

IHN in Europe

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in comparison to VHSV only few current data available:

- literature
- organisations (OIE, FAO)
- databases

➤ Distribution of IHN in Europe

➤ phylogenetic analyses



History

Originally = viral disease of Pacific salmonids (spawn and fingerlings) in river systems at the Pacific coast of North-America

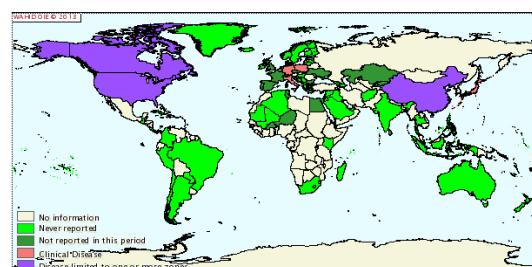
- 1953** Oregon-Sockeye-virus from sockeye salmon (*Oncorhynchus nerka*) hatcheries in **Washington** (Rucker et al. 1953)
- 1960** Sacramento-River-Chinook-Virus from chinook salmon in **California** (*Oncorhynchus tshawytscha*) (Ross et al. 1960)
- 1969** Infectious Hematopoietic Necrosis Virus (IHNV) from rainbow trout (*Oncorhynchus mykiss*) von Amend et al. (1969)
- 1971** IHNV from sockeye salmon (*Oncorhynchus nerka*) in Mori hatchery in Hokkaido **Japan** (eggs from Alaska)
- 1987** **France** (Baudin Laurencin, 1987) and **Italy** (Bovo et al., 1987)
- 1992** **Germany** (Enzmann et al., 1992)

VAHID Interface

OIE Home Page

IHNV distribution maps

01-06/2012



IHNV outbreaks in Europe (Ref.: OIE)

Year	Σ	AT	BE	CH	CZ	DE	ES	FR	HR	IT	NL	PL	RU	SI
1996	11				1			7		3				
1997	45		8	7		11		3		5				11
1998	35	1	1	1		6		1		12				13
1999	35	1		2		9		18		3				3
2000	44				1		6		23	6		1	7	
2001	30	1			1	11		8		7				2
2002	37	1		1	3	13		6		2		1	3	
2003	28	2		1	2	11		4		7			1	
2004	24			1		7		7		7				2
2005	?					5	1	x	x	x				
2006	14				1	12								1
2007	6				1	4								1
2008	28					6		3		18	1			
2009	5					5								
2010	13	1			1	5				5				1
2011	18				1	9		2		2	2	1		1
2012	5					2				2		1		

IHNV distribution: 2005-2012

Austria
Canada
China (People's Rep. Of)
Croatia
Czech Republic
France
Germany
Iran
Italy
Japan
Korea (Rep.of)
Netherlands
Poland
Russia
Slovenia
Switzerland
United States of America

Classification of IHNV**Demonstrated clinical disease**

Country	Latest report date	Domestic	Wild	Note
Germany	Jul - Dec, 2012	✓	✗	
Italy	Jul - Dec, 2012	✓	✗	
Japan	Jul - Dec, 2012	✓	✗	
Poland	Jul - Dec, 2012	✓	✗	
Switzerland	Jul - Dec, 2012	✓	✗	

Disease restricted to certain zone(s) / region(s) of the country

Country	Latest report date	Domestic	Wild	Note
Canada	Jul - Dec, 2012	✓	✓	
China (People's Rep. of)	Jul - Dec, 2012	✓	✗	
United States of America	Jan - Jun, 2012	✓	✓	

- no clear differentiation with classical diagnostic methods

- **close serological relationship** → 1 serogroup

- differentiation only with mAb

- original classification is based on the **geographical origin** of the isolates:

RB (Round Butte; Salmon, Alaska and British Columbia),

WRAC (Western Regional Aquaculture Center, freshwater fish)

SRCV (Sacramento River Chinook Virus, salmon , California)

1980s Phenotypic and genetic diversity among IHNV isolates from different geographical sources
 * protein electrophoresis (Hsu et al. 1986)
 * monoclonal antibody reactivity (La Patra et al 1995, Ristow & Arnzen 1989, Winton et al. 1988)
 * RNase T1 fingerprint analysis (Oshima et al. 1995)
 * ribonuclease protection assay (RPA) N, G, NV

→ Separation into four groups = correlate with the different geographic origin

1987 Identification of glycoprotein gene sequence (Koerner et al. 1987)

→ Increase of data for phylogenetic and comparative investigations
 * highly conservation of G and NV
 * nucleotide divergence of only 3.6 % (G) - 4.4 % (NV)

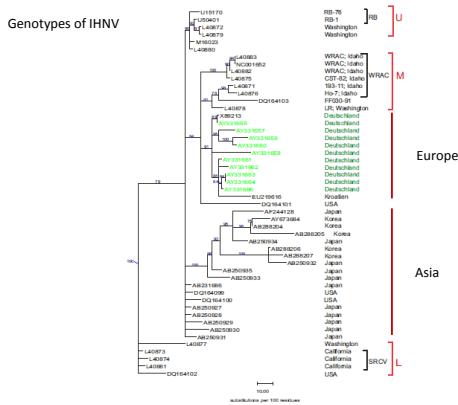
→ Correlation of phylogenetic relationship with the geographic origin of virus isolates rather than with host species or time of isolation (Kurath et al. 2003)

Antiviral Res. 2012, 95(3):257-81

Review

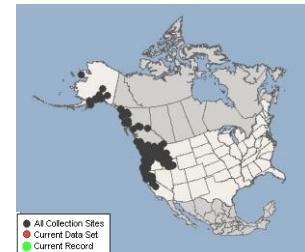
Countermeasures against viral diseases of farmed fish

Frederick S.B. Kibenge^{a,*}, Marcos G. Godoy^b, Mark Fast^c, Samuel Workenhe^c, Molly J.T. Kibenge^c

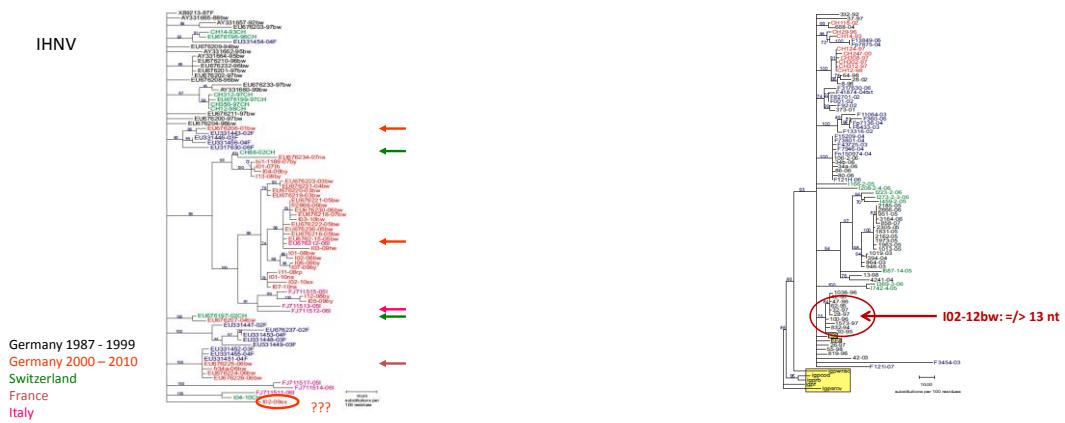
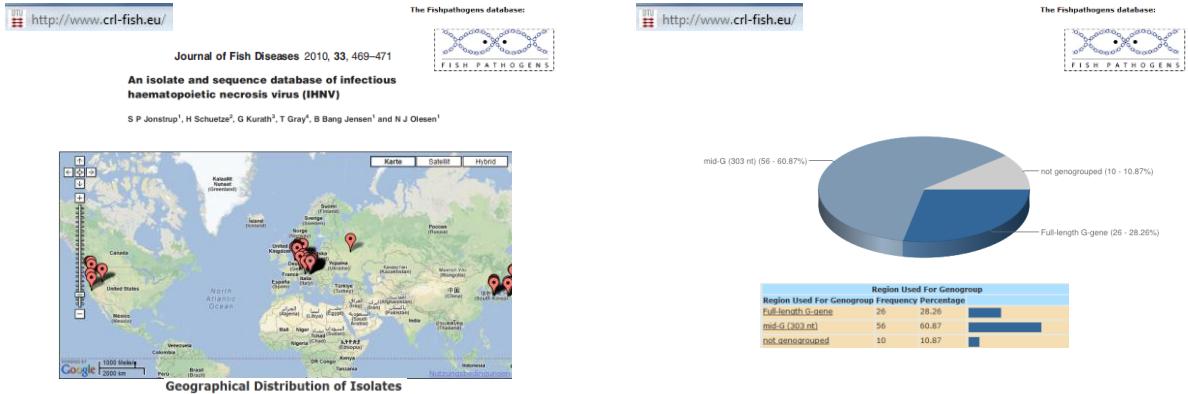


Journal of Fish Diseases 2011, 34, 579–587
Development of an aquatic pathogen database (AquaPathogen X) and its utilization in tracking emerging fish virus pathogens in North America

E.J. Emmerenger¹, E. Kentop², T.M. Thompson¹, S. Pittman³, A. Ryan³, D. Keon³, J.A. Carillo⁴, J. Ranson⁵, R.B. Life¹, R.M. Troyer³, K.A. Garver³ and G. Kurath¹



<http://gis.nace.org/ihnv/#>



Conclusions:

- European IHNV isolates: Genogroup M
- Only few or no sequence information about current IHNV isolates from :
Austria, Croatia, Czech Republic, France, Italy, Netherlands,
Poland, Russia, Slovenia, Switzerland
 - Submission of IHNV isolates to EURL Fish for characterisation
 - Better understanding about distribution and evolution of IHNV isolates
 - Better risk analysis and assessment