



Oxford Cambridge and RSA

Friday 16 October 2020 – Morning

A Level Biology B (Advancing Biology) H422/02

Scientific literacy in biology

Advance Notice

Time allowed: 2 hours 15 minutes



INSTRUCTIONS

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INFORMATION

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The mitochondrial genome has a higher mutation rate (about 100-fold higher) than the nuclear genome. What explains the high mutation rate of mtDNA? Two nuclear genes, TWNK and POLG, encode enzymes for replicating the mitochondrial genome. TWNK encodes a helicase enzyme and POLG encodes DNA polymerase gamma. The POLG protein consists of two regions: a catalytic region that exhibits polymerase activity, and an exonuclease region that is involved in the recognition and removal of DNA base-pair mismatches that occur during DNA replication. A recent study suggests that mitochondria may have a nucleotide imbalance that leads to decreased POLG fidelity and higher mtDNA mutation rates.

What is mitochondrial disease?

When a person has mitochondrial disease, the mitochondria in the cells fail to produce enough energy. They are either inefficient or they do not work at all. There is huge variety in the symptoms and severity of mitochondrial disease. It depends on how many cells are affected and where they are in the body. Each person with mitochondrial disease will have a different combination of functional and non-functional mitochondria within each cell. However, there are times when particular body systems are affected in a recognisable pattern and these diseases have specific names. One example is Alper's disease.

Alper's disease

Alper's disease is a mitochondrial disease that affects the brain and liver. Symptoms of the disease include severe epilepsy, loss of developmental skills and liver failure.

Alper's disease is caused by mutations in the nuclear gene called POLG. The mutations are present in both the catalytic and exonuclease regions of POLG. The faulty product of POLG – the polymerase gamma enzyme – fails to produce sufficient amounts of functioning mtDNA in the liver and brain.

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