



GENETIC VARIATION IN PACIFIC OYSTERS FOR RESISTANCE TO *Ostreid herpesvirus-1*

Peter Kube

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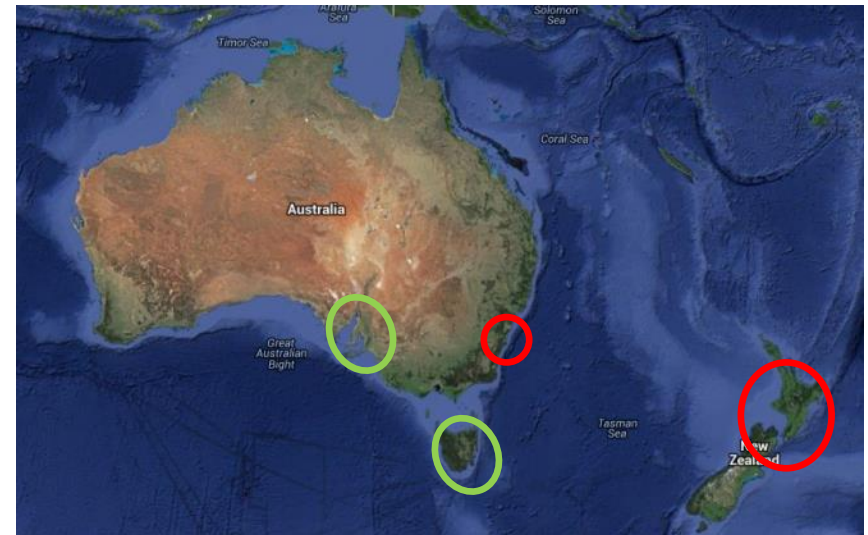
The Spread of Oyster Herpes Virus

Ostreid herpesvirus 1

- Affects only Pacific oysters
- New micro-variant with high virulence

Outbreaks:

- France 2007-08
- New Zealand Mar 2010
- Australia Nov 2010



The Spread of Oyster Herpes Virus

SPREAD IN AUSTRALIA:

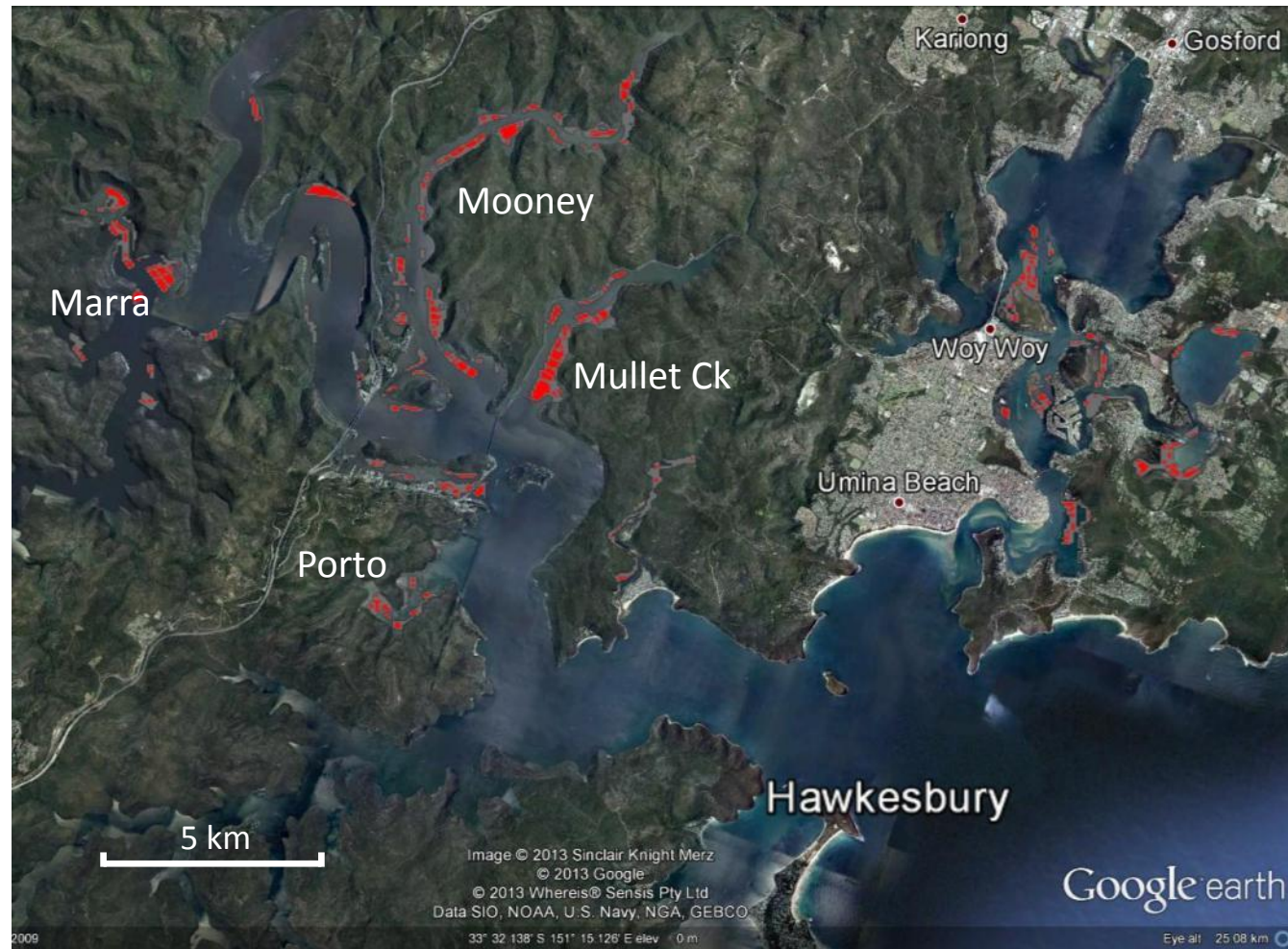
- Georges River Nov 2010
- Sydney Harbour 2011
- Hawkesbury River Jan 2013

Sydney Region New South Wales Central Coast



The Spread of Oyster Herpes Virus

- DAY 1 10 AM:
First sighting
(30% mortality
on one lease)
- DAY 1 5 PM:
mass mortality
on that lease
- DAY 3:
10 million dead
oysters (\$3M loss)
- DAY 8: Entire
system affected



Aims



Breed for resistance to OsHV-1



The science challenge:

- Is there resistance in our population?
- Genetic parameters of resistance?
- How can we measure resistance?
- How long before economically useful resistance?

Data analysis and summary

- Australian breeding population (up to 9 generations)
- Six field challenges and two laboratory challenges
- Two ages (4 to 6 months and 12 months)
- Analysed using ASReml;
Sire model with pedigree structure (binary data)

Number year classes challenged	3	2011, 2012, 2013
Total number of families	175	
Total number of parents	345	
Number animals challenged	56,658	

FIELD CHALLENGE NOV 2012
ADULTS (AGE 12 MONTHS)





A top-down view of a black mesh tray filled with numerous dead oysters. The oysters are scattered across the tray, showing various stages of decay and discoloration. Some shells are open, revealing the interior. The colors range from dark purple and black to light tan and yellowish. In the upper right corner, a portion of an orange-handled oyster knife is visible, with the letters 'SA' written on the handle. In the lower left corner, a white rectangular box contains the text 'SURVIVAL = 0%'.

SURVIVAL = 0%



52%

5%

8%

0%

3%

25%

2%

3%

0%

3%

3%

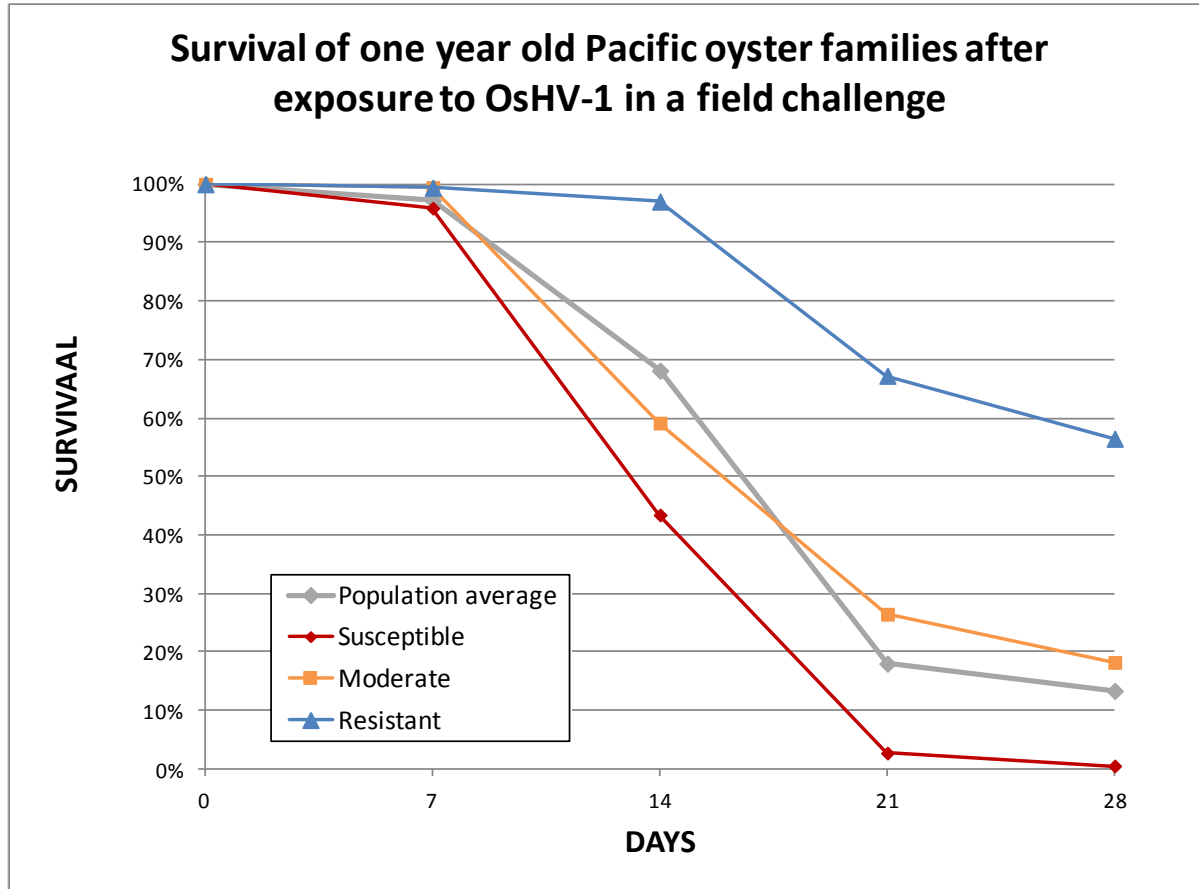
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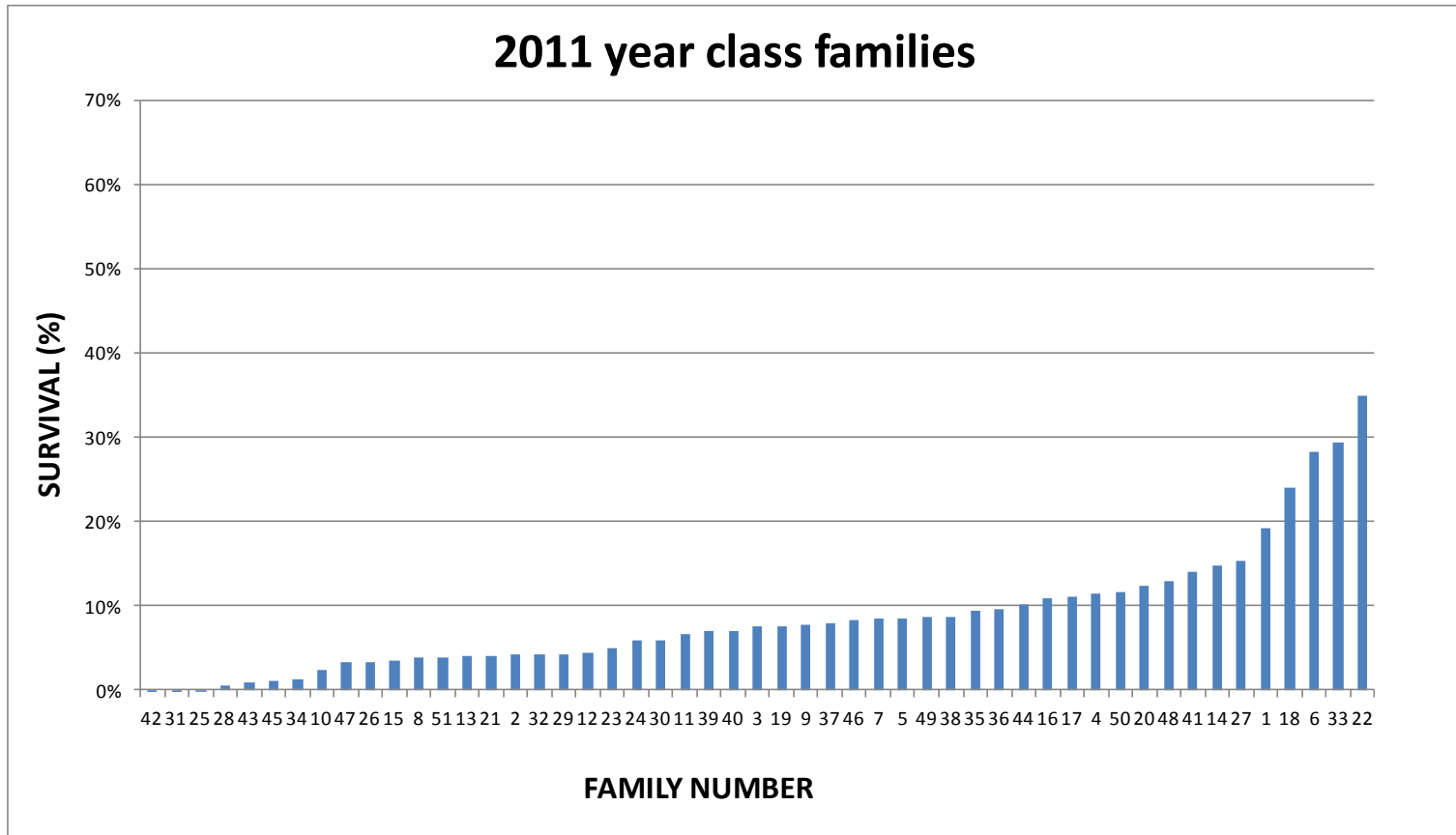
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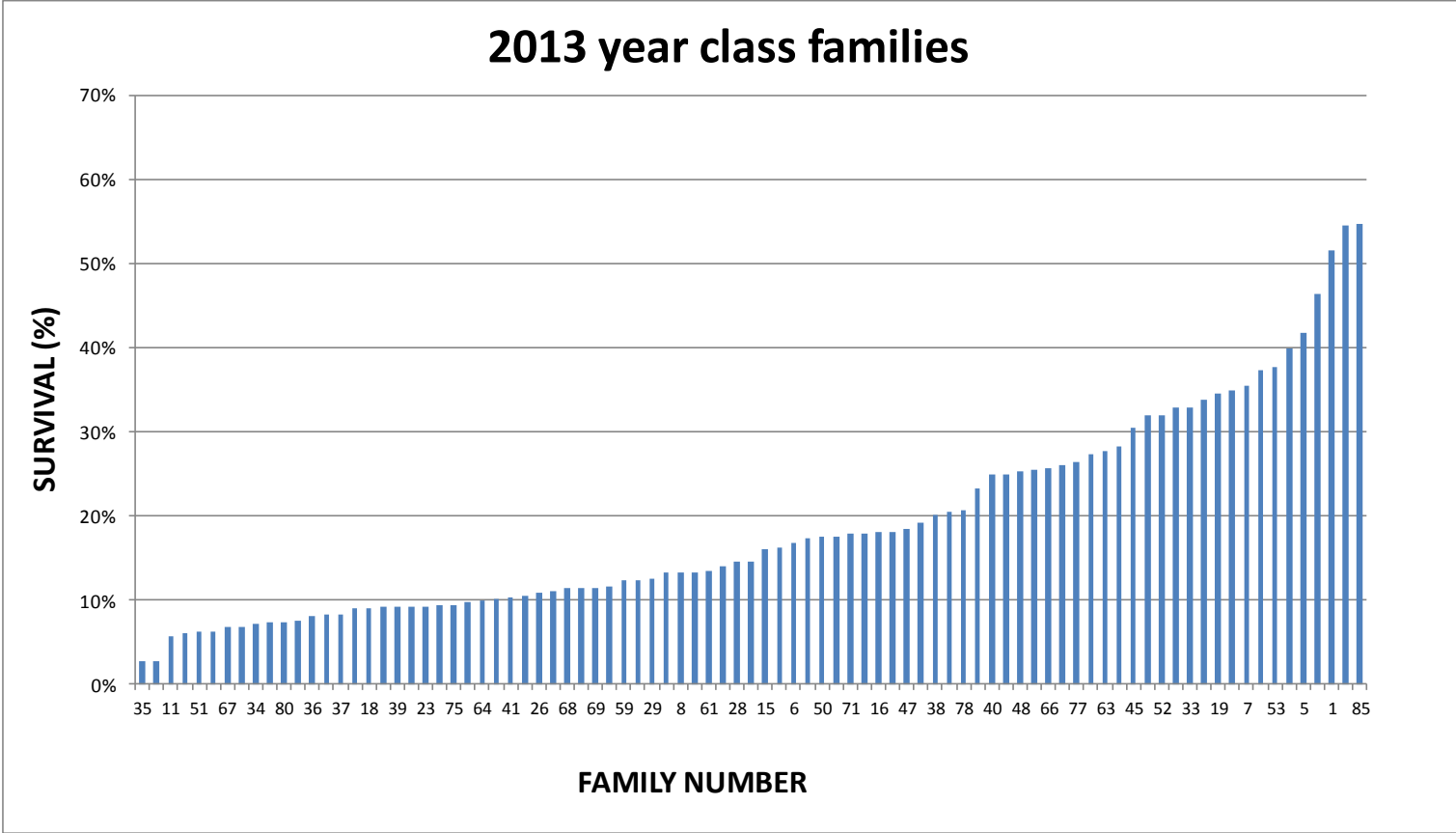
Disease progression in a field challenge



Unselected families



Selected families



Difficult to get a good field challenge

TRIAL	Description	h^2	Test day survival
1	2011 YC spat	0.38	47% <input checked="" type="checkbox"/>
2	2011 YC adults	0.18	20% <input checked="" type="checkbox"/> <input checked="" type="checkbox"/>
3	2012 YC spat	0.53	91% <input checked="" type="checkbox"/>
4	2012 YC adults	0.60	29% <input checked="" type="checkbox"/>
5	2013 YC spat (a)	-	No mortality <input checked="" type="checkbox"/>
6	2013 YC spat (b)	-	No mortality <input checked="" type="checkbox"/>
7	2013YC adults	0.46	18% <input checked="" type="checkbox"/>

Laboratory Disease Challenge

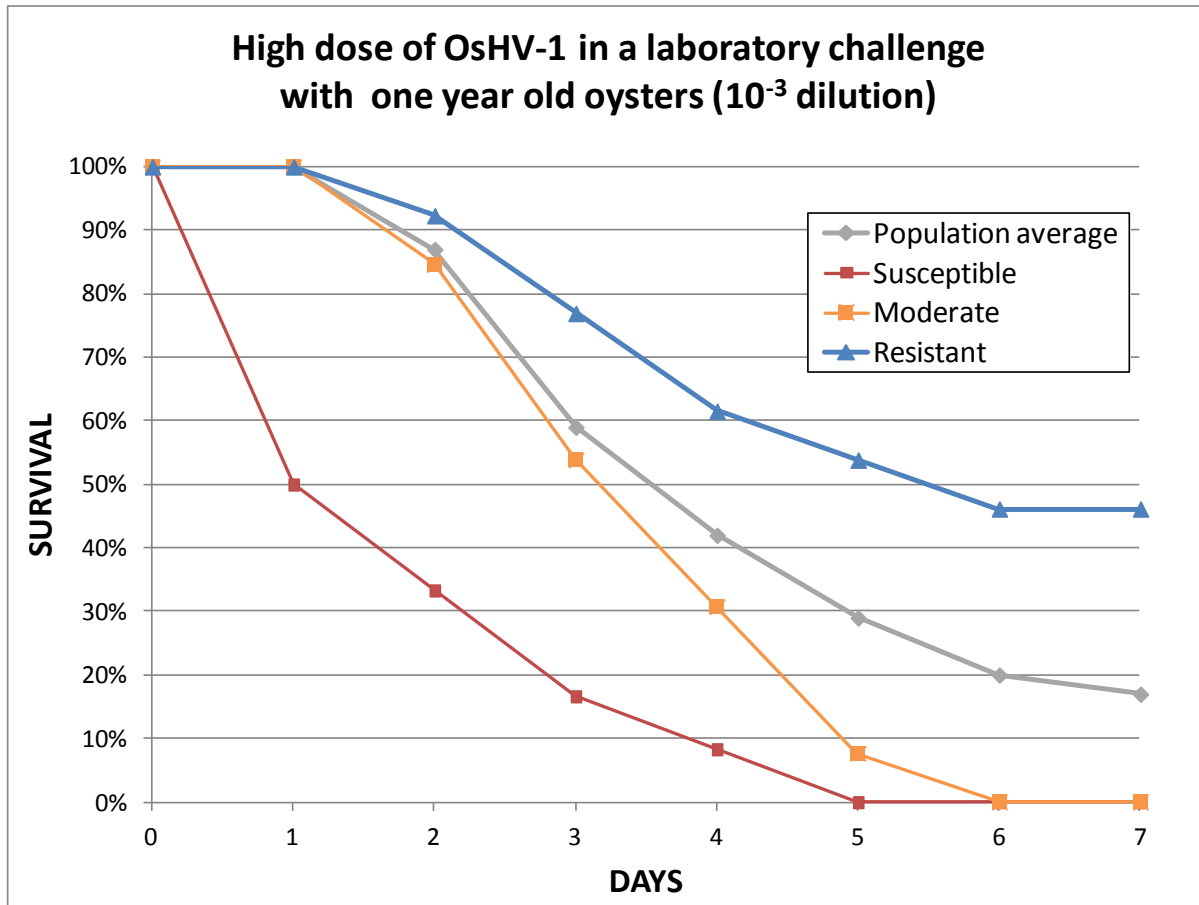
Done in a biosecure laboratory
(NSW Department of Primary Production)

Uses stock virus solution
(cryo-preserved) stock

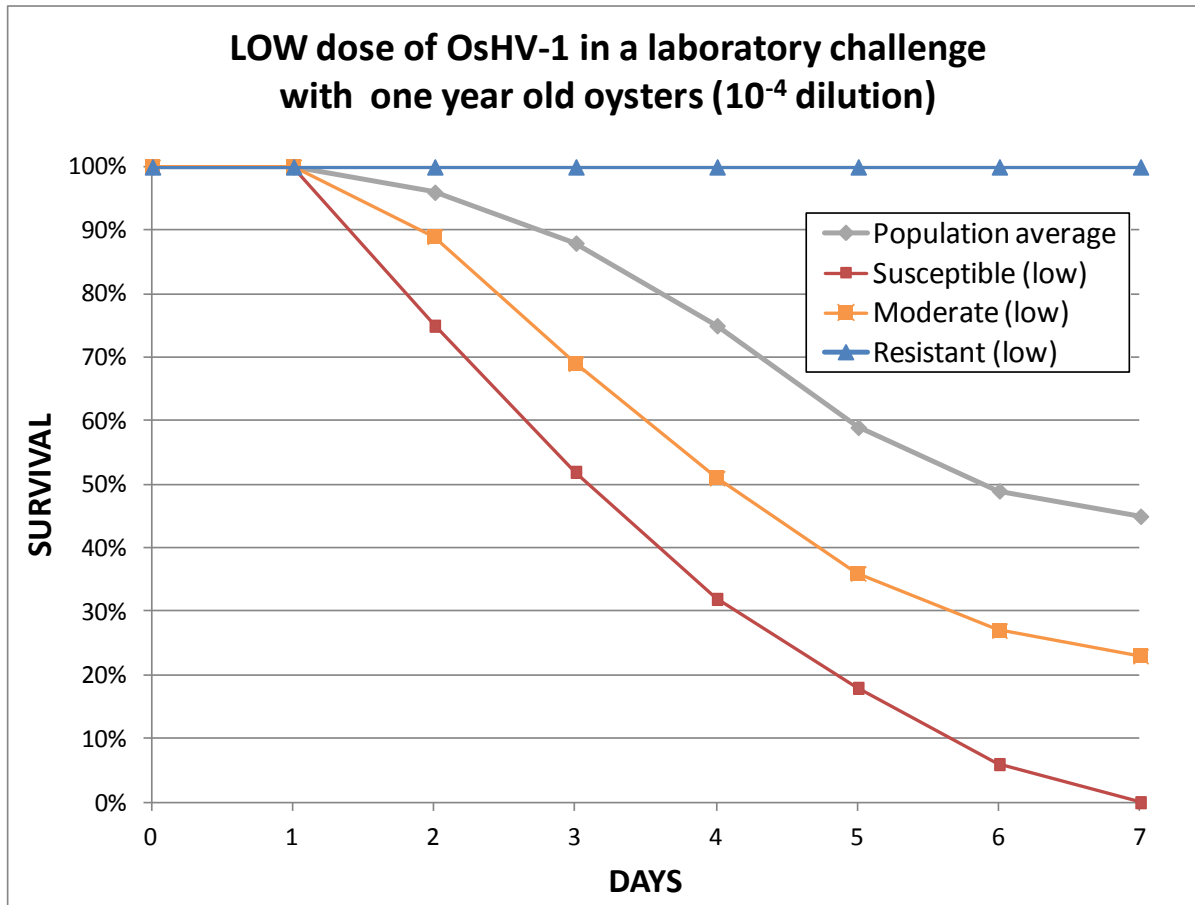
Immersion of relaxed oysters



Disease progression in a lab challenge



Disease progression in a lab challenge (low dose)



Genetic parameters – heritabilities

TRAIT	h^2 (<i>se</i>) observed	h^2 underlying
Field survival (spat)	0.39 (0.09)	0.61
Field survival (adults)	0.27 (0.04)	0.55
Laboratory survival (spat)	0.18 (0.11)	0.28

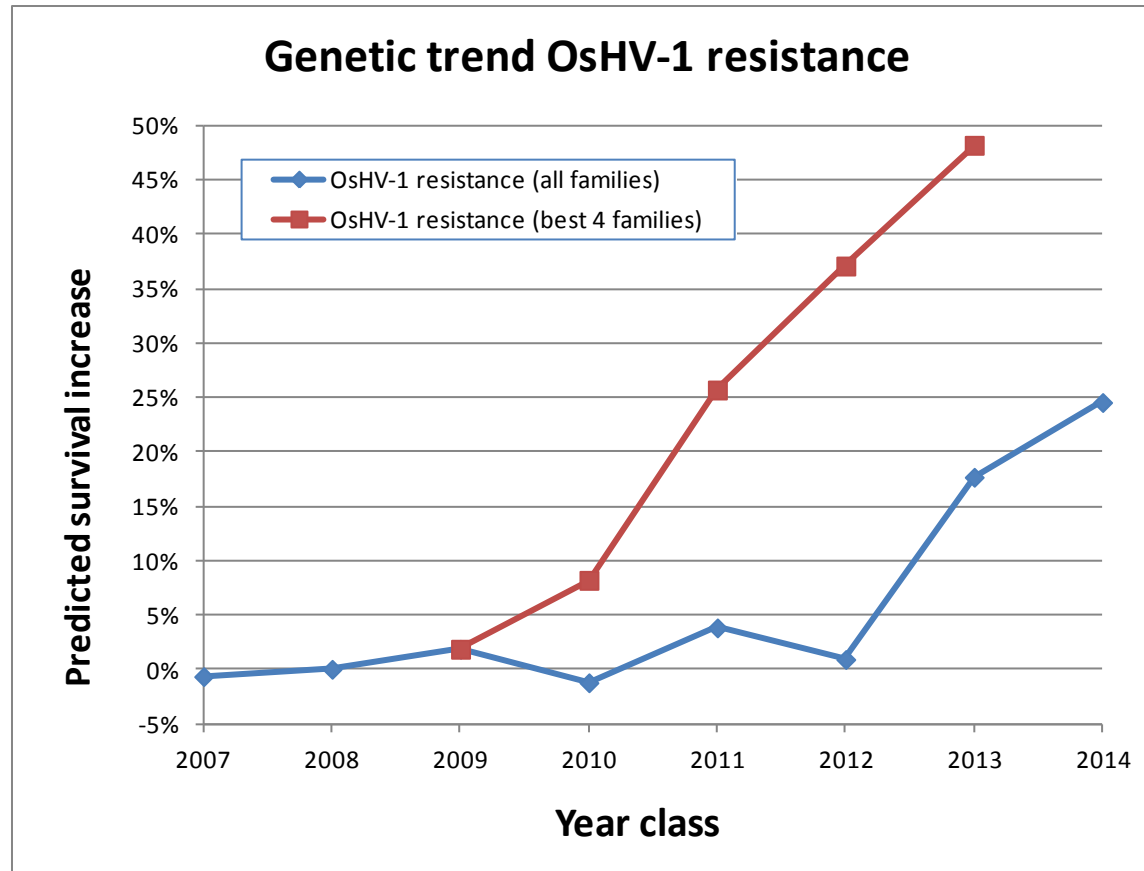
Genetic parameters – genetic correlations

TRAIT	h^2 (observed)	h^2 (underlying)	Genetic correlations r_g (se)	
			Field surv. (spat)	Field surv. (adult)
Field survival (spat)	0.39	0.61		
Field survival (adults)	0.27	0.55	0.85 (0.08)	
Laboratory survival (spat)	0.18	0.28	0.71 (0.30)	0.61 (0.24)

Genetic correlations – test days

	Field challenge survival (r_g)		
	Day 7	Day 14	Day 21
Day 14	0.74		
Day 21	0.02	0.77	
Day 28	0.03	0.77	0.99

Genetic gains



Conclusions

- There is genetic variation for OsHV-1 resistance in our population
- Field challenges presents logistic difficulties for applied breeding, and a reliable laboratory challenge is needed
- Selective breeding is providing a means to mitigate the impact and risk of this disease

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