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QUALIFY FOR THE FUTURE WORLD
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Scholarship 2022 Biology

Time allowed: Three hours
Total score: 24

ANSWER BOOKLET

Check that the National Student Number (NSN) on your admission slip is the same as the number at the top of this page.

Write your answers in this booklet.

Start your answer to each question on a new page. Carefully number each question.

Check that this booklet has pages 2–26 in the correct order. Pages 2–4 are blank and are to be used for planning. Pages 5–26 are lined pages for writing your answers.

Do not write in any cross-hatched area (☒). This area may be cut off when the booklet is marked.

YOU MUST HAND THIS BOOKLET TO THE SUPERVISOR AT THE END OF THE EXAMINATION.

Question	Score
ONE	
TWO	
THREE	
TOTAL	

ASSESSOR'S USE ONLY

PLANNING

Q2

Limited rearing niche behaviors (voluntary).

Breeding — territorial.

Solitary

choose sex choice ↓.

- Speciated, dependent. → changing environment.
as %

* Black footed ferret — prone

• co predation +/-

• extra parasitism +/- o obligate pathogen.
as host size ↓ host ↓, parasite ↓. o
parasite

Breeding:

protection.

metabolic energy demand

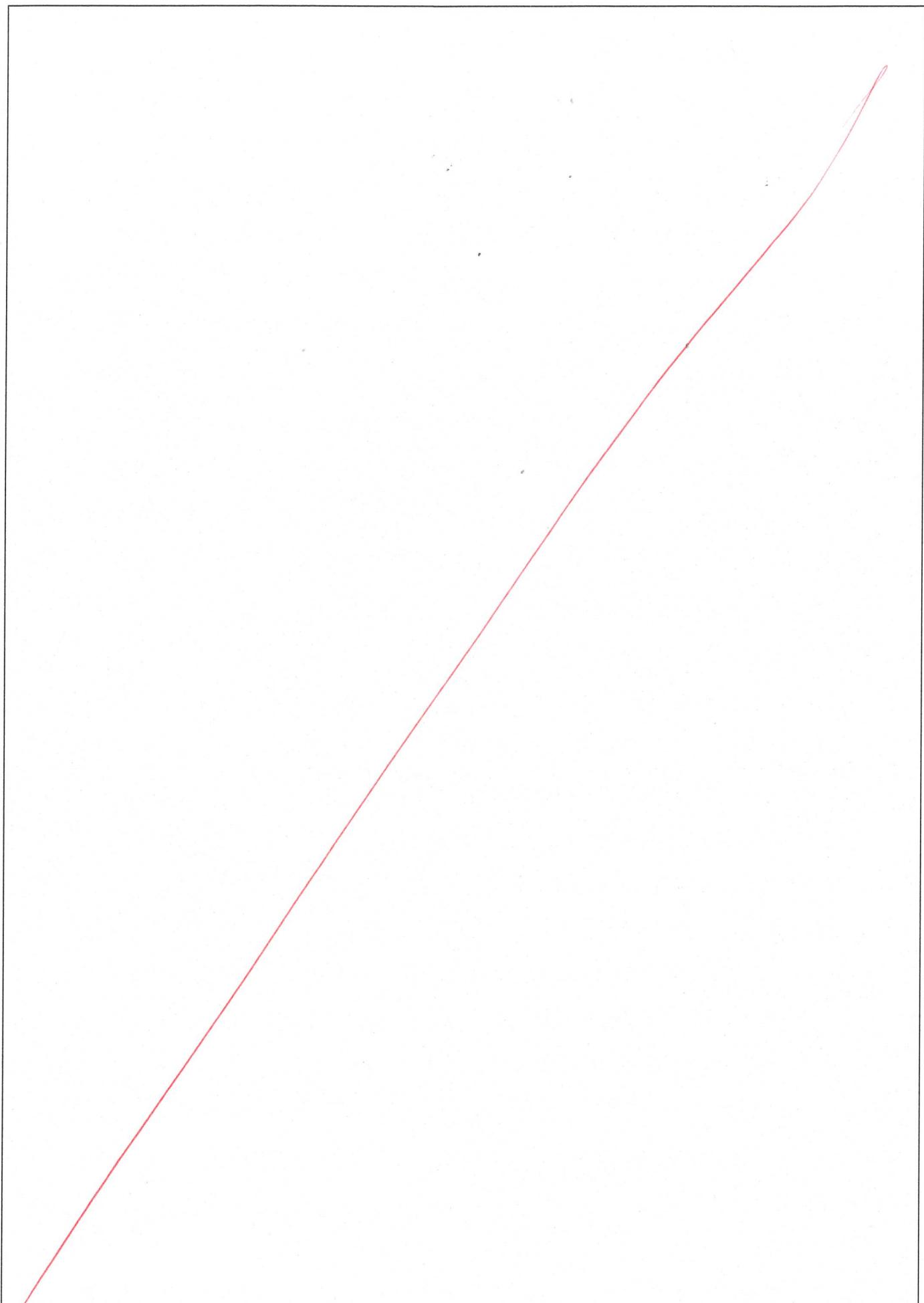
* to select the K selected. (small litter size
↳ litter)

* Bottleneck (as) — disease. — small gene pool — At fixed.
↳ genetic diversity small

Habitat loss., fragmentation

↳ ecological barriers, no corridor.

Prone host-pathogen — poisoning. Bottleneck epidemics.

PLANNING

PLANNING

(Q1)

Galapagos

→ Antarctic & water.

(1)

Divergent evolution: from Procellariiformes

[66 mya]

(Cretaceous mass extinction).

Adaptive radiation. (50 species).

↳ unoccupied niche.

(2)

Large

Competitor marine mammals.

Extinction. 25 mya.

(3)

Living

✓

Divergent directional selection.

↳ Selection pressure

↳ Divergent
colder
↳ smaller PSA↳ Verner
↳ larger SA

islands

① gene flow
② allopatric

no flight.

ACC

(4)

Great Auk

Convergent evolution — similar selection pressures.

— human intervention.

(5)

Common ancestor

— Gondwanaland.

— Areteal features.

- Oceanic carnivore niche cleared by reptiles.
- No need for flight — homologous structure wings.
- bones
- flatter.
- energy.

~~(*) Sympatric speciation may have occurred here via niche & specialization given that so many vacant niches were available.~~

5

Q1

The penguins underwent divergent evolution with the Procellariiformes around 66 mya. This date coincides with the mass extinction of apex predators, during the K-T / Cretaceous Tertiary / Cresta. eon - Paleogene extinction leading to a loss of 45% of species present. Divergent evolution is where a common ancestor gives rise to two or more different species, which is what is seen between penguins and the Procellariiformes. The mass extinction of apex predators such as sharks and marine reptiles opened new and vacant ecological niches which were unoccupied. This led to adaptive radiation of species to occupy and exploit the resources available ~~of penguins and Procellariiformes~~ ^{Scans}.

Mutations may have occurred in the last common ancestor to allow it to ~~occupy~~ habitats underwater. This would confer an adaptive advantage as it would have less interspecific intraspecific and interspecific competition, leading to a higher survival and reproductive fitness. Therefore, the allele frequency selecting for the oceanic carnivorous niche would increase, leading to the accumulation of reproductive isolating mechanisms such as ecological isolation. This would ultimately lead to speciation of penguins from the Procellariiformes clade.

Because of the shared ancestry with flying seabirds, the penguins share homologous features such as modified wings for underwater dives. This further supports the theory of divergent evolution as homologous features suggest the wings were adapted to a new function in the penguins. The loss of flight would have led to less energy spent in locomotion which could be allocated to reproduction and survival to increase its fitness. The feathers in penguins are highly specialised, to with the streamlined contour features allowing efficient diving and swimming when escaping from predators and in hunting. The insulating plumules allow exploitation of colder habitats as they provide dense insulating functionality. Penguins have dense bones, which would suggest ~~it is~~ robust and when fighting against ^{if is} for food intraspecies, and can ~~is capable to~~ suffer significant injury. Genes relating to thermoregulation, osmoregulation and diving capacity were undergoing //

Strong selection to increase in allele frequency as they conferred an adaptive advantage for the penguins to increase their survival and reproductive chances.

extinct and extant

The common ancestor of penguins were likely to have existed when Gondwanaland was a land mass connecting the continents of Africa, Antarctica, Australasia and New Zealand and South America. Because penguins are not capable of aerial flight, their current distribution globally would suggest ~~that~~ ^{support} be biogeographical evidence that Antarctica, Africa, NZ, Australia and South Africa were once connected. This would have made the dispersal across continents more likely to occur. Fossil evidence showed that extinct species had body mass up to 175 kg. whereas living penguins were much smaller ranging from 1kg to 30kg. This suggests that adaptive radiation occurred within the penguin species and ~~as~~ there was a large range of phenotypes present. The extinction of giant ~~penguins~~ penguins coincided with the ~~range~~ time range where footed whales and seals ^{otter} marine mammals increased in abundance. This is likely due to a competitive ~~relationship~~ interspecific relationship where the giant penguins and marine mammals were competing for resources such as food. Because of their similar ecological niches such as oceanic carnivore preying on larger fish, Gause's competitive exclusion principle applies. Eventually, the giant penguins were outcompeted by the marine mammals and driven to extinction. There may have ~~been~~ been a predatory relationship between penguins and larger marine ~~mammals~~ marine organisms such as orca and seals, although this was not as likely to be the reason of their extinction as the smaller-sized penguins survived.

allopatric

Living penguins underwent ~~sympatric~~ speciation via ~~niche differentiation~~. Where disruptive selection occurred to select for larger body types in colder regions and smaller body types in warmer regions. The larger body type reduced surface area to volume ratio to prevent excessive ~~exposure~~ of heat & convection loss of heat through convection and ~~as~~ confirmed //

a selective advantage as to survive in colder habitats. Similarly, small body types with higher surface area to volume ratio allowed loss of heat through convection cooling, and thus conferred adaptive advantage in warmer waters. These penguins were selected for via natural selection as there were different selection pressures in the different environments (cold and warm waters). The diversification of living penguins from common ancestor around 22 mya is an example of ~~at~~ divergent evolution, as multiple species were formed from one ancestor. The evolutionary history of penguins supports the punctuated equilibrium theory mode of evolution as the diversification of species occurred during short periods of time and ~~remained~~^{species} ~~but~~ remained stable for long periods of time ("stasis"). This is seen again in the living penguins where their diversification coincided with the separation of Antarctica from Australia and South America. The formation of Antarctic circumpolar current led to cooling of Antarctic regions and the current allowed dispersal of penguins to newly formed island groups such as the Galapagos Islands. The Galapagos Islands are tropical in nature and fall on the equator, ~~thus~~ because of the Antarctic circumpolar current which brings colder water around, the penguins were able to disperse and occupy the ~~empty~~ vacant niches of the Galapagos Islands.

It is this dispersal which led to the allopatric speciation of penguins. The geographical separation allowed different selection pressures to operate on populations occupying different niches. Over time, the differences accumulated to form reproductive isolating mechanisms (RIMs) such as ecological difference (penguins living in warmer waters unlikely to breed with those in colder waters), temporal difference (breeding seasons may differ), behavioural ~~isolation~~ isolations (courtship displays / rituals may differ) and mechanical isolation (sexual reproductive parts may not structurally fit together). Because the penguins are on different islands, these RIMs can accumulate and act as a barrier against gene flow, and thus speciation occurred. As penguins //

were flightless, the island groups prove are less likely to interbreed as they do not overlap in geographical sense. As natural selection acted on the isolated groups with different selection pressures due to different environments, this produced diversity amongst the ~~set~~ penguins.

The great auk penguins display convergent evolution with the Southern Hemisphere penguins as both were subjected to similar selection pressures being present in ~~Earth~~ similar environment of cold waters and were hunting similar prey of small fish. Despite being unrelated, the Great auk and the Southern Hemisphere penguins show some phenotypes such as ~~at~~ wing modified flippers to dive. This is an example of an analogous structure since the unrelated species evolved to possess similar morphological traits ~~as~~ because of their similar niche. Co-evolution with marine mammals such as orca and seals resulted in selection for more efficient escape from predators which led to their streamlined body shape being selected for. Similar selection pressure of predation applied amongst the Spheniscidae penguins which therefore parallel evolution to a streamlined body form was also seen in the Southern Hemisphere penguins. As the great auk were habitats in the Northern Hemisphere, they were likely to be separated geographically from ~~and~~ the Southern penguins (Laurocid was separated from Gondwana 180 mya). Therefore, the only explanation for their similar traits despite ~~to~~ their little relatedness is that the ~~two~~ derived traits were result of convergent evolution.

events forming
occurring to form the flightless seabirds

Overall, because of the multiple adaptive radiations due to biogeographical events led to a huge diverse range of species, both extinct and living.

Q2.

Black-footed ferret is a solitary species. This reduces chances of successful predatory behaviour associated with cooperative hunting, reducing its survival chances. In addition, black-footed ferrets are territorial, with males and females only coming together during breeding seasons. This reduces the chances of a social bond forming between mates and energy needs to be invested in each year to find a mate and establish a territory. This would have led to decreased reproductive success. The ferrets are a specialist species, which are completely dependent on prairie dogs for food. This means it has a narrow ecological niche, and is subject to higher risk of extinction when environmental conditions change. The ferrets are a ~~parasite of the predator~~ (and is positively impacted in this relationship) to the ~~prairie~~ prairie dogs where ~~ferrets hunt on prairie dogs for food~~ they are negatively impacted and energy and prairie dogs ~~die from hunting~~ However, because 90% of the ferret's diet is from the prairie dog, it is largely obligate predator of the prairie dog. This means that a decrease in the prairie dog population would lead to decrease in the ferret population, as it is unable to change prey to other mustelids. In addition, the ferrets are parasites on the prairie dog species where the ferrets benefit from prairie dog burrows. They gain protection, habitat, ~~so the ferrets are positively impacted~~, ^{from predators, use} from weather, and do not use them as dens, the prairie dog is species is negatively impacted as they are likely to be killed when they return to the burrow. However, similar to above, the ferrets are obligate parasites of the prairie dog species for their burrows. Because ferrets are in a competitive and a parasitic and predatory interspecific relationship with the prairie dog species, it is very likely that the prairie dog species can go extinct and lead to the extinction of the ferrets.

In addition, black-footed ferrets display a K-reproductive strategy where they have low reproductive rates, only giving birth to 3-4 babies each breeding season. The female invests a lot of energy into nursing each litter to its own nearby burrow which increases the risk of female death from predation. This decreases //

the reproductive survival of the female. ~~In fact~~ Additionally, the kits hunt independently late summer, with a short lifespan of only one year. This means ~~not~~ not all females survive to reproductive maturity and pass on their alleles to the gene pool. The slow reproductive capacity limits increase in fennec population density overall.

Furthermore, the fennec population underwent a population bottleneck in 1986, where a small population of around 130 was hit by a disease and shrunk to only 18 individuals. Because of the small population size, it is likely that genetic drift played a large role. Genetic drift is where alleles are fixed or lost due to chance. In a small population, the death/migration of ~~of~~ an individual ~~reduces~~ leads to a large proportion of the gene pool. Therefore, the genetic diversity of the small fennec population is small. Harmful alleles may have been fixed in the fennec population such as vulnerability to CDV and the sylvatic plague. The captive breeding programme involves only 7 who breed successfully out of 18. This impacts large extent of inbreeding, which increases the possibility of harmful ~~alleles~~ recessive alleles coming together. The increase in homozygosity of the population decreases genetic variation, thus ~~is~~ is not sufficient raw material to work on, natural selection could lead to species extinction. Although numbers are now higher at 220 captive & 200 wild, genetic diversity would be largely compromised. These factors have led to the critically endangered status of the fennec.

Moreover, the loss of habitats to agriculture and natural gas extraction degrades habitats of prairie dogs. As ~~never~~ detailed previously, because females are dependent on prairie dogs for food & habitats, the decline in prairie dog habitat would lead to possible extinction of females. ~~Because prairie dog~~ ~~is controlled through poisoning,~~ ~~if~~ The fragmentation of habitat would lead to ecological corridors destroyed, limiting the extent of gene flow which //

decreases the species overall fitness. Epidemics wiping out entire colonies are of concern as they can impose severe bottleneck effects.

There are ~~three main~~^{multiple} options for black-footed ferret management.

Firstly, captive breeding programs with artificial insemination can increase the reproductive success of ferrets. ~~More care is a~~ However, this would apply artificial selection pressure ~~to~~ and could potentially decrease selection pressure for ferrets with better sperm quality; ~~as humans are assisting the breeding.~~ This is seen in genetic studies too.

However, artificial insemination can benefit by ~~in~~ Contrary to genetic analysis to breed individuals with least relatedness to increase genetic diversity of the population.

By analyzing genome of the breeding stock, scientists can identify markers for increased immunity against sylvatic plague and CDV. Individuals with alleles coding for increased immunity can be ~~intensely~~ bred together to increase the allele frequency of beneficial traits.

Secondly, insecticide control of fleas in prairie dog colonies would decrease sylvatic plague through controlling the bacterial disease spread by fleas. This increases the prairie dog colony which would lead to more ~~more~~^{implications} numbers of prey available and habitats available for the foxes. However, ~~an infection of this is~~ the potential of ~~removing all the~~ increasing the prairie dog population density since we are removing ~~the most~~ ~~selection disease~~ a selection pressure on the prairie dogs. As prairie dogs are considered a pest, increasing their numbers could lead to flow-on effects on the ecological food chain that may not be desirable for other native species.

Thirdly, ~~the~~ vaccination of ferrets against plague and CDV may increase the population ~~plus~~^{Increasing ferret's survivability and reproductive success} immunity against the disease. However, a high percentage of the population would need to be vaccinated ($\geq 90\%$) to achieve herd immunity. This would mean considerable efforts into developing vaccines & delivering vaccines to the ferret population. In addition, vaccines against these diseases do not ensure immunity against other diseases. Therefore, outbreaks //

of other diseases may again lead to reduction of ferret population size.

Fourthly, the conservation of prairie and grassland habitats would increase

the ~~gene flow~~ ^{and} reproductive success of ferrets. The restoration

of habitats can allow dispersal across the range of areas.

This would lead to increased genetic diversity as different populations would

be subject to different selection pressure. The higher genetic diversity would

allow for a more ~~pop~~ ^{fit} resistance population against changing environments.

Therefore, in my opinion, ^{of} the key strategies outlined, the scientists should

focus on captive breeding and habitat conservation, as these two provide

most significant advantages, and can ensure a self-sustaining population

without significant negative implications.

Besides the strategies proposed, other options include cloning and transgenic techniques. Willa is a ferret which survived before the disease outbreak in 1981.

She therefore possesses ~~three~~ 3x more genetic diversity than the current

population. If through including her in the captive breeding programme, the

this would increase the ~~at~~ genetic diversity in the current gene pool.

The second cell line from SB2 died from CDV before breeding. It

is likely that ~~the~~ ^{his} DNA includes harmful alleles which increases

susceptibility from CDV. Then adding him to the gene pool would increase

the risk of ~~a~~ offspring dying before reaching sexual

maturity. Therefore, the SB2 cell line should not be included into the

captive breeding program. Cloning in turn can have biological implications

which include limited genetic diversity (as offspring is entirely similar).

However, since in this scenario, the clone will be added into a pre-existing

gene pool, this negative implication is largely negated. //

Transgenic techniques can also be used such as ~~t~~. For example, genes coding for increased immunity (such as alleles for MHC proteins) can be spliced into the breeding stock through techniques such as CRISPR. This would allow innate (inherent) immunity to increase ferret health & and survival, ~~etc~~. This is also inheritable if spliced into the gene producing cells, which needs less effort required to sustain the ~~pop~~ population as the ~~A gene with~~ allele will be inherited naturally in the wild.

Transgenics may bring in negative implications including the ~~etc~~ the hybridization of ferrets with other species leading to the immunity allele spreading into ~~with~~ other species. This can lead to flow on effects on ecosystem food chains and may lead to other pest species outcompeting the ferrets. Hybrid vigour may also be displayed with transgene hybrids, and this could lead to decline in ferret populations.

At In my opinion, cloning & transgenic techniques ~~are~~ definitely have large potential to improve the vulnerable status of blood footed ferrets, but they should be tested on smaller scale to minimize any negative implications that may follow. //

Planning Q3.

H. floresiensis / H. habilis left

Australopithecus gait similar.

14

H. erectus left and then evolved.

large reduction of body size

'Significant' dwarfing

• stegodon supports.

• Similar crania shape

*• Receding & small braincase.
(small brain).*

• Control fire.

Tool Fatty

Q3.

In my opinion, the larger bodied *H. erectus* left Africa, moved through Indonesia and Flores, then evolved into *H. floresiensis*. (second model).

One of the evidence to support the theory is that Flores has always been separated from mainland Asia even at the glacial periods. This means to reach Flores, a significant water crossing had to be made over at least 24 kilometers. This implies that the founding population must have had tools and technology such as rafts capable of making such a significant crossing. At 24 km further away, the island of Flores would have been visually unable to be seen, thus some farfetched and imagination must have been present amongst the founding population.

H. habilis is associated with brains of 614 cm^3 , with limited ability ~~for~~ of imagination and primitive Oldowan tool technology. Therefore, I find it unlikely that *H. habilis* would have been able to migrate out of Africa and make significant crossing into Flores. Although it is possible that a storm may have carried the ~~hominids~~ ~~homins~~ ^{need to} across over, this is less likely as a large population would have been established in order to maintain reproductive ~~about~~ capabilities in the population.

Add: Another piece of evidence to support this model is that Flores is an

island geographically isolated from other habitats of hominids. The reduced energy environment can lead to dwarfing of species when there is limited food source. Directional selection would have occurred to favour individuals requiring less energy to sustain ~~and~~ survive; and this could lead to increase in allele frequencies of alleles ~~decreasing~~ ^{such as presence of} a smaller body size. Dwarfing is an occurrence supported by other evidence where Stegodon, and other dwarfed species were present on Flores and other small ^{islands} evidence. On one hand, this supports the dwarfing hypothesis, and on the other hand, this provides evidence that *H. floresiensis* had ^① food source available to them. Additionally, dwarfing may have been exacerbatd by the small population size. Significant amounts of inbreeding would have occurred, which may lead to homozygous recessive alleles expressed in homozygous individuals. One of these alleles could have been coding for reduced body size, thus the ~~isolated~~ ^{and its allele frequency} high frequencies of inbreeding would have increased its expression. Over time, genetic drift ^{may} ~~would have~~ led to this allele being fixed in the Flores population by chance. ~~The founding population~~ However, this can also be seen as evidence for the first model where a ~~homogeneous~~ ^{species with relatively} small body size migrated to Flores and further inbreeding led to an increase in the allele frequency for the small body size. Having said this, I believe dwarfing hypothesis is supported by the evidence of similarity of dwarfing amongst other animals and the fact that ~~the~~ Flores with its limited food source would lead to selection for smaller body size.

The wealth of evidence in physical morphology and fossil evidence suggest a mosaic of features between ancient and more primitive and ~~more~~ relatively more modern features. (cont next page).

④ Similar dwarfing is seen in *H. florensis* population in *H. naledi* population where the isolated geography and excessive inbreeding along with limited food led to largely reduced body size.

H. floresiensis has a cranial shape similar to *H. erectus* with a receding and small forehead with a flat face. This is evidence to support my chosen model as comparative anatomy can provide evidence of ancestry. Given that their similarities in cranial shape, it may be likely that *H. floresiensis* descended from *H. erectus* rather than *H. habilis*. The *H. floresiensis* possessed a relatively large jaw and teeth also resembling *H. erectus*. However, its bipedal gait is reminiscent of older hominins such as *H. habilis* and australopithecines. This could be seen as evidence for ancestry from a more primitive hominin such as *H. habilis*. However, I think a hominin with such an inefficient gait (flat arch lacks spring mechanism for propulsion, shorter thigh bone) would have been unlikely to migrate a long distance from Africa into Flores. Therefore, I propose that there may have been a reverse evolutionary process where once ~~H. erectus~~ *H. erectus* reached Flores, the lack of selection pressure for bipedal gait (Flores is a tropical island, ^{less} ~~no~~ need to cover long distances) led to a the transition into a more primitive gait. Similarly, the wrist bones could have undergone similar processes. The fact that *H. floresiensis* lack features that evolved with ancestors of modern humans at least 800 kya implies that *H. floresiensis* diverged from the modern human lineage at least 800 kya. However, this can be taken as evidence for both models as modern human lineage only developed ~~at~~ 200 kya. Given that *H. erectus* is seen as consequently believed to be ancestor of modern humans, it means that *H. floresiensis* must have diverged from *H. erectus* 800 kya. This ~~is coincidently~~ coincides with the colonization of Flores and therefore can support the evidence that ~~H. erectus~~ *H. floresiensis* evolved from *H. erectus*.

Other evidence to support the chosen model include cultural evidence.

Stone tools were found during 190 kya to 150 kya including tools anterior but smaller than those carved by *H. erectus*. This is reasonable as *H. floresiensis* had a smaller hand size, and would have made smaller tools //

relatively complex
↑↑↑

to fit their grasp. The use of choppers, radial cores, perforators with cutting edges indicate Acheulean tool technology. This paper does not support that *H. floresiensis* evolved from a more primitive hominin which used Oldowan tool technologies. *H. floresiensis* is proposed to have existed up to 790kya (50kya due to their tools dating to this age. This means they have coexisted with multiple Homo species including *H. sapiens*, or *H. neanderthalensis* etc. Because of its geographical isolation, *H. floresiensis* was able to survive until replacement by *H. sapiens* occurred as proposed by the OoA model. The use of fire in Liang Bua cave provides evidence of controlled fire use in *H. floresiensis*. Because *H. erectus* has been the species believed to first control fire, this again supports my chosen model. *H. floresiensis* was also believed to be hunting small elephants. This would suggest social co-operative behaviour, which requires communication, foraging and planning. The more complex developed *H. erectus* is therefore ~~more~~ likely to be the ancestor of *H. floresiensis*.

~~Fire was also believed to be used for cooking.~~ If the alternative model is correct, this implies that *H. habilis* would have been able to control fire, had a developed tool technology and complex social behaviours. However, I do not believe this to be correct, and thus I believe *H. erectus* is the ancestral population of species of *H. floresiensis*. Compiling all evidence of biological and cultural evolution, I propose that the larger bodied *H. erectus* left Africa during the initial migration. So a subset of the ~~initial~~ ^{initial} population were established on Flores. ~~The island habitats~~, founder effect and genetic drift, and extensive inbreeding led to dwarfism. And finally, the geographical isolation on islands meant *H. floresiensis* could coexist until it was replaced with *H. sapiens* geographically. //