

Dear Fishpathogens.eu user

This is the third newsletter from Fishpathogens.eu. Find previous newsletters at <http://www.fishpathogens.eu/newsletters.php>.

NEW IHNV DATABASE RELEASED

The IHNV database is now public available (www.fishpathogens.eu/ihnv). A publication on this database is available at Journal of Fish Diseases

(<http://www3.interscience.wiley.com/journal/123334558/abstract>). The database so far contains data on 64 isolates on IHNV. 65 sequence reports are currently available. All users of the VHSV database are automatically users of this new database as well. You can search all the public available data by following this link:

<http://www.fishpathogens.eu/ihnv/search.php>

VHSV DATABASE

The VHSV database now covers 434 isolates and 327 sequence reports. Included in this are 179 full length G-gene reports covering all genotypes and subtypes. For most isolates the information available for each isolate is far more abundant than the information found in genbank. You can search all the public available data by following this link:

<http://www.fishpathogens.eu/vhsv/search.php>

FULL-LENGTH SEQUENCES

Since several users wish to have the stop-codon included when retrieving full-length open reading frames we have now included this in the search result. Sequences added without stop-codon will still be included as full-length sequences also.

DO YOU WANT TO START ADDING DATA?

You are always very welcome to contact me on spjo@vet.dtu.dk if you would like to get some help to add your data. We are grateful for all contributions. You are also welcome to send feedback on your experiences with the database to this address. There are also manuals available:

<http://www.fishpathogens.eu/vhsv/help.php>

Thank you for your time
Søren Peter Jonstrup