <td style="width: 15%;">$\frac{d}{d} \frac{s}{s}$</td> <td style="width: 15%;">$\frac{D}{d} \frac{S}{s}$</td> <td style="width: 15%;">$\frac{d}{d} \frac{s}{s}$</td> <td style="width: 15%;">$\frac{D}{d} \frac{s}{s}$</td> <td style="width: 15%;">$\frac{d}{d} \frac{S}{s}$</td> </tr> <tr> <td style="width: 15%;">100%</td> <td style="width: 15%;">45% [D, S]</td> <td style="width: 15%;">38% [d, s]</td> <td style="width: 15%;">9% [D, s]</td> <td style="width: 15%;">8% [d, S]</td> </tr> </table>	$\gamma^{\text{♀}}$	$\frac{D}{d} \frac{S}{s}$	$\frac{d}{d} \frac{s}{s}$	$\frac{D}{d} \frac{s}{s}$	$\frac{d}{d} \frac{S}{s}$	$\gamma^{\text{♂}}$	45%	38%	9%	8%	$\frac{d}{d} \frac{s}{s}$	$\frac{D}{d} \frac{S}{s}$	$\frac{d}{d} \frac{s}{s}$	$\frac{D}{d} \frac{s}{s}$	$\frac{d}{d} \frac{S}{s}$	100%	45% [D, S]	38% [d, s]	9% [D, s]	8% [d, S]	0.25x2
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100%	45% [D, S]	38% [d, s]	9% [D, s]	8% [d, S]																			
4		<p>The map gene of two studied genes:</p> <p>The percentage of the recombinant phenotype is 17%, so the distance between two genes is 17cMg.....</p> <p>scale: 1cm → 2 cMg (Accept any appropriate scale)</p> <div style="text-align: center; margin-top: 20px;"> <p>17cMg</p> </div>	0.25																				

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