Assessing the Effects of Parental Traits On the Production of

Hatchery Spring Chinook Salmon Minijacks

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Study questions:

Is the proportion of minijacks produced by natural origin (NO) Yakima River spring Chinook broodstock affected by parental age and other phenotypic traits?

Is body size of fry at emergence (swim up) and at smolt stage affected by parental age or other parental phenotypic traits?

Study Design:

- 3 broodyears: 2014 to 2016
- Factorial mating of NO broodfish of different ages:
 - o females (Age 4 or 5)
 - o males (3 [jack], 4 or 5); added Age 1 NO microjacks in BY 2015 and 2016

Hatchery Origin Age 2 Minijack

Natural Origin Age 1 Microjack Matures in first year

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BY14	Females				
Males	Age 4	Age 5			
Age 3 Jack	200 eggs	200 eggs			
Age 4	200 eggs	200 eggs	1 des		
Age 5	200 eggs	200 eggs			

BY15 & BY16	Females			
Males	Age 4	Age 5		
Age 1 (micro-jack)	200 eggs	200 eggs		
Age 3 Jack	200 eggs	200 eggs		
Age 4	200 eggs	200 eggs		
Age 5	200 eggs	200 eggs		

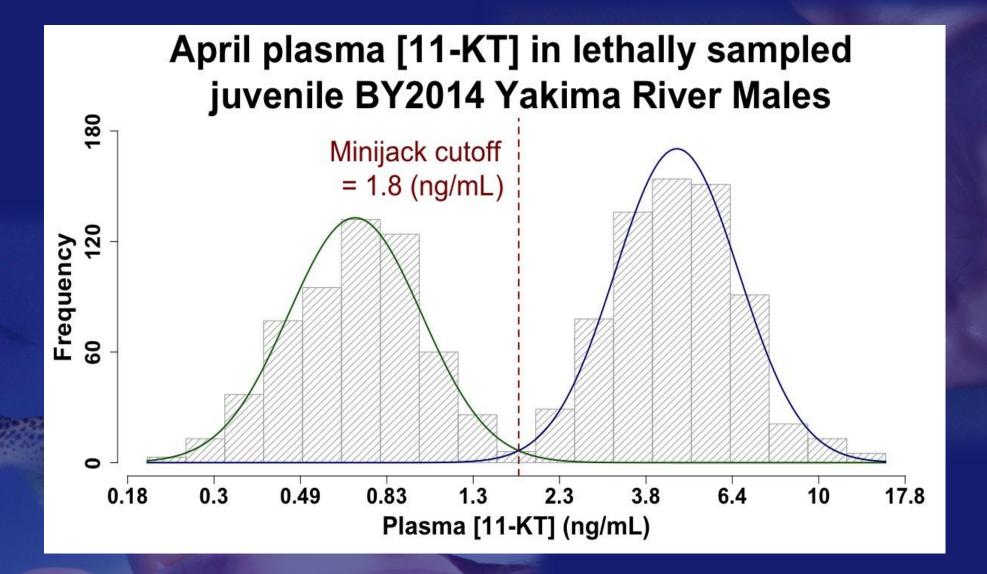
Females

		Age	4	5	4	5	4	5	4	5
	Age	Carc ID	F14-01	F14-02	F14-03	F14-04	F14-05	F14-06	F14-07	F14-08
	3	M14-01								
	4	M14-02								
	5	M14-03								
in.	3	M14-04								
	4	M14-05								
Males	5	M14-06								
	3	M14-07								
and the second	4	M14-08								
1	5	M14-09								
	3	M14-10								
	4	M14-11								
6	5	M14-12								

Study Design:

- 3 broodyears: 2014 to 2016
- Factorial mating of NO broodfish of different ages:
 - o females (Age 4 or 5)
 - o males (3 [jack], 4 or 5); added Age 1 NO microjacks in BY 2015
- Rear juveniles (50 fry per mating) to smolt stage (April) in common environment
- Blood samples collected for 11-KT assays
- Tissue samples collected for genotyping and parentage analysis
- One Time for BY 2014
- Live sampled juveniles 11KT, PIT tagged and then reared to full maturity (Sept)

to confirm April 11-KT measures reflect actual male maturation (minijack) status in September –
draft manuscript in prep with L. Medeiros lead author



Established a maturation threshold ("cutoff") of 1.8 ng/mL in fish lethally sampled in April of 2016 (n = 1,224). [Slide produced by L. Medeiros.]

BY 2014 Sampling Completed

<u>Number</u>

- 1,254 smolts sampled April 2016
 - 459 PIT tagged April, sacrificed Sept 2016
- 1,414 smolts with data for both 11-KT + genotypes
 - 57 full sib progeny groups with both parents of known age, (n=1,170; average = 21 males/progeny group)

BY 2015

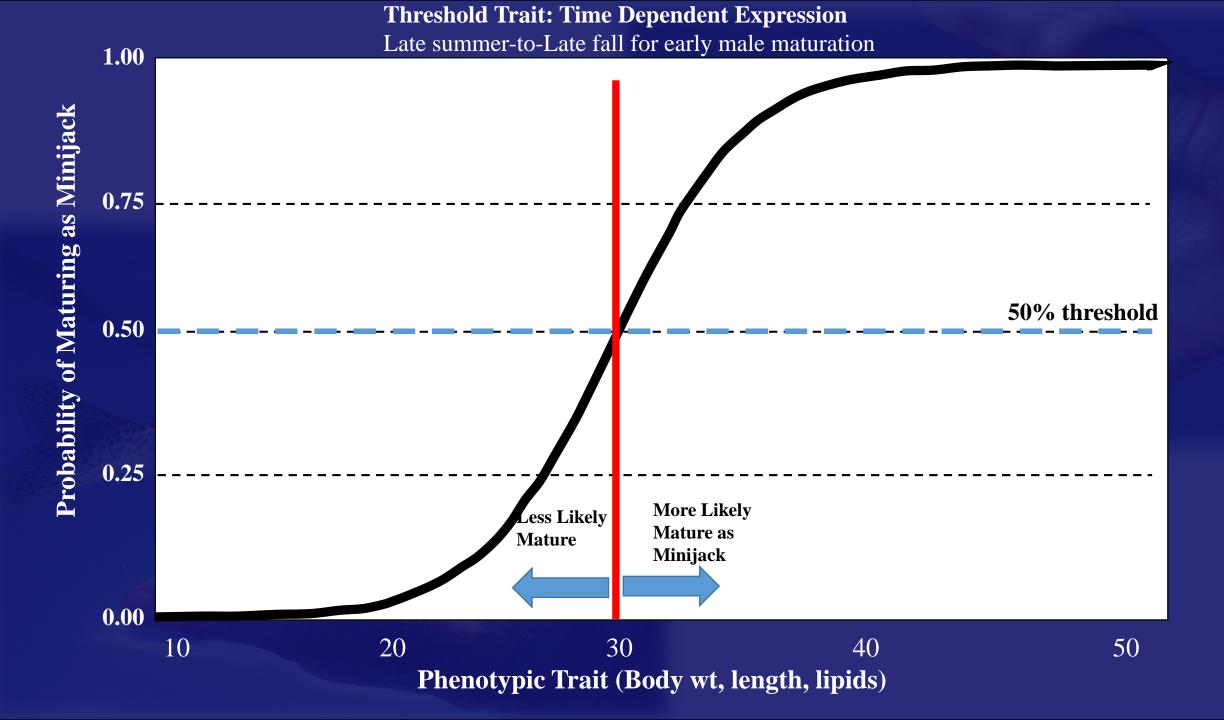
- All factorial crosses made
- Smolts were sampled April 2017
- 11-KT and genotype analyses completed by Fall 2017

BY 2016

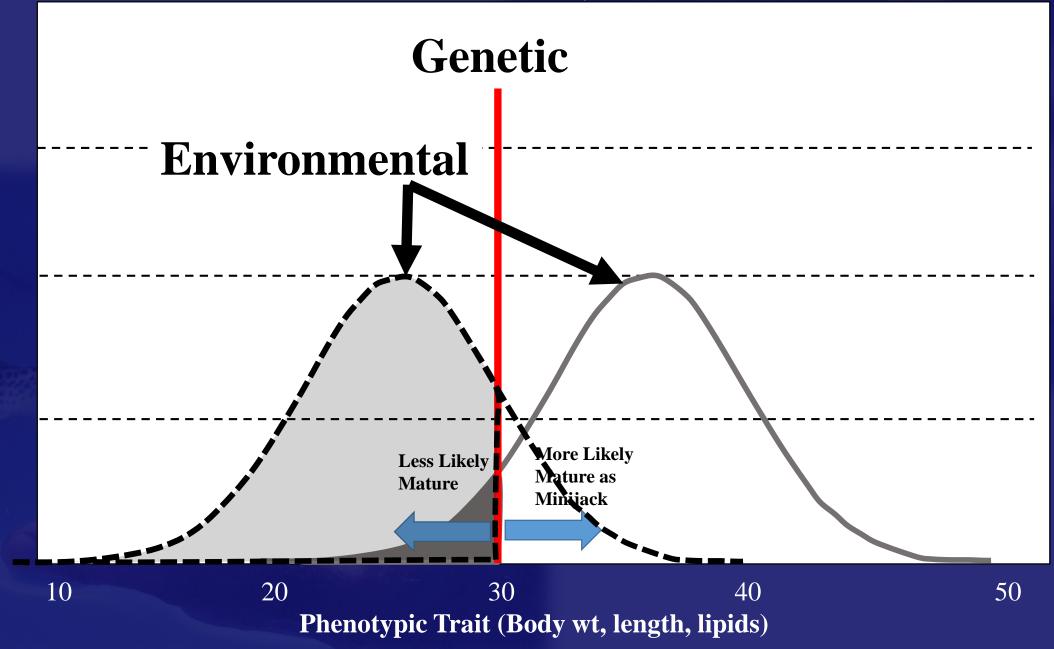
- All factorial crosses made
- Smolts will be sampled April 2018
- 11-KT and genotype analyses completed by Fall 2018

Fry Samples

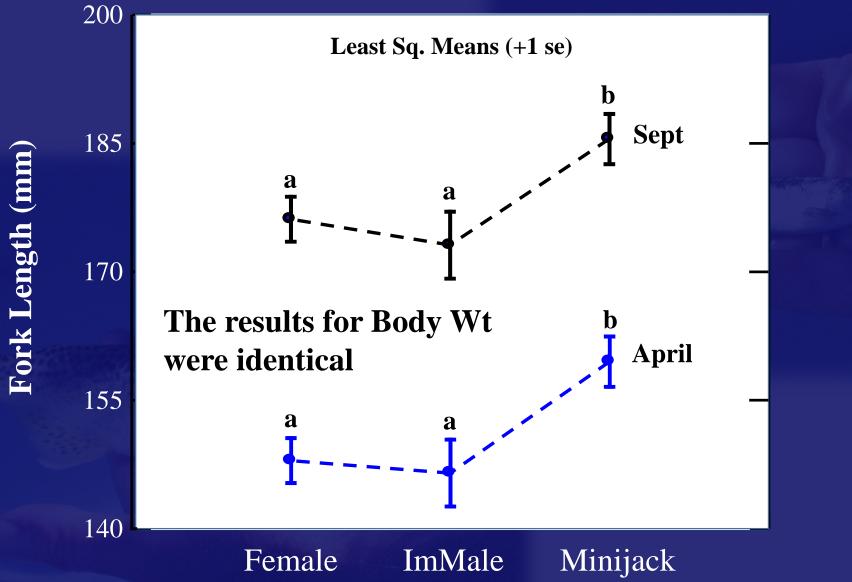
BY 2014 Number **370** - NO fry sampled March 2015 5 fry per factorial mating; parents of known age, length, body wt, egg size, etc. **BY 2015** Number **305** - NO fry sampled March 2016 **BY 2016** Number xxx - NO fry sampled March 2017

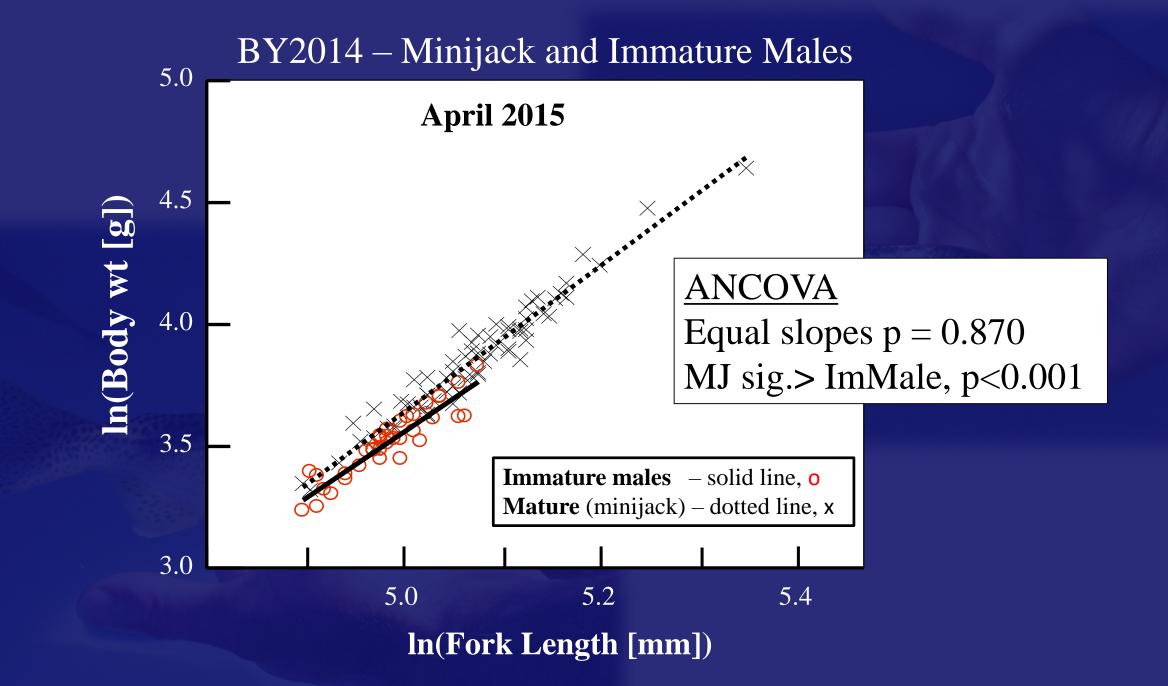


Threshold Trait: Time Dependent Expression Late summer-to-Late fall for early male maturation



BY14 Smolt Data

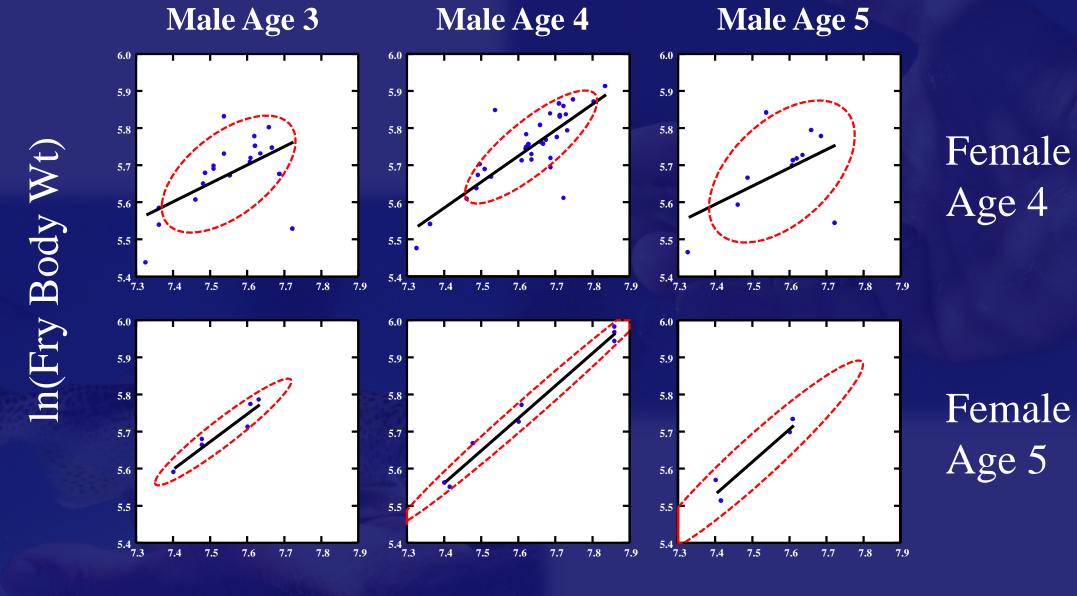




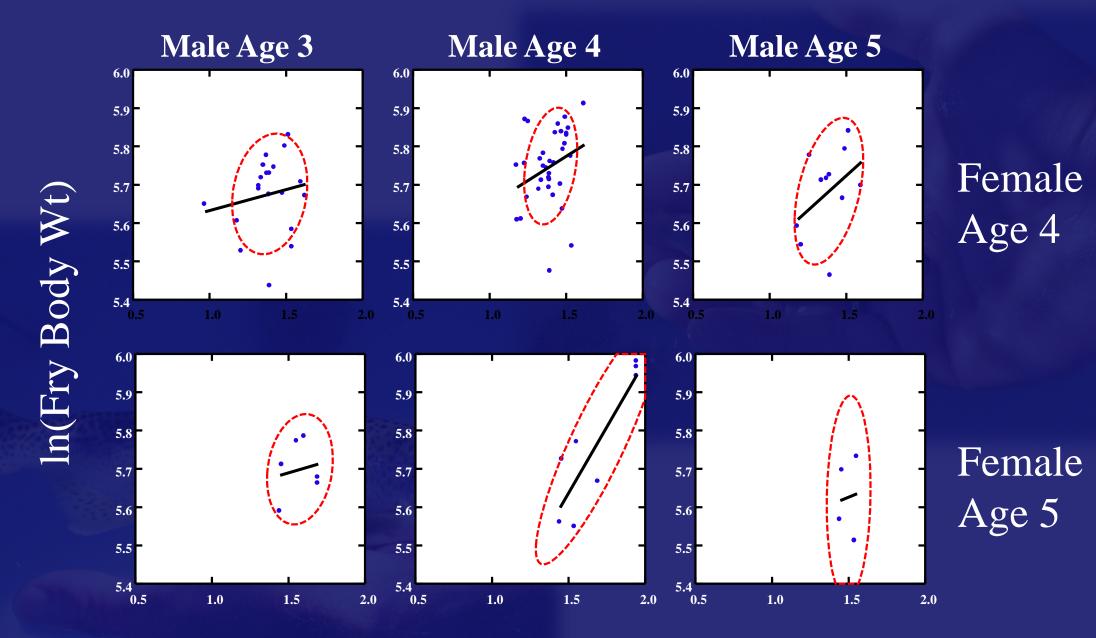
Linear Model for Fry Analyses: Female and Male Age Effects

Fry Body wt ~

Female Age + Male Age + In(Egg wt) + In(Female Length) + In(Female Body wt) + (Male Age*Fem Age)



ln(Egg wt)



ln(Female Body Wt)

Linear Model for Fry

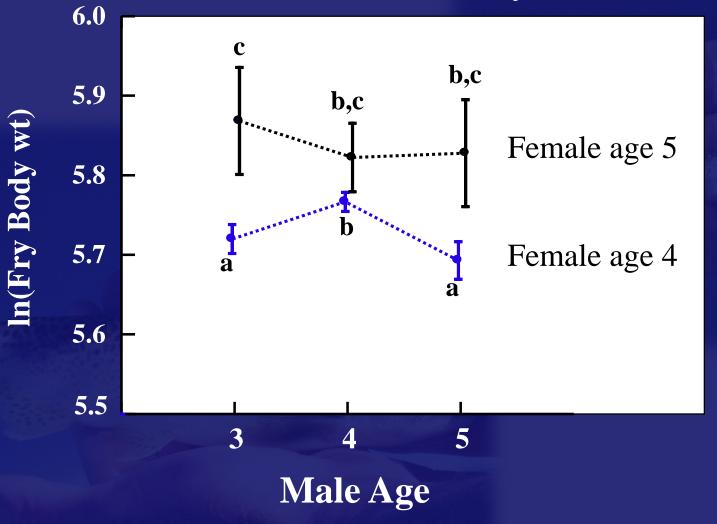
Fry Body wt ~

Female Age + Male Age + In(Egg wt) + In(Female Length) + In(Female Body wt) + (Male Age*Fem Age)

BY2014

Analysis of Variance								
Source	Type III SS	lf I	Mean Squares	F-ratio	p-value			
Female Age	0.09039	1	0.09039	21.65248	0.00001			
Male Age	0.01498	2	0.00749	1.79431	0.16846			
ln(Egg W)	0.13667	1	0.13667	32.73858	<0.00001			
LN_POHP (female)	0.18481	1	0.18481	44.27144	<0.00001			
LN_BW (female)	0.34511	1	0.34511	82.66974	<0.00001			
Male Age*Fem Age	0.03724	2	0.01862	4.46057	0.01253			
Error	1.00606	241	0.00417					

BY2014 Fry



Logistic Regression Model Minijack:

Pr(Mature as MJ) ~ JuvMale FL* + JuvMale BdWt*

* These are phenotypic data from individual juvenile males. Once parentage analysis is completed we can identify all the Adult related traits for that individual male.

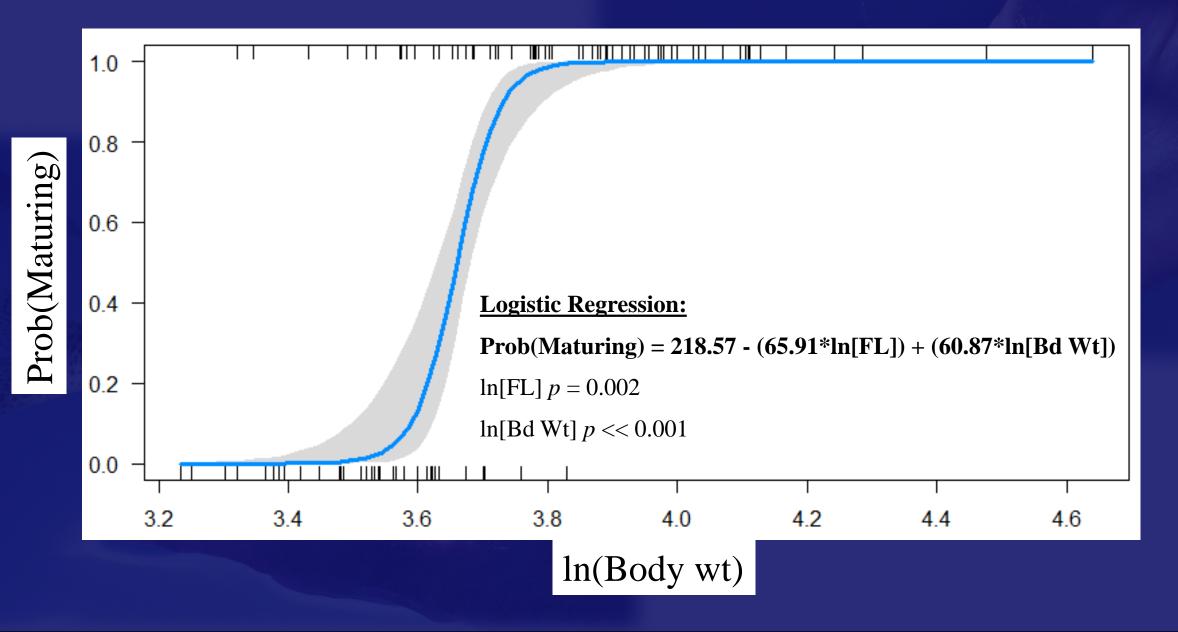
Full Juvenile Model: Prob(Mature as MiniJk) ~ ln(Male FL) + ln(Male Bd Wt)

<u>BY2014</u> Logistic Regression Model selection table:

Intercept	lnBW	lnFL df	logLik	AICc	delta	weight
218.6000	30.870	-65.91 3	-34.831	75.9	0.00	0.998
-34.7300	9.644	2	-42.052	88.2	12.32	0.002
-106.9000		21.40 2	-51.364	106.8	30.95	0.000
0.5671		1	-68.723	139.5	63.59	0.000

Output from *dredge* option in *MuMin* package (R software)

BY14 Minijack April Samples

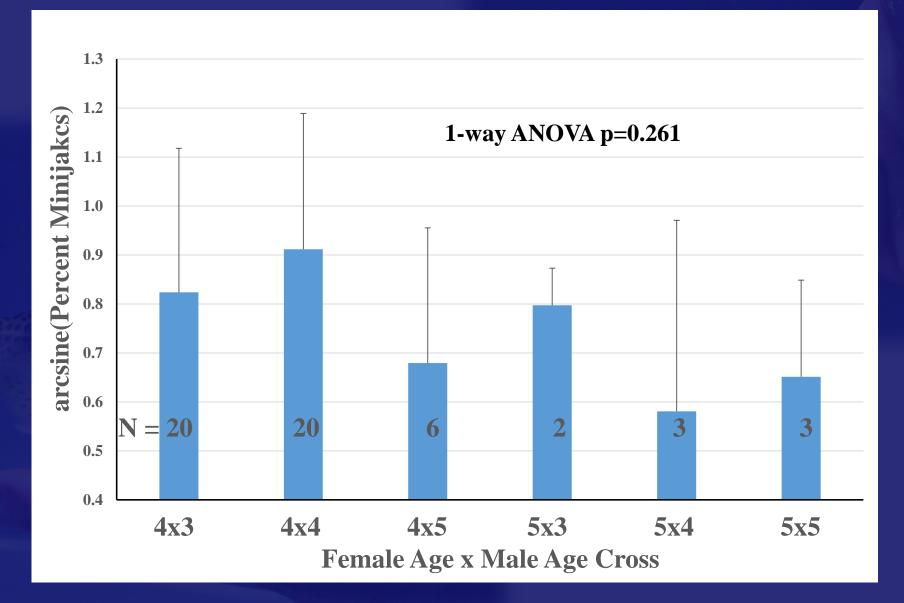


Logistic Regression Model:

Pr(Mature as MJ) ~ JuvMale FL + JuvMale BdWt +

Next StepIncludeAdultSire age + Dam age + Sire FL+ Sire BdWt + Dam FL + Dam BdWt +
Dam Egg wt + Fry Condition FactorTraits

BY2014 arcsin(Percent Minijacks) + 1 sd



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CRITFC personnel:

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