

Assessment Schedule – 2014**Scholarship Biology (93101)****Evidence Statement****Question One Bulldogs: Evidence Statement**Human manipulation in evolution of bulldogs from wolves (**E**)

	Evidence		Justification
EW	wolves obtain scraps / humans adopted wolf pups	EW_J	Wolves followed / stayed close to humans
EB	Selectively bred / encouraged non aggressive wolves to humans	EB_J	Benefit explained eg keep other predators away, don't harm / attack humans, for hunting
EG	Small gene pool / founder effect (genetic drift)	EG_J	Unfavourable traits / alleles lost / favourable traits / alleles fixed in wolves OR favourable traits / alleles lost / unfavourable traits / alleles fixed / high frequency in bulldogs
	Bulldogs ferocious and insensitive to pain selected OR Thick-set, low in stature, broad and powerful selected OR Short broad face inclined upwards / wrinkles / powerful jaw selected	EP_{J1} EP_{J2}	Enables the bulldog to subdue animals / drive cattle / bullbaiting without being injured / killed Causes frequency of the alleles for these phenotypes to increase
ET	Selectively bred for docile temperament after 1835 / bullbaiting	ET_J	Enabled them to be suitable family pets that would not be a risk (to children) / for dog shows
EI	Inbreeding within bulldogs	EI_J	Increased chances of harmful recessive alleles coming together
		EL_J	Linked genes: a gene with disorder with a gene humans have selected for

how selective breeding (**B**) could be further used to eliminate **each** of the named disorders **and** evaluate how effective this might be

	Evidence		evaluation
BS	Brachycephalic Syndrome selectively breed for a long / longer face OR Dermatitis breed for no folds / wrinkles or fewer / less deep folds / wrinkles	BS_J	Will be effective but removes / reduces bulldog characteristic from standard (so impractical)
BR	Brachycephalic Syndrome breed only short faced dogs with limited respiratory problems		
BH	Hip dysplasia only breed those who show no symptoms of dysplasia	BH_J	Limited success with many genes causing disorder (so may be impossible to eliminate)
BV	VSD only breed individuals who do not have VSD	BV_J	Heterozygote parents will have 25% chance of VSD offspring / 75% chance of no VSD so will be very effective (as inheritance is due to a recessive allele)
BT	VSD carry out test cross / genetic testing to identify homozygous dominant individuals	BT_J	Will be most effective but takes time / is expensive
BC	Cryptorchidism do not breed males who have the disorder / females with male offspring with the disorder	BC_J	Since females are carriers this will be effective over time / generations
BO	Out breeding / cross breeding with other dog breeds	BO_J	Effective as long as bulldog standard can be maintained / Ineffective as will only maintain dominant bulldog standard traits
		BL_J	Disorder gene may be on same chromosome as bulldog standard gene / linked genes and so may be impossible to / can't eliminate without losing standard

Judgement statement (2 areas are E and B)

8	8J's OR 7J's and 2 descriptions Must have 3 J's from each E and B area Answer displays: <ul style="list-style-type: none"> • perception and insight • sophisticated integration and abstraction • independent reflection and extrapolation • convincing communication
7	7J's OR 6J's and 2 descriptions Must have 2 J's from each E and B area Answer displays aspects of: <ul style="list-style-type: none"> • perception and insight • sophisticated integration and abstraction • independent reflection and extrapolation • convincing communication
6	6J's OR 5J's and 2 descriptions OR 4J's and 4 descriptions Must have 2 J's from each E and B area Answer displays: <ul style="list-style-type: none"> • analysis and critical thinking; • integration, synthesis and application of highly developed knowledge, skills and understanding • logical development, precision and clarity of ideas
5	5J's OR 4J's and 2 descriptions OR 3 J's and 4 descriptions Must have 1 J from each E and B area Answer displays aspects of: <ul style="list-style-type: none"> • analysis and critical thinking • integration, synthesis and application of highly developed knowledge, skills and understanding • logical development, precision and clarity of ideas
4	4J's OR 3J's and 2 descriptions OR 2 J's and 4 descriptions
3	3J's OR 2J's and 2 descriptions OR 1 J and 4 descriptions
2	2J's OR 1J and 2 descriptions OR 4 descriptions
1	1J OR 2 descriptions
0	Lack of relevant evidence

Question Two: North Island Weta: Evidence Statement

Evolutionary processes and patterns resulting in Two groups of weta (T)

	Evidence		Justification
TE	Weta evolutionary rate is punctuated equilibrium	TE_J	Many / 70 species arose in the last 90 million years / since Gondwana split OR Many species of Giant and tree weta in last 2-5 million years
TA	Weta groups / species have undergone an adaptive radiation in NZ OR Giant weta and tree weta groups / species diverged from a common ancestor	TA_{J1} TA_{J2}	Different groups / species inhabit different habitats / niches available in NZ with two named examples such as urban, coastal, lowland forests, food sources allopatric speciation due to named geographical barrier OR Sympatric speciation due to niche differentiation
TS	Different selection pressures (e.g. types of plants for food) in each environment	TS_J	Caused changes in allele frequency between the populations leading to divergence
TR	RIM's develop between giant weta and tree weta groups / species as a result of changes to behaviours / courtship pheromones / mating sounds	TR_J	Different numbers of stridulatory ridges causes different sounds not recognised by the other group so no gene flow OR Different pheromones meant tree weta did not recognise giant weta so no gene flow OR Behaviour such as burrowing explained preventing gene flow OR Physical differences in mating organs would prevent gene flow (leading to speciation)
TF	Founder / bottleneck effect on offshore islands	TF_J	changes in gene pool / allele frequency (due to genetic drift)
TM	Mutations occurred in each (isolated) population		

Distribution difference (**D**)

	Evidence		Justification
DR	Large, widespread ancestral population of north island giant weta was reduced by mammalian predation / competition or habitat destruction		
DV	Giant weta have little amount of genetic variation in their population compared to tree weta OR tree weta have a large amount of genetic variation in their population compared to giant weta	DV_J	Prevents giant weta populations adaptation to a wide range of habitats unlike the tree weta
DA	Giant weta show no aggressive response to being threatened OR Tree weta show an aggressive response to being threatened: leg spikes up, jaws open, biting	DA_J	Giant weta less likely to survive attack by a predator OR tree weta more likely to survive an attack by a predator
DP	Giant weta are more likely to be eaten by (introduced mammalian) predators (such as rats, stoats, cats, hedgehogs) due to TWO of ; • Relatively slow moving • Cannot jump (to escape predation) / freeze • are larger (so more visible / appealing meal for predators) • are more exposed during the day / not in burrows • mate locked together for 4 hours / move sluggishly afterwards OR Tree weta are less likely to be eaten by predators due to TWO of: faster moving, can jump, are smaller, not exposed, mate quickly.	DP_J	leads to decreased survival / reproductive success of giant weta and so only found as three isolated populations OR leads to increased survival / reproductive success of tree weta and so have a widespread distribution
DE	Giant weta eggs are more likely to be predated upon as on forest floor (rather than in the ground) OR Tree weta eggs less likely to be predated upon as in the ground	DE_J	leads to decreased survival of giant weta and so only found as three isolated populations OR leads to increased survival of tree weta and so widespread
DH	Giant weta mating behaviour / no “harem” compared with tree weta mating behaviour / ‘harems’	DH_J	Strongest / dominant males are more likely to mate and pass on alleles beneficial for survival
DL	Giant weta don’t eat lichen so are competing with introduced pests for food OR Tree weta eat lichen so not competing with introduced pests for food	DL_J	Leads to decreased survival of giant weta so not widespread OR leads to increased survival of tree weta and so widespread

Judgement statement (2 areas are T, and D)

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Question Three DNA analysis in Evolution: Evidence Statement

	Evidence		Justification
ML	MYH16 allowed alarger brain	ML_J	allowed increased communication / abstract thought / make better tools / plan hunts linked to a consequence
MR	MYH16 allowed reduction in mandible / zygomatic arch / saggital crest	MR_J	Energy saved from the reduction can be used for brain development aiding cultural evolution
S	SRGAP2 Connections increased brain complexity / developed areas of the brain	S_J	Greater brain complexity allowed: more successful cooperative hunting or tool making / greater creativity (art, spirituality) / the start of tool use / teaching and learning CAN ONLY GET 1 OF S_J AND F_{J1}
F	FOXP2 Allowed speech / language	F_{J1} F_{J2}	Speech / language allowed greater communication related to 1 of: making or using tools / clothing / food gathering or hunting / building shelter From birth able to communicate facilitating learning (cultural evolution)
GB	Glucose transporters Greater energy production / supply to the brain through respiration	GB_J	larger / greater development of brain led to 1 of cultural evolution: make tools, use fire, clothing, building shelter, hunting, planning (communicating / art)
GM	Glucose Transporters led to smaller muscles	GM_J	Bipedalism becomes more efficient (compared to quadrapedalism)
HH	HACNS1 Greater manipulation of hand / opposable thumb	HH_J	precision grip / fine motor control contributed to being able to make more refined tool / shelters / clothing / art
HF	HACNS1 Foot arched / big toe larger / aligned	HF_J	Enables greater bipedal efficiency
A	Salivary Amylase Allows more starch digestion (in the mouth)	A_{J1} A_{J2}	Greater efficiency of starch digestion gave domestication of plants / wheat / rice / agriculture an adaptive advantage Greater supply of simple sugar (glucose) for growing / larger brain
SP	Any named mutation(s) are selected for by the environment eg living in an area with high starch foods selects for salivary amylase mutation	SP_J	Increases allele frequency of mutation / prevalence of mutation in gene pool
	Links 2 of MYH16 / SRGAP2 / FOXP2	L_{J1}	Explains how the combined effects of two of the mutations increase communication / cultural evolution
	Links glucose transporters to 1 of MYH16 / SRGAP2	L_{J2}	Explains how glucose is required for energy via respiration for the larger brain due to MYH16 / greater number of neurons due to SRGAP2
	Links salivary amylase to glucose transporters	L_{J3}	Explains how more efficient breakdown of starch by amylase means more glucose is available to the brain via glucose transporters

	Links MYH16 / SRGAP2 to salivary amylase	L₄	Explains how larger brain / greater neuron connectivity led to use of fire to cook starchy food which could be better utilised due to more amylase
	Links HACNS1 with MYH16 / SRGAP2	L₅	Explains how the greater manipulation of the hand / precision grip facilitates the making of eg tools enabled by the larger brain

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