

Notes

These results are provided by the Molecular Genetics Lab at PBS, Nanaimo (DFO)

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Samples were collected, dissected, homogenized by Hakai staff

Nucleic Acid Extraction and Pathogen analysis by:

Tobi Ming

Data processing by:

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OTHER:

Export parameters:

Program Hakai
Current TRUE

Export Description:

Here are the Worksheets in this file...

1. Results CT – shows the cycle number from the qPCR.

a. NO LOD limits have been applied

b. 999 – negative result

c. -999 – duplicated results did not match. This is most often due to the high CT (low load) samples on the edge of the detection limits

2. Results Copy - shows the copy number as determined by the Biomark software using the MGL APC standards.

a. NO LOD limits have been applied

b. 0 – negative result

c. -999 – duplicate results did not match. This is most often due to the high CT (low load) samples on the edge of the detection limits \

A BLANK Result indicates that the Assay was not run for that sample, or that the reaction failed

3. Assay Information

- a. Table of abbreviations used for tested agents.
- b. LOD is the Limit of Detection for that specific assay (95% detection level)

Please let me know if you have any questions.