

Supplemental Table 3: Comparisons of GR mRNA levels for each tissue between midshipman reproductive morphs

Morph	Breeding Season Tissue	Norm. mean	Standard error	ANOVA values
Male Type I	OB/Tel/POA	0.0152* ¹	0.0015	p=0.0001
Male Type II		0.0353* ¹	0.0102	dF=2, 11
Female		0.0071	0.0002	Fratio=22.60
Male Type I	Mid/Di/Cbl	0.0131	0.0024	p=0.0020
Male Type II		0.0266* ²	0.0063	dF=2, 11
Female		0.0065	0.0007	Fratio=11.55
Male Type I	Vocal Hindbrain – SC	0.007	0.0006	p<0.0001
Male Type II		0.042* ³	0.0091	dF=2, 11
Female		0.0051	0.0005	Fratio=67.79
Male Type I	Vocal Muscle	0.0275* ⁴	0.0062	p=0.0137
Male Type II		0.0113	0.002	dF=2, 11
Female		0.0133	0.0013	Fratio=6.49
Male Type I	Liver	0.0398* ⁴	0.0081	p=0.0011
Male Type II		0.0163	0.0026	dF=2, 10
Female		0.0129	0.0013	Fratio=14.62
Male Type I	Testis	0.0191	0.0056	p=0.6453
Male Type II		0.0215	0.006	dF=1, 8
¹ II>I>F, ² II>F, ³ II>I and F, ⁴ I>II and F				Fratio=0.2287

Supplemental Table 3: Glucocorticoid receptor (GR) mRNA levels for tissues sampled in the three midshipman reproductive morphs. Mean normalized values (see Fig. 3 legend) and standard errors for mRNA levels are listed according to midshipman reproductive morph (column 1) for each tissue sampled (column 2, see Fig. 3 for abbreviations). ANOVA values (p value, degrees of freedom-dF, and Fratio) are also listed for each analysis performed for each tissue. Asterisks indicate significantly greater values over one or both other morphs for each tissue (see footnotes).