

5 degrees of separation -

does a reduced propagation temperature for SAV contribute towards higher titers and increased challenge model mortality?

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Topics for the talk



Acquired SAV isolates from across Norway – SAV₂ and SAV₃
Sequencing - MinION vs Illumina approach

Isolates subjected to *in vitro* studies – optimal temperature

Cohabitation challenge model – mortality seen affected by propagation temperature?

Handling of heart tissues from confirmed PD outbreaks

Tissue homogenates were inoculated on CHH-1 cells

→ cultures with high SAV transcript levels were chosen

Passaged until visible CPE → 2-7 passages

Selected some isolates for sequencing

	Heart tissue	Cell culture	Passage #
SAV ₃	VXo1	VXo7	7
	VXo2	VXo8	3
	VXo3	VXo9	5
	-	VX13	4
SAV ₂	VXo4	VX10	3
	VXo5	VX11	2
	VXo6	VX12	3



Illumina Sequence Capture vs. MinION Genome Sequencing

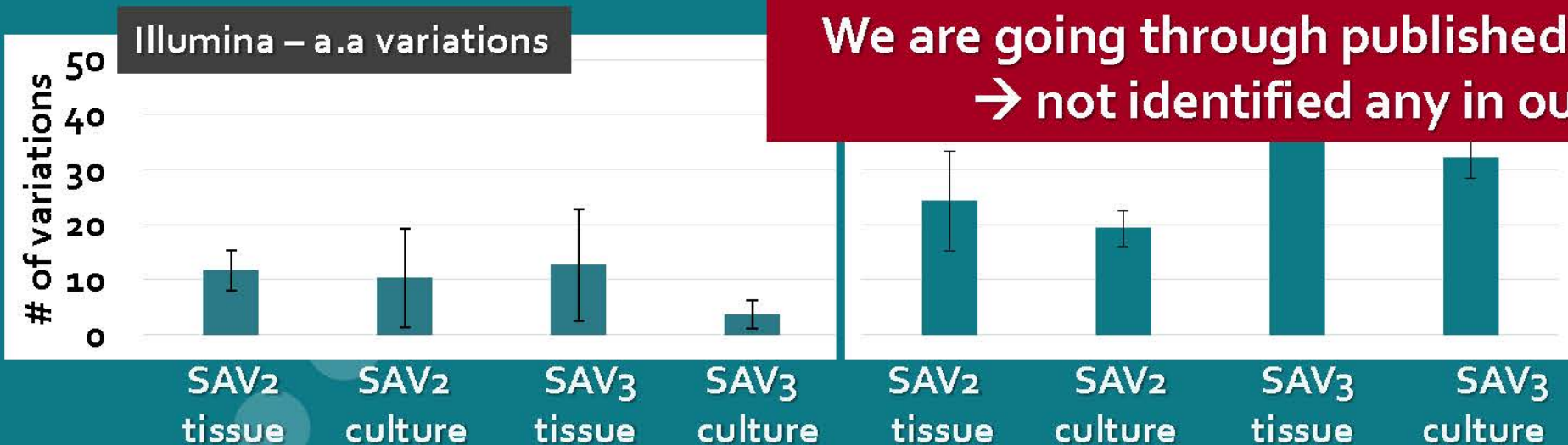
The consensus sequences for Illumina and MinION were identical

Phylogenetically the parallel tissue and cell culture samples grouped together

- The subtypes fall into distinct clusters
- In general very similar sequences with few genetic variations

MinION seem to overestimate single nucleotide variations

- Illumina showed limited SNV's in both tissues and cultures
- SAV2 indicates less variation than SAV3
- Bottleneck effect; reduced variation in cultured samples than tissues



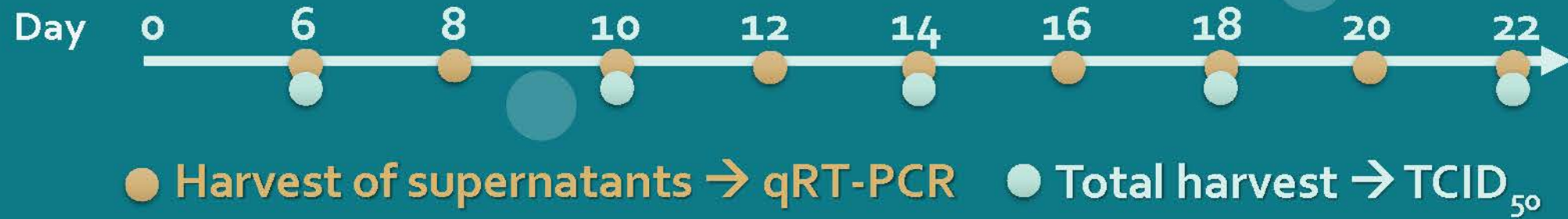
We are going through published amino acid mutations
→ not identified any in our sequences yet

In vitro characterization of SAV2 and 3 isolates

SAV₂: VX010

SAV₃: VX07, VX08 and VX09

Optimal culture conditions verified by SAV transcripts & TCID₅₀



- Incubation at 10, 15 and 18°C
- Three independent experiments
 - present data from individual experiments

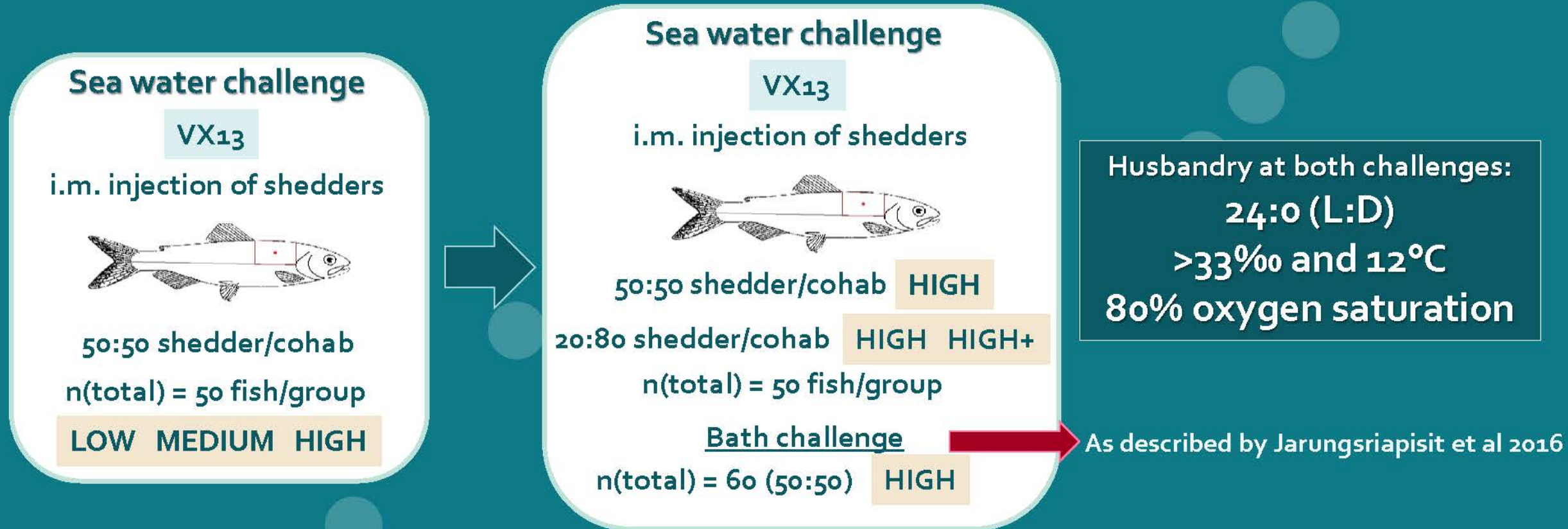
In vitro characterization of SAV isolates

Incubation at 10°C gave higher SAV transcript levels than 15 and 18°C

Infectivity were in general negatively affected over time at 15°C

Cultivation at 10°C gave highest titer for both SAV₂ and SAV₃

SAV₃ challenge model development



VX₁₃ has given mortality in a sea water challenge model (Taksdal et al 2013)

→ could we repeat that?

Moribund fish sampled for qRT-PCR of heart tissue

Challenge terminated after 6-8 weeks with sampling for histology and qRT-PCR (n=10/group)

Sea water challenge model development

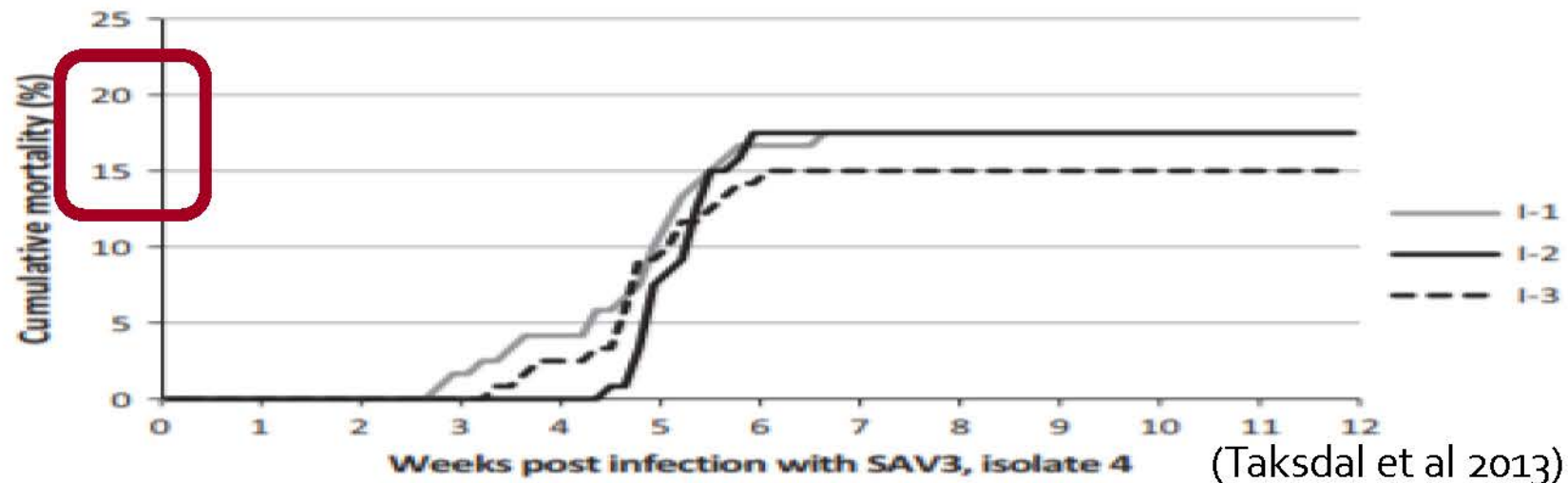


Table 3 Cumulative mortality in Atlantic salmon carrier fish injected with three isolates of salmonid alphavirus (SAV) 2 and three isolates of SAV 3 and in sham-injected fish in control tanks

Viral isolate number	SAV 2			SAV 3			Control
	1	2	3	4	5	6	
Number of dead fish	0	0	0	1	1	1	1
Percentage mortality	0	0	0	0.3	0.3	0.3	0.3

Differences:

Dose? Our low dose is lower

Propagation? CHSE vs CHH-1

Temperature? 15 vs 10°C

Shedder ratio? 50:50 vs 20:80

Lower cumulative mortality for the second challenge – any indications in the PD infection status?

Conclusions

Norwegian SAV₂ and SAV₃ isolates

- SAV₃ yields higher TCID₅₀ than SAV₂
- SAV₃ show slightly higher genetic variation than SAV₂
 - both in tissues and after propagation in cell culture

For both SAV₂ and SAV₃, propagation at 10°C was superior to 15 and 18°C

- Higher SAV transcript levels and titers at 10°C
- Reduction of infectious particles with increase in temperature

SAV₃ challenge models induced mortalities between 35-60%

- Similar mortality rates in both cohabitation and bath challenge models
- Has been shown previously for the same isolate (Taksdal et al. 2013)
 - May cultivation temperature play a role?

The literature show that temperature is important – the degree of separation is $\pm 5^\circ$?

Moriette et al (2006) – sleeping disease virus

- *In vivo* challenge after propagation at 10 and 14°C of a WT and recombinant SD virus
- Increased mortality for the strains propagated at 14°C for both WT and rSD



Taksdal propagate at 15°C – we at 10°C → variation in cumulative mortality

- Not only propagation temperature that contribute;
 - fish
 - challenge doses and challenge model
 - onset of challenge after smoltification

We have an abundance of sequencing data to dig through...

- Compare our isolates to previously known mutations that may be relevant for infectivity
 - Sequence samples from 10 and 15°C cultivated samples - genetic differences?

Thank you for your attention!

• norway
vaxx:inova