Urszula Golebiewska, Associate Professor

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Dear Annotation Control Team,

We are submitting the Queensborough Community College annotations of the genome of Loser. Loser has high homology to other A2 subcluster members, there are no unique genes. We identified GP 27 and 28 as part of programmed frameshift.

There is one tRNA (tryptophan), gp10 that we confirmed with web-based Aragorn, and tRNA-scanSE.

GPs 37, 38, 54, 56, 62, 63, 66, 69, 83 did not have 1:1 BLAST matches and were challenging in assigning the starts.

Sincerely,

Urszula Golebiewska