WJEC (Eduqas) Biology A-level Topic 2.5: Inheritance Questions by Topic - Mark Scheme

1.	Q	uesti	on	Marking details	Marks Available
	1	(a)	(i)	NnGg for both; NG Ng nG ng for both; correct completion of punnet square; correct ratio 9:3:3:1; correct phenotypes matched to ratio;	5
			(ii)	Correct expected number column 72:24:24:8;	1
		(b)	(i)	0.667/0.67/3/5;	1
			(ii)	7.82 circled;	1
			(iii)	Accept because χ^2 value is to left of {critical value/7.82}/ Accept because χ^2 value has probability higher than {0.05/5%}/ Accept because χ^2 value falls between {0.9/90%} and {0.8/80%} probability/ Accept because the probability lies between 80-90% that it is due to chance alone; If not circled any answer for (ii) must refer to 7.82 in (iii) ECF from chi squared table	1
		(c)		{Common phenotypes/red grey and scarlet ebony} are due to linkage/ description of linkage; {Rare phenotypes/red ebony and scarlet grey} due to {crossing over/recombinants};	2
				Question 1 Total	[11]

2.

Question			Marking details	Marks Available
2	(a)	(i)	Correct parental genotypes: BbRr BbRr; Correct gametes: BR Br bR br; Correct headings on punnet square: BR Br bR br; Correct completion of genotypes in punnet square;(ecf)	4
		(ii)	Chocolate ³ / ₁₆ / 18.75% and yellow ⁴ / ₁₆ / ½ / 25%; both for 1(ecf)	1
		(iii)	1⁄4 /25%;	1
		(iv)	All the noses should be brown; Accept fur is black and skin is brown	1
	(b)	(i)	BBrr;	1
		(ii)	Cross with {yellow, brown nose dog / homozygous recessive / bbrr} / back cross / test cross;	1
		(iii)	Yellow with a brown nose;	1
			Question 2 Total	[10]

3.

	0	41	Mayling dataile				Marks Available							
	Question		Marking details				A01	AO2	AO3	Total	Maths	Prac		
3	(a)	(i)	Purple White and Aa aa (1) Aa Aa aa (1) correct completion of punnett square				2		2					
		(ii)			is no significanumbers/rat E (1) 25 25 (1)		(O-E) ² 49 49	(O-E) ² E (1) 1.96 1.96	-	4		4	4	4
		(iii)	value (at 0.	ni ² value {is 05)} (1) nypothesis not due to c and E(1)	greater tha						4	4	4	4
	(b)	(i)		npletion (1)	Ab AaBb Aabb (from their ple : 5 white		a a s]	b aBb abb		3		3		
	(ii) Hypothesis 2 and The chi squared value is {to t value/no significant difference is between {0.8-0.9/above 0.4 alone				e is {to the le	eft/less t	han} the and E/p	robability			1	1	1	
	(c)		Hypothesis 1 white x white could only produce white offspring/hypothesis 2 white x white could give purple/or description with genotypes		or			1	1		1			
			Question 3	total			<u> </u>		0	9	6	15	9	9

4.

Question				Marks Available				
1.	(a)	(i)	Reject sex	chromosor	chromosomes; mes x chromosomes	not on the	autosomes	1
		(ii)	Parents		X ^H Y	*	X ^H X ^h :	4
		1.0	Gametes	XH	Y	XH	X ⁿ :	
			Offspring	XHXH	X ^H X ^h	X ^H Y	X ^h Y;	
				Normal female	Normal/ Carrier female	Normal male	Haemophiliac/ sufferer/ affected} male;	
		(iii)	NAME OF TAXABLE PARTY.		cf apart from phe r a haemoph <mark>i</mark> liac		3 A	1
		(iv)	0.25 / 25% Accept 1 in Reject 1:3					1
	(b)		AB AABB 3:1;	ab AaBb	AB AaBb	ab aab	12	3
			tetween a	and b for	ow some corre ecf oss is completed		tion of linkage	

Question	Marking details			
(c)	Incomplete linkage;	Max 2		
	Genes (further/ far) apart on same chromosome;			
	{Crossing over/ chiasmata} can occur;			
	Four types of gametes produced(but not in equal numbers);			
	Small numbers of recombinants / large numbers parental			
	types;			
	Recombinants equal in numbers / parental equal in numbers;			
	Question 4 Total	[12]		

Question		on	Marking details	Marks Available
5	(a)	(i)	CGC is replaced by TGC/C is replaced by T; Amino acid cys has replaced arg;	2
		(ii)	Change in {protein/tertiary} structure/different protein is made; MC1R will not be stimulated (by the hormone); {Less/no} eumelanin will be produced;	Max 2
			Question 6 Total	[4]

5.

6. **(a)** (j)

Gametes	HW	HR	hW	hR.
HW	HW	HR	hW	hR
	HW	HW	HW	HW
HR	HW	HR	hW	hR
	HR	HR	HR	HR
hW	HW	HR	hW	hR
	hW	hW	hW	hW
hR.	HW	HR	hW	hR
	hR.	hR	hR	hR

	1 mark per column; if wrong- gametes-1, letters -1;	4
(ii)	Phenotypes	
	Horn less roan 6	
	Horn less red 3	
	Horn less white 3	
	Horned roan 2	
	Horned red 1	,
	Horned white 1	
	3 marks all phenotypes + 3 for all correct matching ratio -1 per error, no ecf from (i)	6