

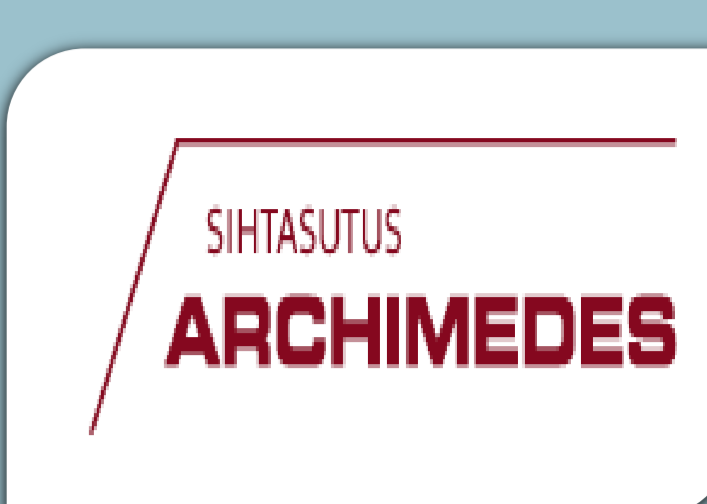
# Comparison of redox enzyme activities in two Atlantic salmon (*Salmo salar* L.) populations: Insight to the trade-offs between physiology and survival during early life stages



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## Introduction

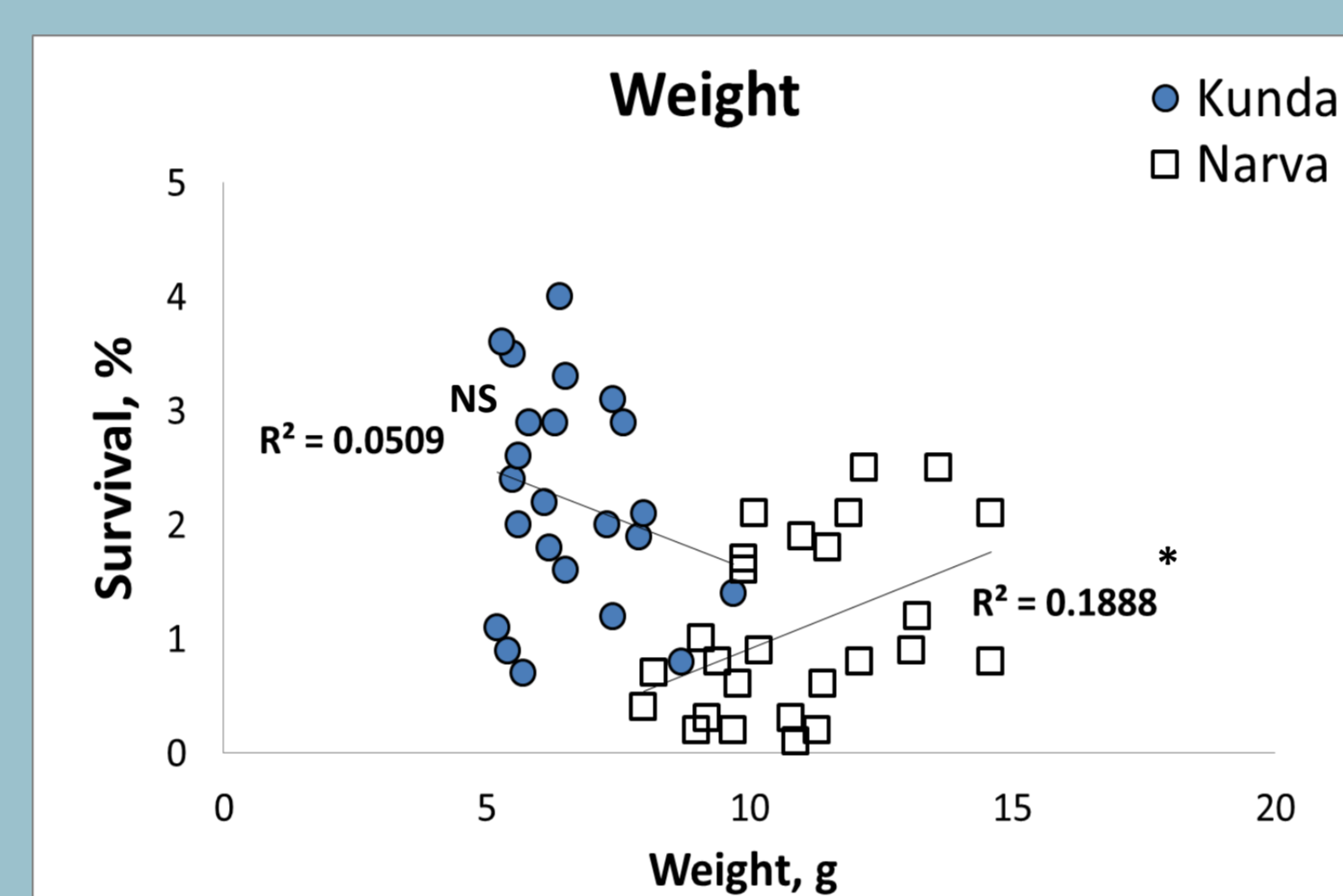
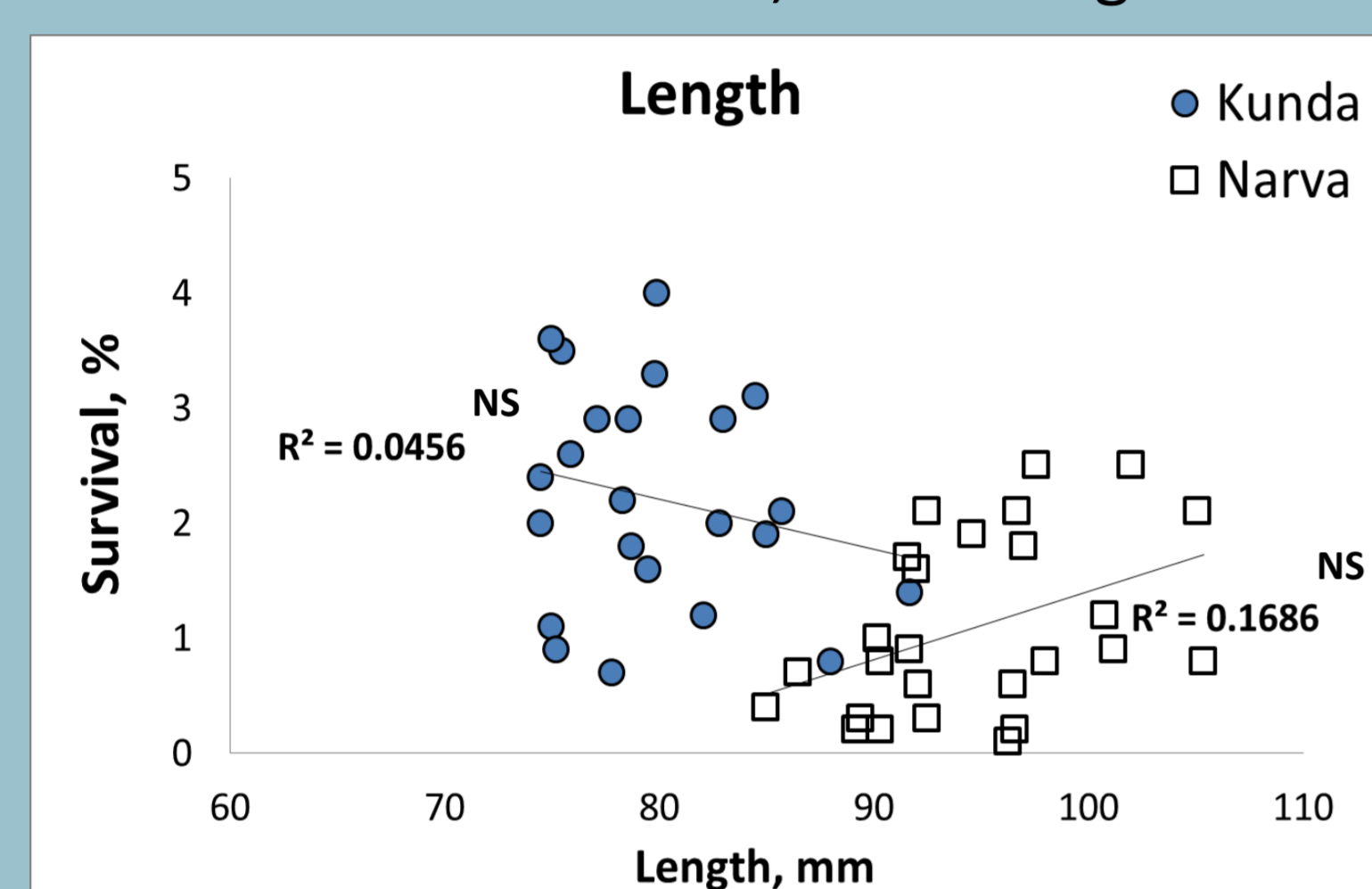
- Oxidative stress (OS) is an imbalance of prooxidants and antioxidants
- OS during early development may have long term consequences on various fitness related traits and ultimately affect their survival
- Currently very little is known about the heritability of OS and the relationships between OS, fitness and survival of natural populations in wild

## Aims

1. Assess the survival and growth of 0+ salmon in multiple natural environment
2. Determine the heritability of three redox enzyme activities (Glutathion Reductase; GR, Glutathion-S-Transferase; GST and Superoxide Dismutase; SOD) in pre feeding fry of two genetically different Baltic Sea salmon populations
3. Evaluate the possible linkage between the physiological traits (e.g. redox status/oxidative stress) and fitness (survival, growth) in wild

## Results

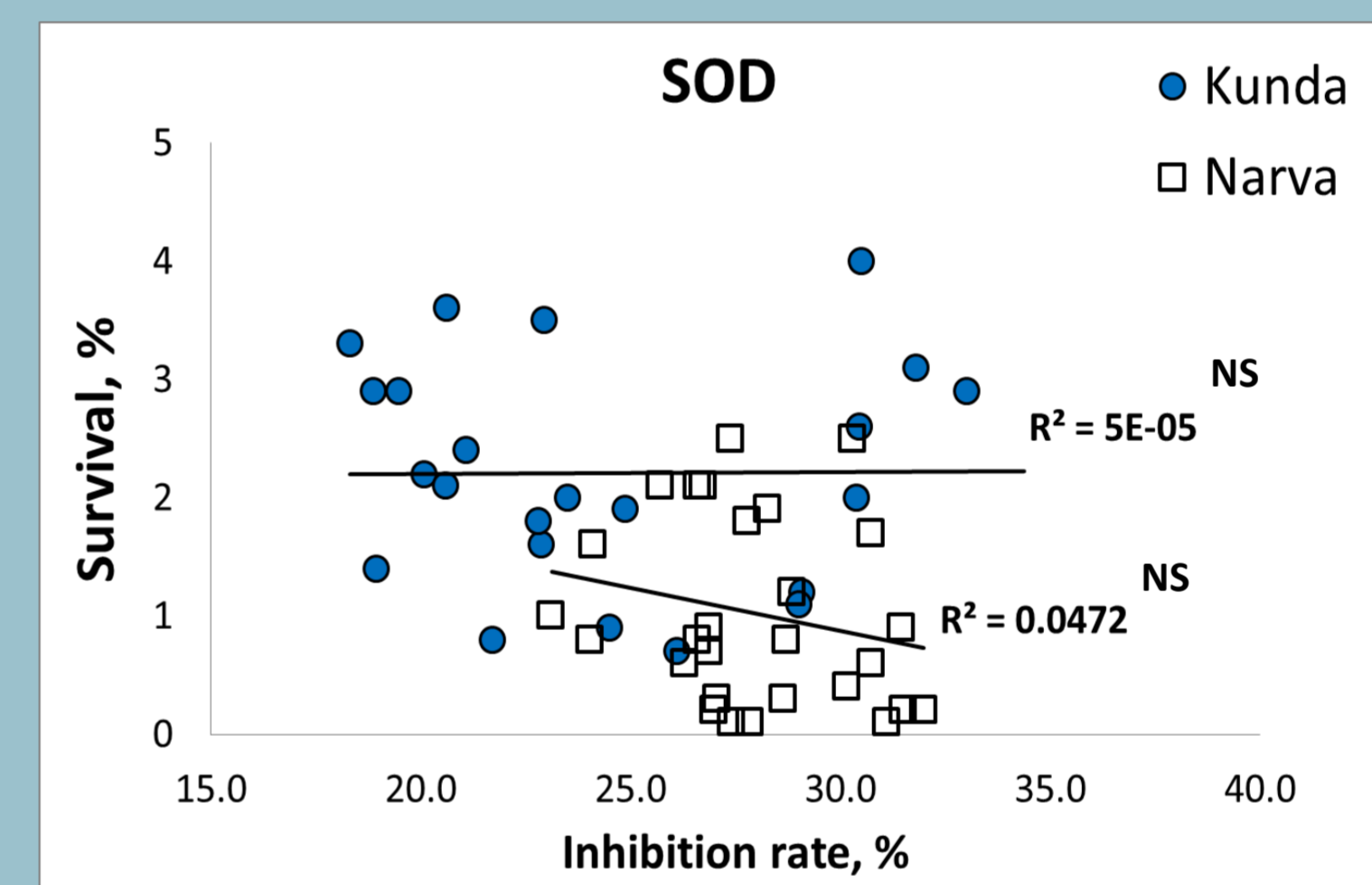
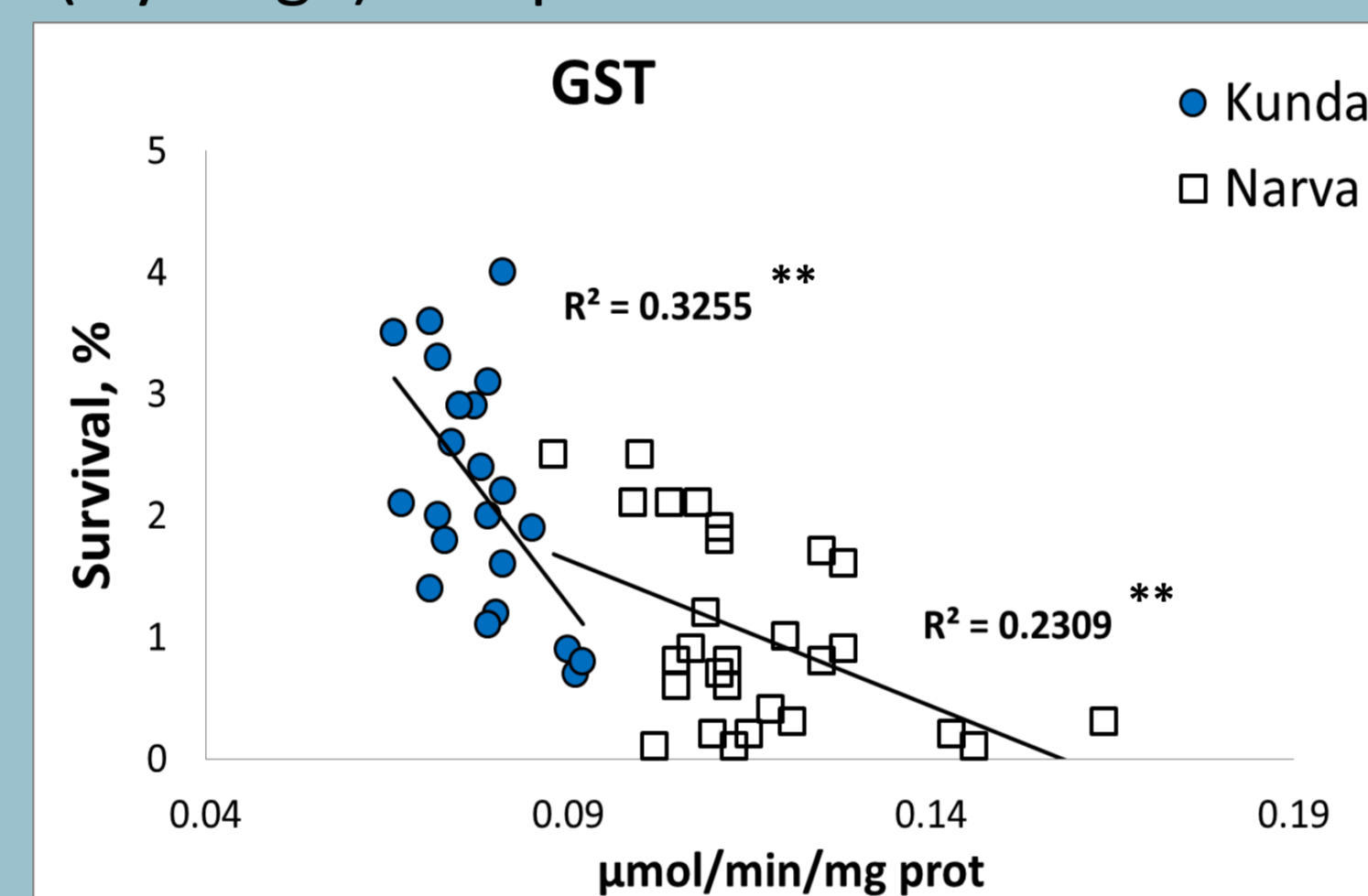
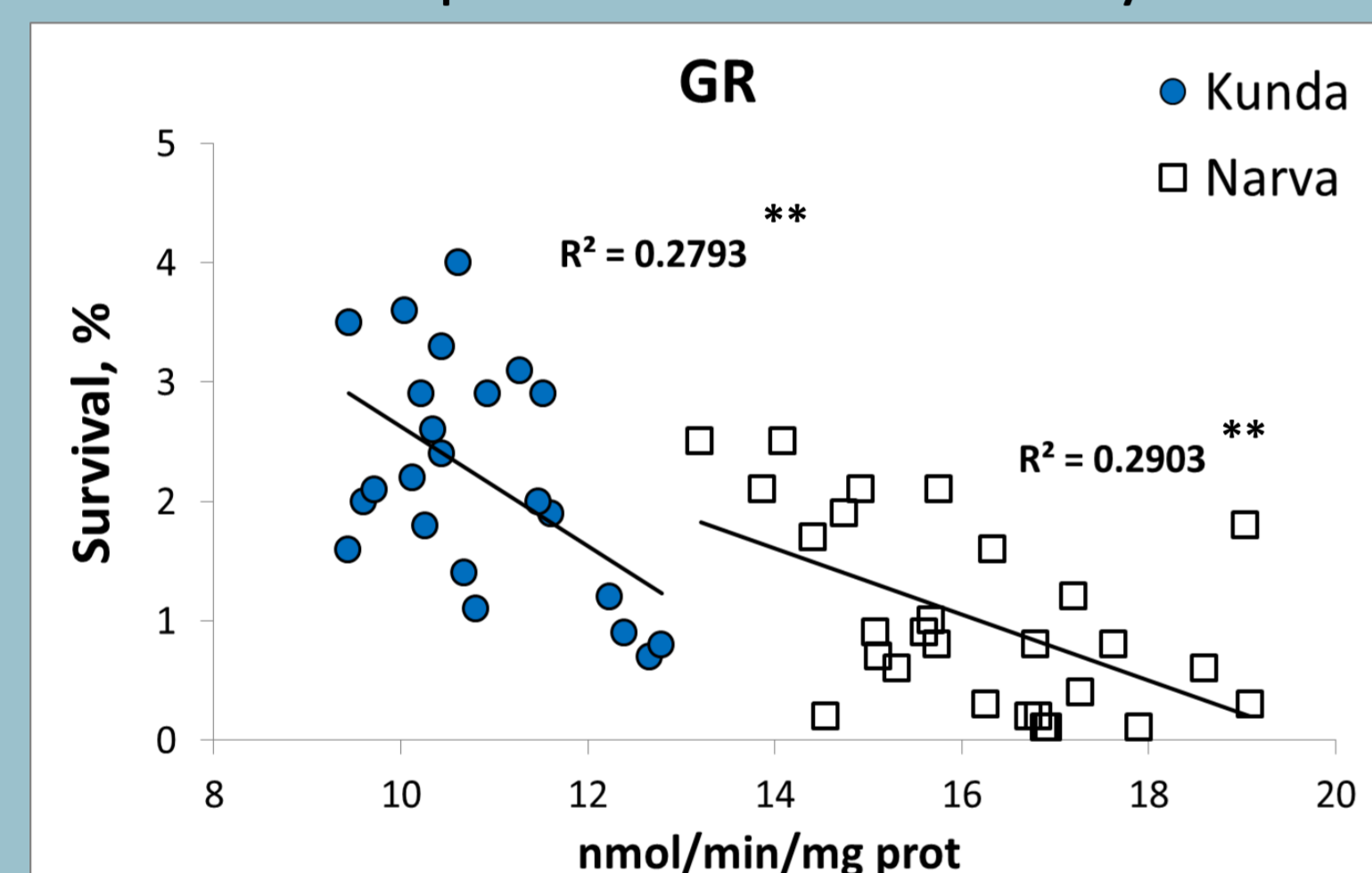
1. Significant differences in growth and survival were found after first summer in wild; p-values are marked as bold stars, NS - no significance
2. The narrow-sense heritability ( $h^2$ ) estimates for three redox enzyme activities in two Baltic salmon population



Population	Enzyme	Model	DIC	$h^2$	p value
Kunda	GR	$V_p = V_a + V_y + V_r$	1533	0,08	NS
<b>Kunda</b>	<b>GST</b>	<b><math>V_p = V_a + V_r</math></b>	<b>-2559</b>	<b>0,31</b>	<b>***</b>
Kunda	SOD	$V_p = V_a + V_y + V_r$	2733	0,07	NS
<b>Narva</b>	<b>GR</b>	<b><math>V_p = V_a + V_y + V_r</math></b>	<b>1451</b>	<b>0,26</b>	<b>***</b>
<b>Narva</b>	<b>GST</b>	<b><math>V_p = V_a + V_r</math></b>	<b>-2714</b>	<b>0,72</b>	<b>***</b>
Narva	SOD	$V_p = V_a + V_y + V_r$	2840	0,10	NS

The variance components reported are total phenotypic variance ( $V_p$ ), additive variance ( $V_a$ ), variance described by yolk sac area ( $V_y$ ) and residual variance ( $V_r$ ). The best model (shown in bold text) was chosen according to DIC – Deviance Information Criterion calculated from MCMCglimm run.

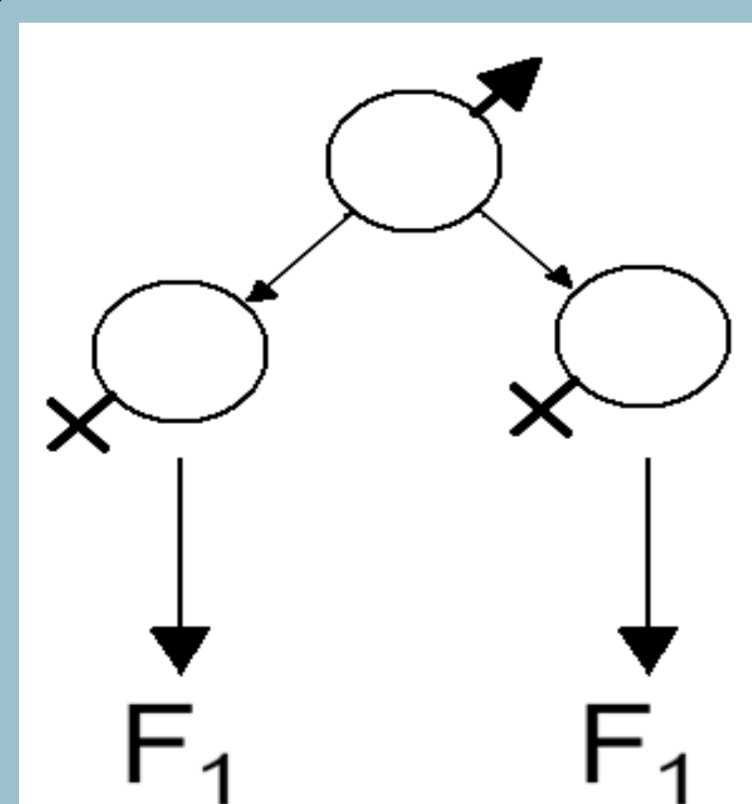
3. Relationships between redox enzyme activities (fry stage) and parr survival



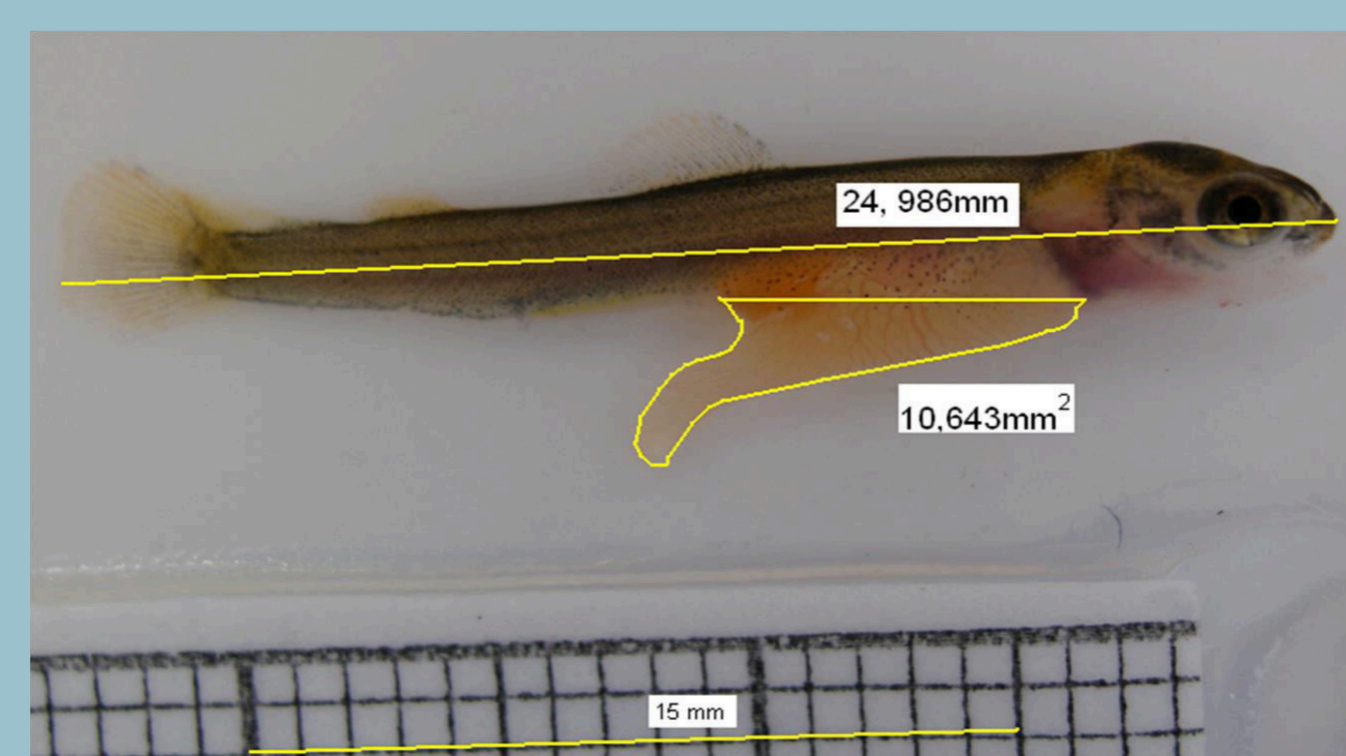
## Conclusions

- Two studied salmon populations differ significantly already in their early developmental period in terms of redox enzyme activities
- Heritable genetic variation exists in two redox status/oxidative stress related enzymes (GR and GST but not in SOD) in pre feeding fry
- GR and GST enzyme activities are negatively correlated with growth and survival in natural habitat
- The inter-population differences in redox status during early life stages may explain considerable proportion of variation in fitness and survival in wild

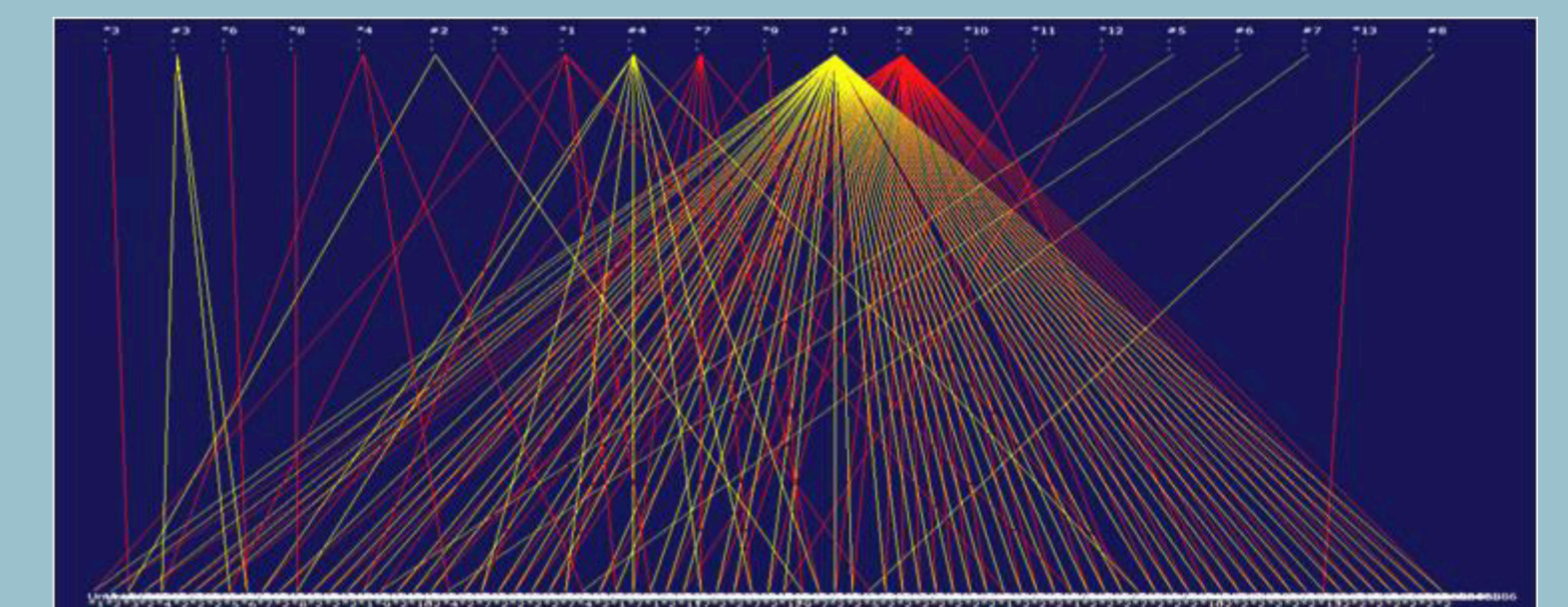
## Experimental design



- 52 half-sib families were created
- Length and yolk-sac area was measured in 20 unfed fry per family
- Enzyme activities were measured in quadruplicates in 384-well plate format



- ~ 221 000 pre feeding fry in total were stocked into 2 natural rivers (2 different sites per river)
- 1869 0+ salmon parr were sampled in total after one summer in wild
- Individuals were finclipped for DNA analyses and later assigned to families utilizing 18 microsatellite markers



## Acknowledgements

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