Urszula Golebiewska, Associate Professor

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Dear Annotation Control Team,

We are submitting the Queensborough Community College annotations of the genome of Smeadley. Smeadley has high homology to other A8 subcluster members, there are no unique genes. We identified GP 19 and 20 as part of programmed frameshift.

There is one tRNA (tryptophan), gp5 that we confirmed with web-based Aragorn, and tRNA-scanSE. We recommended deleting gp59 not enough support and it is a forward gene in a long stretch of reverses genes.

Sincerely,

Urszula Golebiewska