How do OsHV-1 µvars stack up? A comparison of multiple variants through experimental challenges

Colleen Burge

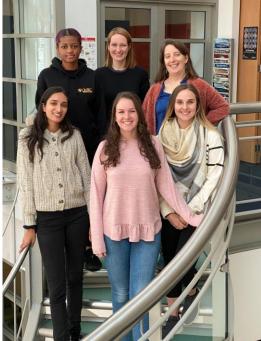
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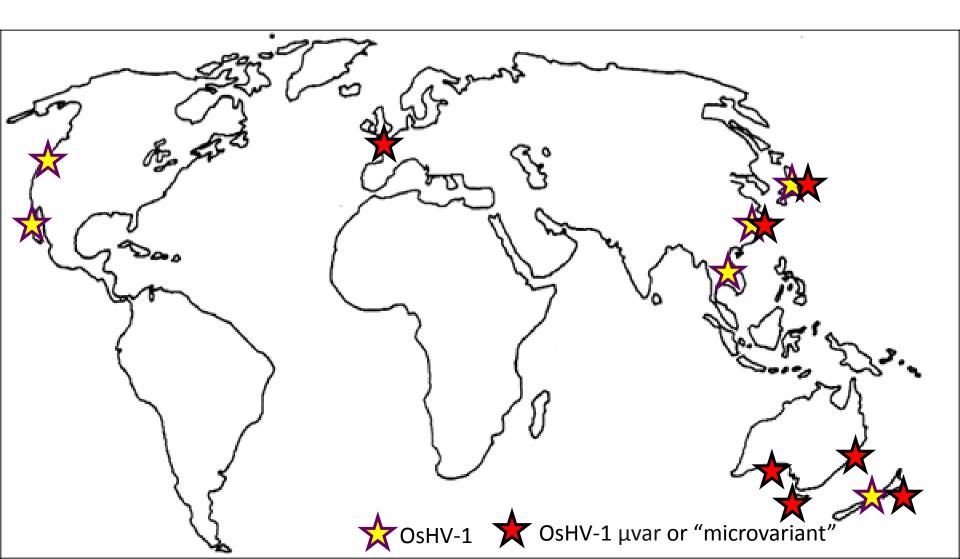
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What is OsHV-1 and why do we care?

- The Ostreid herpesvirus 1 (OsHV-1) causes mass mortalities of Pacific oysters
- Host range not restricted to Pacific oysters
- Disease progresses rapidly and can kill up to 100%
- Once established OsHV-1 persists in the environment (likely in surviving oysters), causing seasonal mass mortalities
- Virus can be transmitted within species and life stages; both horizontally and vertically
- Multiple variants of OsHV-1 exist including the emerging OsHV-1 µvars; a cause of concern for oyster growers globally

Reviewed in Pernet et al 2016, Arzul et al 2017, Burge et al 2018







Research Objectives

- Conduct laboratory trials to examine differential survival of oysters to OsHV-1 μvars and OsHV-TB using multiple stocks and species
- Initial characterization of a new OsHV-1 from San Diego, California



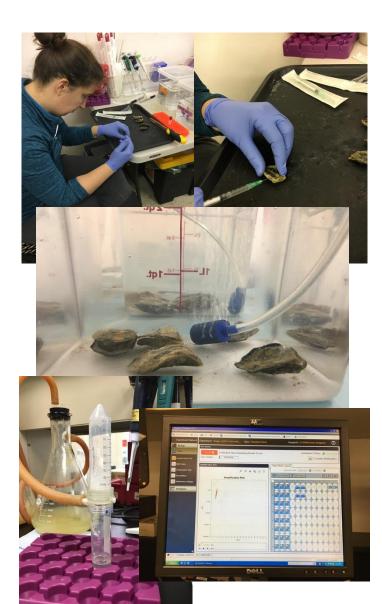


Laboratory Trials in Tucson, AZ

Experiments were conducted in Arizona over 200 miles from the ocean and in a hot, dry desert in strict quarantine

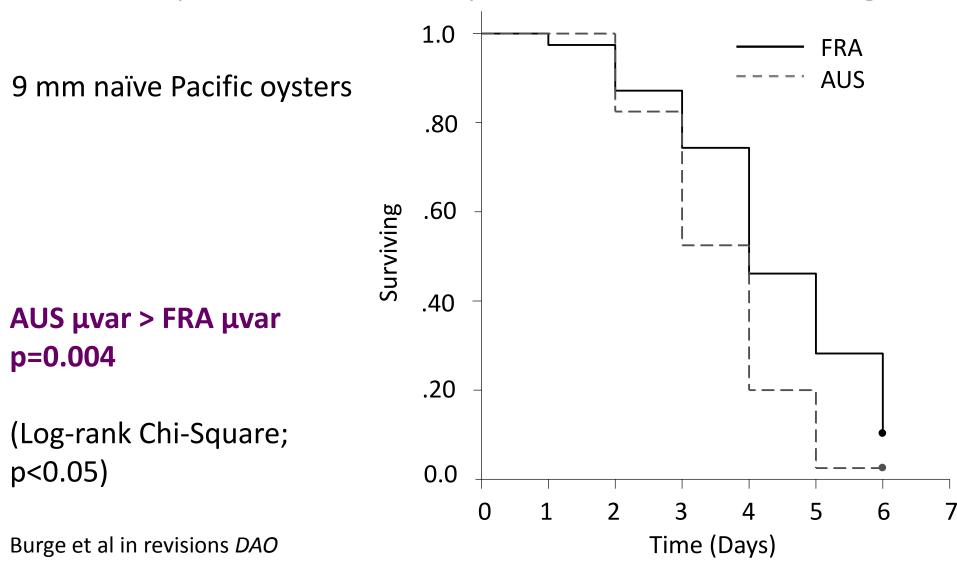
Laboratory methods

- Received infectious virus stocks from France & Australia
- Used susceptible animals to create larger stocks of homogenates
- Sampled moribund animals to create filtered tissue homogenates
- Three challenge types
 - Bath
 - Injection
 - Natural water exposure



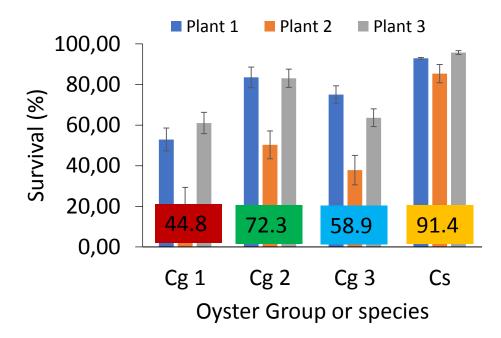


First comparison of two "µvars" in bath challenge





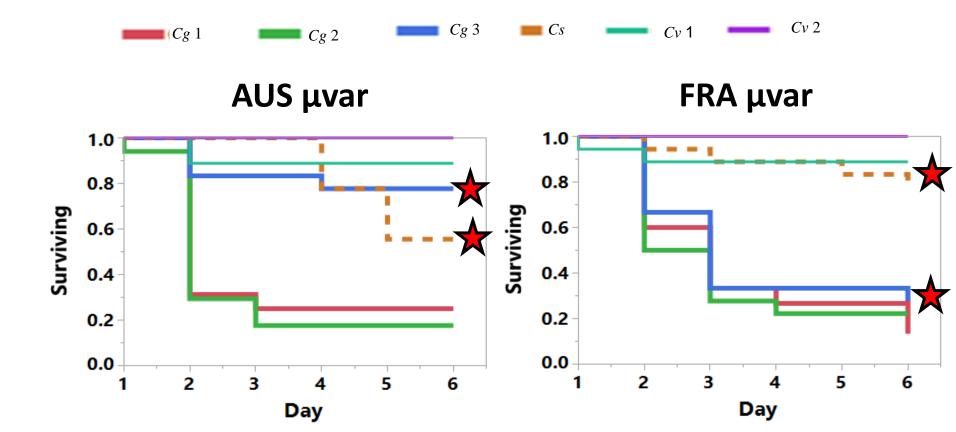
Injection trial (from field to lab)



Field data: Burge *et al.* in prep

- *Cg*1=low survival (52.9 ± 0.7 mm)
- Cg2=high (53.2 ± 0.8)
- Cg3= moderate (44.8 ± 0.6)
- *Cs*= high (41.5 ± 0.5)
- *Cv*1=high (40.8 ± 0.50
- *Cv*2=high (39.1 ± 0.6)

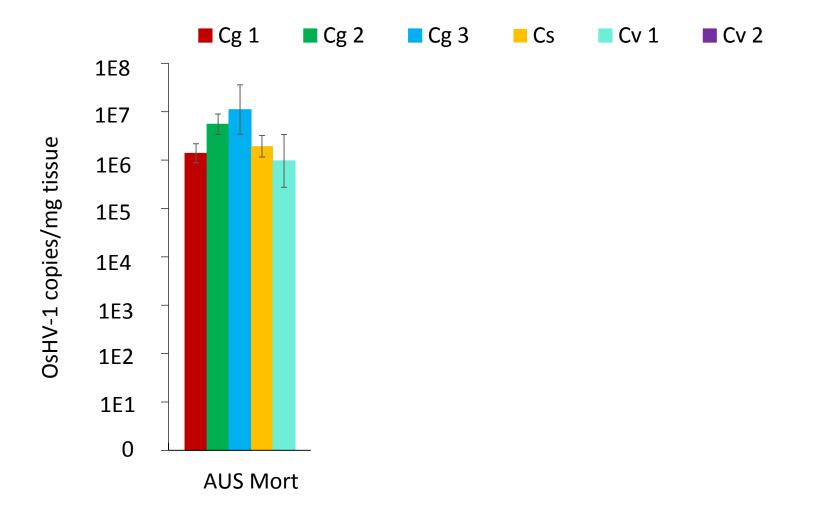
Cg= Crassostrea gigas; Tazzie, Midori & Wild X MBP Cs=Crassostrea sikamea Cv=Crassostrea virginica; DEBY, LOLA Injection challenge: Survival of US oyster species and stocks



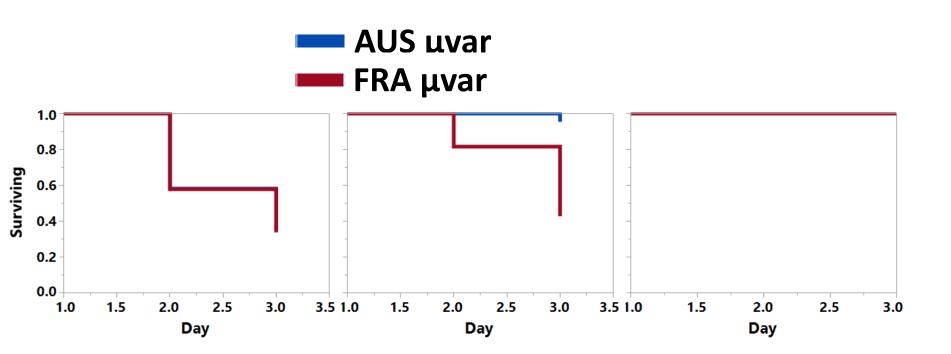
Friedman, Burge, Reece et al in prep



Injection challenge: Viral load







 Cg1 highly susceptible to both µvars Cg3 more
Cv 2 no
susceptible to
FRA μvar Friedman, Burge, Reece et al unpub data



'Natural' exposure technique

- Injected 20 juvenile (~1 yr old) *C. gigas* with 1 million copies of virus
 - OsHV-1 µvar FRA
 - OsHV-1 µvar AUS
 - OsHV-1 TB (California)
- Virus replicated for ~24 hours
 - > 5.25 x 10⁶ copies of virus added per plate
 - Relative concentration: FRA>AUS>TB at 24 hours







'Natural' exposure technique

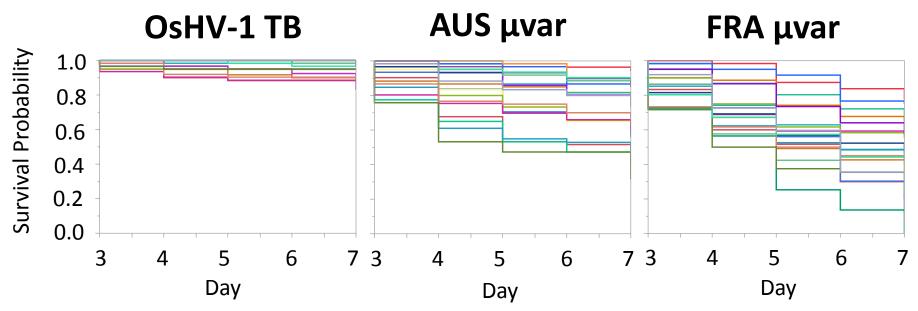
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Method of Schikorski et al 2011



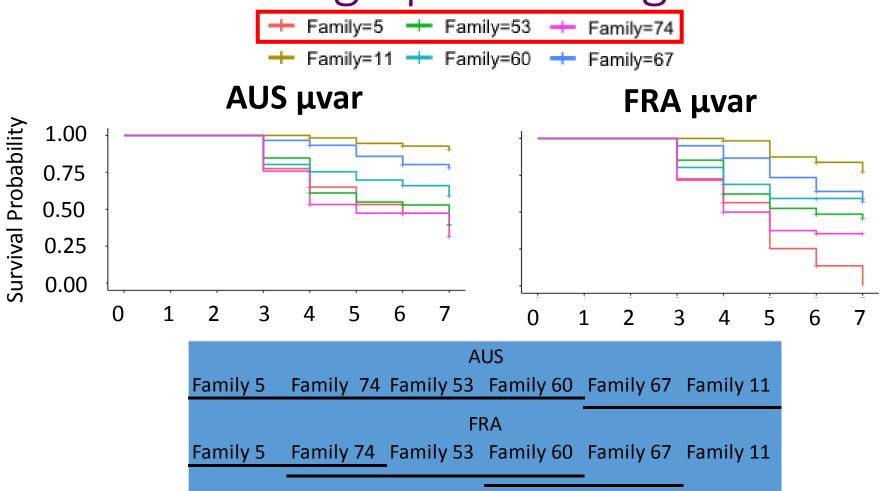


Increased transmission in Pacific oysters



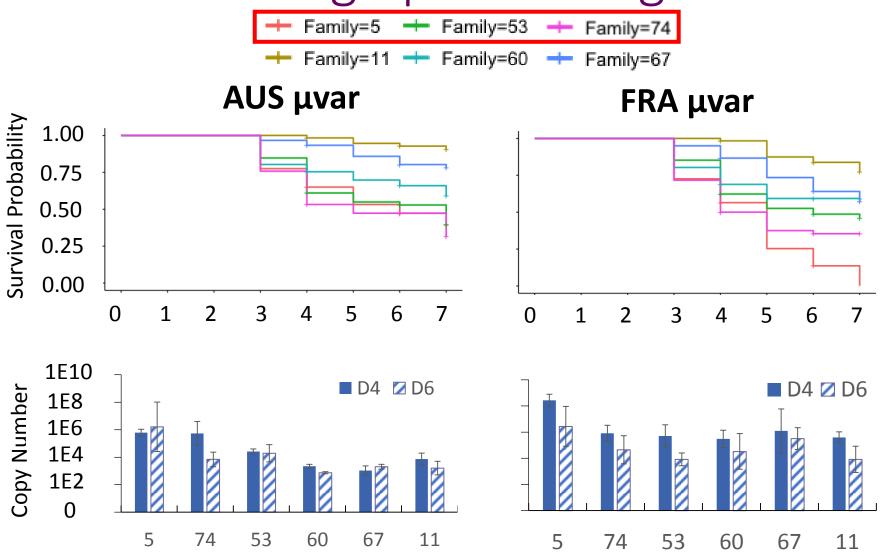
20 Molluscan Broodstock Program families 10 low/10 high based on data from Divilov *et al* in press OsHV-1 TB <AUS μvar < FRA μvar (p<0.001) Agnew, Burge, Friedman *et al* unpub data

Six low and high performing families



Log-rank Chi-Square statistics p<0.05 Using R packages survival & survminer

Six low and high performing families



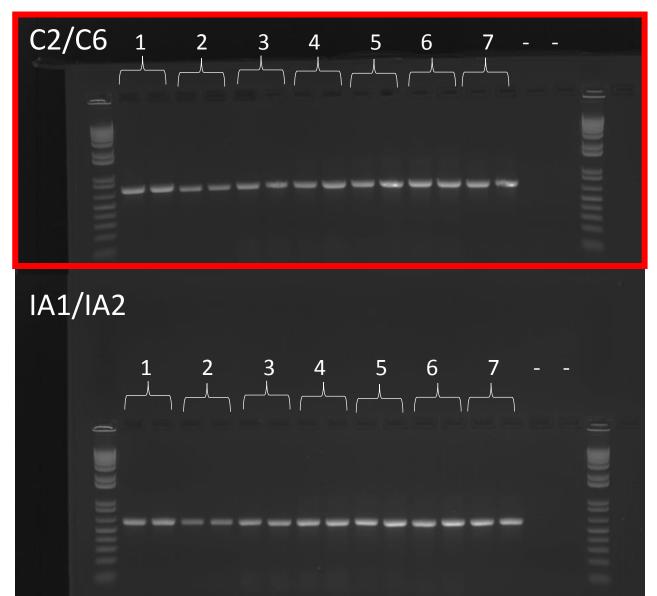
OsHV-1 in San Diego: late 2018

- First detection of OsHV-1 outside of Tomales Bay area in the US (since likely establishment in early 1990's)
- In new aquaculture venture establishing health history
- Nursery depopulated and shutdown
- Original seed source tested and "heat" test of juveniles coming from source bay
- Ongoing "sentinel" outplants by California Department of Fish and Wildlife both in San Diego & S. California

Burge, Moore, Elston, Friedman unpub data







- 1) OsHV-1 TB (2017)
- 2) OsHV-1 μ var AUS
- 3) San Diego
- 4) San Diego
- 5) San Diego
- 6) OsHV-1 μvar FRA
- 7) OsHV-1 TB (2003)

Percent Identity Matrix from Clustal 2.1 "C region"/ORF 4

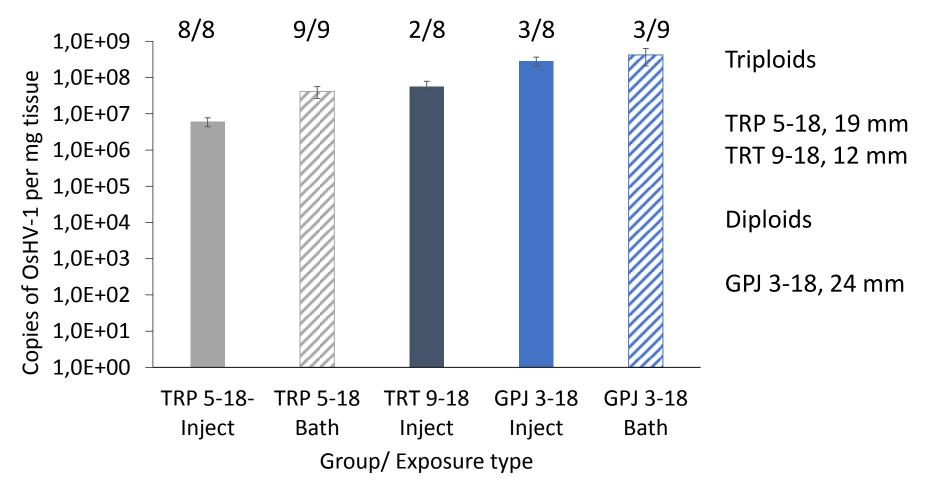
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San Diego (n=3; 3-5)		
Japanese variant	99.36	
New Zealand µvar	99.68	X
OsHV-1 μvar FRA	99.84	*
Italy μvar full genome	99.84	ズ
OsHV-1 µvar AUS	99.84	×
OsHV-1 ref full genome	95.30	\Rightarrow
OsHV-1 TB (2003)	94.86	*
La Cruz, Mexico	95.06	\bigstar



Pilot transmission study: February 2019

Injected with 25,000 copies* or exposed by bath with 260,000* total copies; 8-9 animals per trt



Next Steps Needed

- Though data indicate non-Pacific oyster species are less susceptible, more life stages and lines of oyster species should be tested & assessed as potential vectors
- Additional characterization of OsHV-1 from San Diego
- Development of a high-throughput assay to differentiate variants
- Gene expression analysis of Eastern vs. Pacific oysters
- Pair laboratory/field trials with quantitative genetic studies to help identify markers of resistance/susceptibility for QTL development
- <u>Education Goal to reduce potential for introduction</u>



Acknowledgements





















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Thank you!





Results: CA OsHV-1

