

	MutSpec vs Temperature (FishBase)	MutSpec vs Maturation (FishBase)		MutSpec vs MaximalLifespan (AnAge)	MutSpec vs Temperature + Maturation (Fish Base)	
		Lm	Tm		Lm	Tm
Co re lat io n	<p>Spearman's rank correlation rho:</p> <p>data: TemperMut\$A_G and TemperMut\$Temperature</p> <p>S = 474663, p-value = 3.321e-05 alternative hypothesis: true rho is not equal to 0 sample estimates: rho -0.3581037</p> <p>data: TemperMut\$T_C and TemperMut\$Temperature</p> <p>S = 256954, p-value = 0.002522 alternative hypothesis: true rho is not equal to 0 sample estimates: rho 0.2648037</p> <p>AFTER NORMALIZATION Spearman's rank correlation rho</p> <p>data: TemperMut\$T_C.NormalOnlyByT and TemperMut\$Temperature S = 325684, p-value = 0.4446 alternative hypothesis: true rho is not equal to 0 sample estimates: rho 0.06815333</p> <p>Spearman's rank correlation rho</p> <p>data: TemperMut\$A_G.NormalOnlyByA and TemperMut\$Temperature S = 385666, p-value = 0.146 alternative hypothesis: true rho is not equal to 0 sample estimates:</p>	<p>Spearman's rank correlation rho:</p> <p>data: MATULmmut\$G_C and MATULmmut\$Lm</p> <p>S = 192684, p-value = 0.008235 alternative hypothesis: true rho is not equal to 0 sample estimates: rho -0.266859</p>	<p>Spearman's rank correlation rho:</p> <p>data: MATUTmmut\$G_C and MATUTmmut\$Tm</p> <p>S = 242544, p-value = 0.0222 alternative hypothesis: true rho is not equal to 0 sample estimates: rho -0.2219755</p>	<p>Spearman's rank correlation rho</p> <p>data: AnAgeMut\$T_C and AnAgeMut\$Maximum.longevity..yr s.</p> <p>S = 223911, p-value = 0.9219 alternative hypothesis: true rho is not equal to 0 sample estimates: rho -0.009449996</p>	<p>Call: lm(formula = T_C ~ scale(Temperature) * scale(Lm), data = allparameters)</p> <p>Residuals: Min 1Q Median 3Q Max -0.150980 -0.040899 -0.009959 0.043756 0.150034</p> <p>Coefficients: value Pr(> t) (Intercept) 0.133998 0.008373 16.003 <2e-16 *** scale(Temperature) 0.019695 0.008442 2.333 0.023 * scale(Lm) -0.001413 0.009040 -0.156 0.876 scale(Temperature):scale(Lm) -0.003960 0.010934 -0.362 0.719 --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</p> <p>Residual standard error: 0.06685 on 60 degrees of freedom Multiple R-squared: 0.08655, Adjusted R-squared: 0.04088 F-statistic: 1.895 on 3 and 60 DF, p-value: 0.1401</p> <p>Call: lm(formula = T_C ~ scale(Temperature) + scale(Lm), data = allparameters)</p> <p>Residuals: Min 1Q Median 3Q Max -0.148531 -0.043140 -0.008988 0.044211 0.146448</p> <p>Coefficients: Estimate Std. Error t value Pr(> t) (Intercept) 0.134190 0.008297 16.174 <2e-16 ***</p>	<p>Call: lm(formula = T_C ~ scale(Temperature) * scale(Tm), data = allparameters)</p> <p>Residuals: Min 1Q Median 3Q Max -0.115771 -0.040550 -0.007716 0.039935 0.136724</p> <p>Coefficients: value Pr(> t) (Intercept) 0.131507 0.008055 16.325 <2e-16 *** scale(Temperature) 0.024866 0.008083 3.076 0.00313 ** scale(Tm) -0.008903 0.008722 -1.021 0.31142 scale(Temperature):scale(Tm) -0.000811 0.009005 -0.090 0.92854 --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</p> <p>Residual standard error: 0.062 on 61 degrees of freedom Multiple R-squared: 0.1815, Adjusted R-squared: 0.1412 F-statistic: 4.508 on 3 and 61 DF, p-value: 0.006387</p> <p>Call: lm(formula = T_C ~ scale(Temperature) + scale(Tm), data = allparameters)</p> <p>Residuals: Min 1Q Median 3Q Max -0.11592 -0.04111 -0.00788 0.03984 0.13706</p> <p>Coefficients: Estimate Std. Error t value Pr(> t) (Intercept) 0.131723 0.007628 17.269 <2e-16 *** scale(Temperature) 0.024931 0.007985 3.122 0.00273 **</p>

<div><div>rho</div><div>-0.1297411</div><div></div><div>WHAT WITH A_G?</div><div></div><div>Call: lm(formula = A_G ~ scale(Temperature) * scale(Tm), data = allparameters)</div><div>Residuals: Min 1Q Median 3Q Max -0.081267 -0.026615 -0.007707 0.014225 0.135117</div><div>Coefficients: Estimate Std. Error t value Pr(> t) (Intercept) 0.067337 0.006327 10.643 1.56e-15 *** scale(Temperature) -0.014702 0.006348 -2.316 0.0239 * scale(Tm) -0.009556 0.006851 -1.395 0.1681 scale(Temperature):scale(Tm) -0.005560 0.007073 -0.786 0.4349 --- Signif. codes: 0 '***' 0.001 '*' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div><div>Residual standard error: 0.04869 on 61 degrees of freedom Multiple R-squared: 0.0903, Adjusted R-squared: 0.04556 F-statistic: 2.018 on 3 and 61 DF, p-value: 0.1207</div><div>MOREOVER: Call: lm(formula = Temperature ~ scale(T_C) * scale(A_G), data = allparameters)</div><div>Residuals: Min 1Q Median 3Q Max</div></div>				<div><div>scale(Temperature) 0.019552 0.008373 2.335 0.0228 * scale(Lm) -0.002592 0.008373 -0.310 0.7579 --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div><div>Residual standard error: 0.06637 on 61 degrees of freedom Multiple R-squared: 0.08455, Adjusted R-squared: 0.05454 F-statistic: 2.817 on 2 and 61 DF, p-value: 0.06758</div><div>Call: lm(formula = scale(T_C) ~ scale(Temperature) + scale(Lm), data = allparameters)</div><div>Residuals: Min 1Q Median 3Q Max -2.1759 -0.6320 -0.1317 0.6477 2.1454</div><div>Coefficients: Estimate Std. Error t value Pr(> t) (Intercept) 2.199e-16 1.215e-01 0.000 1.0000 scale(Temperature) 2.864e-01 1.227e-01 2.335 0.0228 * scale(Lm) -3.798e-02 1.227e-01 -0.310 0.7579 --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div><div>Residual standard error: 0.9723 on 61 degrees of freedom Multiple R-squared: 0.08455, Adjusted R-squared: 0.05454 F-statistic: 2.817 on 2 and 61 DF, p-value: 0.06758</div><div>Call: lm(formula = scale(T_C) ~ 0 + scale(Temperature) + scale(Lm), data = allparameters)</div><div>Residuals: Min 1Q Median 3Q Max -2.1759 -0.6320 -0.1317 0.6477 2.1454</div></div>	<div><div>scale(Tm) -0.008600 0.007985 -1.077 0.28564 --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div><div>Residual standard error: 0.0615 on 62 degrees of freedom Multiple R-squared: 0.1814, Adjusted R-squared: 0.1549 F-statistic: 6.867 on 2 and 62 DF, p-value: 0.002023</div><div>Call: lm(formula = scale(T_C) ~ scale(Temperature) + scale(Tm), data = allparameters)</div><div>Residuals: Min 1Q Median 3Q Max -1.7327 -0.6145 -0.1178 0.5956 2.0488</div><div>Coefficients: Estimate Std. Error t value Pr(> t) (Intercept) -4.629e-16 1.140e-01 0.000 1.00000 scale(Temperature) 3.727e-01 1.194e-01 3.122 0.00273 ** scale(Tm) -1.286e-01 1.194e-01 -1.077 0.28564 --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div><div>Residual standard error: 0.9193 on 62 degrees of freedom Multiple R-squared: 0.1814, Adjusted R-squared: 0.1549 F-statistic: 6.867 on 2 and 62 DF, p-value: 0.002023</div><div>Call: lm(formula = scale(T_C) ~ 0 + scale(Temperature) + scale(Tm), data = allparameters)</div><div>Residuals: Min 1Q Median 3Q Max -1.7327 -0.6145 -0.1178 0.5956 2.0488</div><div>Coefficients:</div></div>
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<div><div><div><div><div>-19.1572</div><div>-4.7552</div><div>0.7703</div></div><div>4.6829</div><div>11.5930</div></div></div><div><div>Coefficients:</div><div><div><div>Estimate</div><div>Std.</div><div>Error</div><div>t value</div><div>Pr(> t)</div></div><div>(Intercept)</div><div>16.0049</div><div>0.8609</div><div>18.590</div><div>< 2e-16</div><div>***</div></div><div>scale(T_C)</div><div>3.0589</div><div>0.8804</div><div>3.475</div><div>0.000948</div><div>***</div></div><div>scale(A_G)</div><div>-1.1890</div><div>0.9333</div><div>-1.274</div><div>0.207495</div></div> <div>scale(T_C):scale(A_G)</div> <div>0.8792</div> <div>0.9322</div> <div>0.943</div> <div>0.349346</div> <div>---</div> <div>Signif. codes: 0 '***' 0.001</div> <div>'**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div> <div><div>Residual standard error: 6.885</div><div>on 61 degrees of freedom</div><div>Multiple R-squared: 0.2167,</div><div>Adjusted R-squared:</div><div>0.1782</div><div>F-statistic: 5.625 on 3 and 61</div><div>DF, p-value: 0.001802</div></div> <div><div>Call:</div><div>lm(formula = Temperature ~</div><div>scale(T_C) + scale(A_G), data</div><div>= allparameters)</div></div> <div><div>Residuals:</div><div><div>Min</div><div>1Q</div><div>Median</div><div>3Q</div><div>Max</div></div><div>-19.3869</div><div>-4.7812</div><div>0.4572</div><div>4.5021</div><div>11.5473</div></div> <div><div>Coefficients:</div><div><div><div>Estimate</div><div>Std.</div><div>Error</div><div>t</div><div>value</div><div>Pr(> t)</div></div><div>(Intercept)</div><div>15.9015</div><div>0.8532</div><div>18.638</div><div>< 2e-16</div><div>***</div></div><div>scale(T_C)</div><div>2.9136</div><div>0.8660</div><div>3.364</div><div>0.00132</div><div>**</div></div> <div>scale(A_G)</div> <div>-1.5153</div> <div>0.8660</div> <div>-1.750</div> <div>0.08511</div> <div>.</div> <div>---</div> <div>Signif. codes: 0 '***' 0.001</div> <div>'**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div> <div><div>Residual standard error: 6.879</div><div>on 62 degrees of freedom</div></div> <div></div> <div></div> <div></div> <div><div><div>Coefficients:</div><div><div><div>Estimate</div><div>Std.</div><div>Error</div><div>t value</div><div>Pr(> t)</div></div><div>scale(Temperature)</div><div>0.28642</div><div>0.12166</div><div>2.354</div><div>0.0217</div><div>*</div></div><div>scale(Lm)</div><div>-0.03798</div><div>0.12166</div><div>-0.312</div><div>0.7560</div><div>---</div><div>Signif. codes: 0 '***' 0.001</div><div>'**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div></div><div><div>Residual standard error: 0.9645 on 62</div><div>degrees of freedom</div><div>Multiple R-squared: 0.08455,</div><div>Adjusted R-squared: 0.05502</div><div>F-statistic: 2.863 on 2 and 62 DF,</div><div>p-value: 0.06466</div></div><div><div>Call:</div><div>lm(formula = scale(T_C) ~ 0 +</div><div>scale(Temperature), data =</div><div>allparameters)</div></div><div><div>Residuals:</div><div><div>Min</div><div>1Q</div><div>Median</div><div>3Q</div><div>Max</div></div><div>-2.1557</div><div>-0.6148</div><div>-0.1217</div><div>0.6598</div><div>2.1147</div></div><div><div>Coefficients:</div><div><div><div>Estimate</div><div>Std.</div><div>Error</div><div>t value</div><div>Pr(> t)</div></div><div>scale(Temperature)</div><div>0.2883</div><div>0.1206</div><div>2.39</div><div>0.0199</div><div>*</div></div><div>---</div><div>Signif. codes: 0 '***' 0.001</div><div>'**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div></div><div><div>Residual standard error: 0.9575 on 63</div><div>degrees of freedom</div><div>Multiple R-squared: 0.08311,</div><div>Adjusted R-squared: 0.06856</div><div>F-statistic: 5.711 on 1 and 63 DF,</div><div>p-value: 0.01986</div></div></div> <div><div><div>Estimate</div><div>Std.</div><div>Error</div><div>t value</div><div>Pr(> t)</div></div><div>scale(Temperature)</div><div>0.3727</div><div>0.1184</div><div>3.147</div><div>0.00252</div><div>**</div></div> <div>scale(Tm)</div> <div>-0.1286</div> <div>0.1184</div> <div>-1.086</div> <div>0.28176</div> <div>---</div> <div>Signif. codes: 0 '***' 0.001</div> <div>'**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div> <div><div>Residual standard error: 0.9119 on 63</div><div>degrees of freedom</div><div>Multiple R-squared: 0.1814,</div><div>Adjusted R-squared: 0.1554</div><div>F-statistic: 6.978 on 2 and 63 DF,</div><div>p-value: 0.00183</div></div> <div><div>Call:</div><div>lm(formula = scale(T_C) ~ 0 +</div><div>scale(Temperature), data =</div><div>allparameters)</div></div> <div><div>Residuals:</div><div><div>Min</div><div>1Q</div><div>Median</div><div>3Q</div><div>Max</div></div><div>-1.67386</div><div>-0.63249</div><div>-0.04528</div><div>0.58369</div><div>2.15754</div></div> <div><div>Coefficients:</div><div><div><div>Estimate</div><div>Std.</div><div>Error</div><div>t value</div><div>Pr(> t)</div></div><div>scale(Temperature)</div><div>0.4075</div><div>0.1142</div><div>3.57</div><div>0.000684</div><div>***</div></div><div>---</div><div>Signif. codes: 0 '***' 0.001</div><div>'**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</div></div> <div><div>Residual standard error: 0.9132 on 64</div><div>degrees of freedom</div><div>Multiple R-squared: 0.166,</div><div>Adjusted R-squared: 0.153</div><div>F-statistic: 12.74 on 1 and 64 DF,</div><div>p-value: 0.0006844</div></div>
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	Multiple R-squared: 0.2053, Adjusted R-squared: 0.1796 F-statistic: 8.007 on 2 and 62 DF, p-value: 0.0008065					
N of species	128	97	106	110	64	65
Script	https://github.com/polarsong/mtDNA_mutspectrum/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.ALL_RANK_CORR.R	https://github.com/polarsong/mtDNA_mutspectrum/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.ALL_RANK_CORR.R	https://github.com/polarsong/mtDNA_mutspectrum/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.ALL_RANK_CORR.R	https://github.com/polarsong/mtDNA_mutspectrum/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.ALL_RANK_CORR.R	https://github.com/polarsong/mtDNA_mutspectrum/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.MultipleReg_Maturity~Temp.R	https://github.com/polarsong/mtDNA_mutspectrum/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.MultipleReg_Maturity~Temp.R
Figures						

	WholeGenome vs Temperature (FishBase)	WholeGenome vs Maturation (FishBase)	WholeGenome vs MaximalLifespan (AnAge)	WholeGenome vs Maturation (AnAge)	WholeGenome vs Body Mass (AnAge)	WholeGenome vs Temperature(FishBase) + Maturation(FishBase)
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		Lm	Tm				Lm	Tm
C or re la ti o n	Pearson's product-moment correlation data: log2(AGG\$FemaleMaturityDays) and AGG\$FrT t = -2.7871, df = 300, p-value = 0.005657 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: -0.26694950 -0.04684676 sample estimates: cor -0.1588715 data: log2(AGG\$FemaleMaturityDays) and AGG\$FrG t = -3.6769, df = 300, p-value = 0.0002796 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: -0.31318440 -0.09707033 sample estimates: cor -0.2076599	Pearson's product-moment correlation data: log2(AGG\$FemaleMaturityDays) and AGG\$FrA t = 2.5363, df = 190, p-value = 0.01201 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: 0.04038784 0.31450996 sample estimates: cor 0.1809612 data: log2(AGG\$FemaleMaturityDays) and AGG\$FrT t = -4.7702, df = 190, p-value = 3.657e-06 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: -0.4479050 -0.1944362 sample estimates: cor -0.3270399	Pearson's product-moment correlation data: log2(AGG\$FemaleMaturityDays) and AGG\$FrT t = -4.0306, df = 186, p-value = 8.105e-05 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: -0.4099044 -0.1462413 sample estimates: cor -0.28342 data: log2(AGG\$FemaleMaturityDays) and AGG\$FrT t = -7.3526, df = 204, p-value = 4.624e-12 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: -0.5593992 -0.3424218 sample estimates: cor -0.4576984 data: log2(AGG\$FemaleMaturityDays) and AGG\$FrC t = 3.0345, df = 204, p-value = 0.002723 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: 0.07319492 0.33500116 sample estimates: cor 0.2078168	Pearson's product-moment correlation data: log2(AGG\$FemaleMaturityDays) and AGG\$FrA t = 2.9679, df = 204, p-value = 0.003357 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: 0.06865682 0.33094580 sample estimates: cor 0.2034487 data: log2(AGG\$FemaleMaturityDays) and AGG\$FrT t = -7.3526, df = 204, p-value = 4.624e-12 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: -0.5593992 -0.3424218 sample estimates: cor -0.4576984 data: log2(AGG\$FemaleMaturityDays) and AGG\$FrC t = 3.0345, df = 204, p-value = 0.002723 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: 0.07319492 0.33500116 sample estimates: cor 0.2078168	Pearson's product-moment correlation data: log2(AGG\$FemaleMaturityDays) and AGG\$FrT t = -5.0863, df = 89, p-value = 2.01e-06 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: -0.6199221 -0.2977171 sample estimates: cor -0.4745648 data: log2(AGG\$FemaleMaturityDays) and AGG\$FrC t = 2.4411, df = 89, p-value = 0.01662 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: 0.04698102 0.43405429 sample estimates: cor 0.2505021	Pearson's product-moment correlation data: log2(AGG\$FemaleMaturityDays) and AGG\$FrT t = -3.7031, df = 124, p-value = 0.0003191 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: -0.4648112 -0.1488685 sample estimates: cor -0.3155589 data: log2(AGG\$FemaleMaturityDays) and AGG\$FrC t = 2.7439, df = 124, p-value = 0.006972 alternative hypothesis: true correlation is not equal to 0 95 percent confidence interval: 0.0671575 0.3975257 sample estimates: cor 0.2392541	Call: lm(formula = FrT ~ scale(Temperature) * scale(Maturity), data = AGGTEMPE) Residuals: Min 1Q Median 3Q Max -0.125078 -0.022805 0.002719 0.019496 0.115140 Coefficients: Estimate Std. Error t value Pr(> t) (Intercept) 0.215970 0.003838 56.273 < 2e-16 *** scale(Temperature) -0.015026 0.003812 -3.942 0.000138 *** scale(Maturity) -0.016741 0.003966 -4.221 4.85e-05 *** scale(Temperature):scale(Maturity) -0.001854 0.004659 -0.398 0.691467 --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 Residual standard error: 0.03978 on 116 degrees of freedom Multiple R-squared: 0.1882, Adjusted R-squared: 0.1672 F-statistic: 8.962 on 3 and 116 DF, p-value: 2.179e-05 Call: lm(formula = FrT ~ scale(Temperature) +	Call: lm(formula = FrT ~ scale(Temperature) * scale(Maturity), data = AGGTEMPE) Residuals: Min 1Q Median 3Q Max -0.088755 -0.024531 -0.000606 0.020775 0.139422 Coefficients: Estimate Std. Error t value Pr(> t) (Intercept) 0.219415 0.004024 54.520 < 2e-16 *** scale(Temperature) -0.021204 0.003880 -5.464 3.10e-07 *** scale(Maturity) -0.023345 0.005685 -4.106 7.93e-05 *** scale(Temperature):scale(Maturity) -0.008171 0.005267 -1.551 0.124 --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 Residual standard error: 0.03812 on 106 degrees of freedom Multiple R-squared: 0.2674, Adjusted R-squared: 0.2467 F-statistic: 12.9 on 3 and 106 DF, p-value: 3.014e-07 Call: lm(formula = FrT ~ scale(Temperature) +

						<p>scale(Maturity), data = AGGTEMPE)</p> <p>Residuals:</p> <table><tr><td>Min</td><td>1Q</td></tr><tr><td>Median</td><td>3Q</td></tr><tr><td>Max</td><td></td></tr><tr><td>-0.124858</td><td>-0.023289</td></tr><tr><td>0.002424</td><td>0.019246</td></tr><tr><td>0.114993</td><td></td></tr></table> <p>Coefficients:</p> <table><tr><td>Estimate</td><td></td></tr><tr><td>Std. Error</td><td>t value</td></tr><tr><td>Pr(> t)</td><td></td></tr><tr><td>(Intercept)</td><td></td></tr><tr><td>0.216464</td><td>0.003619</td></tr><tr><td>59.819</td><td>< 2e-16 ***</td></tr><tr><td>scale(Temperature)</td><td></td></tr><tr><td>-0.014851</td><td>0.003772</td></tr><tr><td>-3.937</td><td>0.000141 ***</td></tr><tr><td>scale(Maturity)</td><td></td></tr><tr><td>-0.016271</td><td>0.003772</td></tr><tr><td>-4.313</td><td>3.38e-05 ***</td></tr><tr><td>---</td><td></td></tr><tr><td>Signif. codes: 0 ‘***’</td><td></td></tr><tr><td>0.001 ‘**’</td><td>0.01 ‘*’</td></tr><tr><td>0.05 ‘.’</td><td>0.1 ‘ ’ 1</td></tr></table> <p>Residual standard error: 0.03964 on 117 degrees of freedom Multiple R-squared: 0.1871, Adjusted R-squared: 0.1732 F-statistic: 13.46 on 2 and 117 DF, p-value: 5.475e-06</p> <p>Call: lm(formula = scale(FrT) ~ scale(Temperature) + scale(Maturity), data = AGGTEMPE)</p> <p>Residuals:</p> <table><tr><td>Min</td><td>1Q</td></tr><tr><td>Median</td><td>3Q</td></tr><tr><td>Max</td><td></td></tr></table>	Min	1Q	Median	3Q	Max		-0.124858	-0.023289	0.002424	0.019246	0.114993		Estimate		Std. Error	t value	Pr(> t)		(Intercept)		0.216464	0.003619	59.819	< 2e-16 ***	scale(Temperature)		-0.014851	0.003772	-3.937	0.000141 ***	scale(Maturity)		-0.016271	0.003772	-4.313	3.38e-05 ***	---		Signif. codes: 0 ‘***’		0.001 ‘**’	0.01 ‘*’	0.05 ‘.’	0.1 ‘ ’ 1	Min	1Q	Median	3Q	Max		<p>scale(Maturity), data = AGGTEMPE)</p> <p>Residuals:</p> <table><tr><td>Min</td><td>1Q</td></tr><tr><td>Median</td><td>3Q</td></tr><tr><td>Max</td><td></td></tr><tr><td>-0.087127</td><td>-0.023234</td></tr><tr><td>0.000743</td><td>0.018729</td></tr><tr><td>0.141510</td><td></td></tr></table> <p>Coefficients:</p> <table><tr><td>Estimate</td><td></td></tr><tr><td>Std. Error</td><td>t value</td></tr><tr><td>Pr(> t)</td><td></td></tr><tr><td>(Intercept)</td><td></td></tr><tr><td>0.222096</td><td>0.003659</td></tr><tr><td>60.706</td><td>< 2e-16 ***</td></tr><tr><td>scale(Temperature)</td><td></td></tr><tr><td>-0.020756</td><td>0.003895</td></tr><tr><td>-5.329</td><td>5.53e-07 ***</td></tr><tr><td>scale(Maturity)</td><td></td></tr><tr><td>-0.016883</td><td>0.003895</td></tr><tr><td>-4.335</td><td>3.31e-05 ***</td></tr><tr><td>---</td><td></td></tr><tr><td>Signif. codes: 0 ‘***’</td><td></td></tr><tr><td>0.001 ‘**’</td><td>0.01 ‘*’</td></tr><tr><td>0.05 ‘.’</td><td>0.1 ‘ ’ 1</td></tr></table> <p>Residual standard error: 0.03837 on 107 degrees of freedom Multiple R-squared: 0.2508, Adjusted R-squared: 0.2368 F-statistic: 17.91 on 2 and 107 DF, p-value: 1.956e-07</p> <p>Call: lm(formula = scale(FrT) ~ scale(Temperature) + scale(Maturity), data = AGGTEMPE)</p> <p>Residuals:</p> <table><tr><td>Min</td><td>1Q</td><td>Median</td></tr><tr><td>3Q</td><td>Max</td><td></td></tr><tr><td>-1.9837</td><td>-0.5290</td><td></td></tr><tr><td>0.0169</td><td>0.4264</td><td></td></tr><tr><td>3.2219</td><td></td><td></td></tr></table>	Min	1Q	Median	3Q	Max		-0.087127	-0.023234	0.000743	0.018729	0.141510		Estimate		Std. Error	t value	Pr(> t)		(Intercept)		0.222096	0.003659	60.706	< 2e-16 ***	scale(Temperature)		-0.020756	0.003895	-5.329	5.53e-07 ***	scale(Maturity)		-0.016883	0.003895	-4.335	3.31e-05 ***	---		Signif. codes: 0 ‘***’		0.001 ‘**’	0.01 ‘*’	0.05 ‘.’	0.1 ‘ ’ 1	Min	1Q	Median	3Q	Max		-1.9837	-0.5290		0.0169	0.4264		3.2219		
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						<p>scale(Temperature) -0.34067 0.08617 -3.954 0.000132 *** scale(Maturity) -0.37325 0.08617 -4.332 3.13e-05 *** --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</p> <p>Residual standard error: 0.9054 on 118 degrees of freedom Multiple R-squared: 0.1871, Adjusted R-squared: 0.1733 F-statistic: 13.58 on 2 and 118 DF, p-value: 4.936e-06</p> <p>Call: lm(formula = scale(FrT) ~ 0 + scale(Temperature), data = AGGTEMPE)</p> <p>Residuals: Min 1Q Median 3Q Max -2.5877 -0.4248 0.0315 0.5867 2.7099</p> <p>Coefficients: Estimate Std. Error t value Pr(> t) scale(Temperature) -0.24042 0.08898 -2.702 0.0079 ** --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</p> <p>Residual standard error: 0.9707 on 119 degrees of freedom Multiple R-squared: 0.0578, Adjusted R-squared: 0.04988</p>	<p>scale(Maturity) -0.38439 0.08827 -4.355 3.04e-05 *** --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</p> <p>Residual standard error: 0.8696 on 108 degrees of freedom Multiple R-squared: 0.2508, Adjusted R-squared: 0.2369 F-statistic: 18.08 on 2 and 108 DF, p-value: 1.693e-07</p> <p>Call: lm(formula = scale(FrT) ~ 0 + scale(Temperature), data = AGGTEMPE)</p> <p>Residuals: Min 1Q Median 3Q Max -2.4959 -0.4454 -0.0535 0.4359 3.4630</p> <p>Coefficients: Estimate Std. Error t value Pr(> t) scale(Temperature) -0.34529 0.08989 -3.841 0.000206 *** --- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1</p> <p>Residual standard error: 0.9385 on 109 degrees of freedom Multiple R-squared: 0.1192, Adjusted R-squared: 0.1111 F-statistic: 14.75 on 1 and 109 DF, p-value: 0.000206</p>
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							F-statistic: 7.3 on 1 and 119 DF, p-value: 0.007903	
N of species	302	192	188	206	91	126	120	110
Script	https://github.com/polarson/mtDNA_mutspectrum/blob/WholeGenomesBranch/Head/2Scripts/WholeGenomeAnalyses.EcologyAndMutSpecChordata.NoOverlap.ActinopterygiiOnlyFishBase.R	https://github.com/polarson/mtDNA_mutspectrum/blob/WholeGenomesBranch/Head/2Scripts/WholeGenomeAnalyses.EcologyAndMutSpecChordata.NoOverlap.ActinopterygiiOnlyFishBase.R	https://github.com/polarson/mtDNA_mutspectrum/blob/WholeGenomesBranch/Head/2Scripts/WholeGenomeAnalyses.EcologyAndMutSpecChordata.NoOverlap.ActinopterygiiOnlyFishBase.R	https://github.com/polarson/mtDNA_mutspectrum/blob/WholeGenomesBranch/Head/2Scripts/WholeGenomeAnalyses.EcologyAndMutSpecChordata.NoOverlap.ActinopterygiiOnlyFishBase.R	https://github.com/polarson/mtDNA_mutspectrum/blob/WholeGenomesBranch/Head/2Scripts/WholeGenomeAnalyses.EcologyAndMutSpecChordata.NoOverlap.ActinopterygiiOnlyFishBase.R	https://github.com/polarson/mtDNA_mutspectrum/blob/WholeGenomesBranch/Head/2Scripts/WholeGenomeAnalyses.EcologyAndMutSpecChordata.NoOverlap.ActinopterygiiOnlyFishBase.R	https://github.com/polarson/mtDNA_mutspectrum/blob/WholeGenomesBranch/Head/2Scripts/WholeGenomeAnalyses.EcologyAndMutSpecChordata.NoOverlap.ActinopterygiiOnlyFishBase.R	https://github.com/polarson/mtDNA_mutspectrum/blob/WholeGenomesBranch/Head/2Scripts/WholeGenomeAnalyses.EcologyAndMutSpecChordata.NoOverlap.ActinopterygiiOnlyFishBase.R
Figures								

	Maturation Lm vs TM (FishBase)	Temperature vs Maturation Lm (FishBase)	Temperature vs Maturation Tm (FishBase)	Temperature (FishBase) vs MaximalLifespan (AnAge)	Maturation (FishBase) vs MaximalLifespan (AnAge)
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C o r r e l a t i o n	<p>Spearman's rank correlation rho</p> <p>data: MATULMTM\$Lm and MATULMTM\$Tm S = 3838500, p-value < 2.2e-16 alternative hypothesis: true rho is not equal to 0 sample estimates: rho 0.6823805</p>	<p>Spearman's rank correlation rho</p> <p>data: TEMPMATULM\$Lm and TEMPMATULM\$Temperature S = 29562841, p-value = 1.282e-12 alternative hypothesis: true rho is not equal to 0 sample estimates: rho -0.3062001</p>	<p>Spearman's rank correlation rho</p> <p>data: TEMPMATUTM\$Tm and TEMPMATUTM\$Temperature S = 21243590, p-value = 1.141e-11 alternative hypothesis: true rho is not equal to 0 sample estimates: rho -0.3095067</p>	<p>Spearman's rank correlation rho</p> <p>data: ANAGETEMP\$Temperature and ANAGETEMP\$Maximum.longevity..yrs. S = 10501627, p-value = 3.03e-05 alternative hypothesis: true rho is not equal to 0 sample estimates: rho -0.2141839</p>	<p>Lm Spearman's rank correlation rho</p> <p>data: ANAGEMATLM\$Lm and ANAGEMATLM\$Maximum.longevity..yrs. S = 1020722, p-value < 2.2e-16 alternative hypothesis: true rho is not equal to 0 sample estimates: rho 0.5399325</p> <p>Tm Spearman's rank correlation rho</p> <p>data: ANAGEMATTM\$Tm and ANAGEMATTM\$Maximum.longevity..yrs. S = 969449, p-value < 2.2e-16 alternative hypothesis: true rho is not equal to 0 sample estimates: rho 0.7321639</p>
N o f s p e c i e s	417	514	460	376	<p>Lm 238</p> <p>Tm 281</p>
S c r i p t	https://github.com/polarson/mtDNA_mutspec/mtDNA_mutspec/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.ALL_RANK_CORR.R	https://github.com/polarson/mtDNA_mutspec/mtDNA_mutspec/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.ALL_RANK_CORR.R	https://github.com/polarson/mtDNA_mutspec/mtDNA_mutspec/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.ALL_RANK_CORR.R	https://github.com/polarson/mtDNA_mutspec/mtDNA_mutspec/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.ALL_RANK_CORR.R	https://github.com/polarson/mtDNA_mutspec/mtDNA_mutspec/blob/Actinopterygii/Head/2Scripts/VertebratePolymorphisms.MutSpecComparisons.Analyses.Ecology.Actinopterygii.FishBaseData.ALL_RANK_CORR.R
F i g u r e s					