Supplemental Table 4: Comparisons of MR mRNA levels for each tissue between midshipman reproductive morphs

Morph	Breeding Season Tissue	Norm. mean	Standard error	ANOVA values
Male Type I		0.001	0.0002	p=0.0331
Male Type II	**OB/Tel/POA**	0.027*[1]	0.022	dF=2, 10
Female		0.0014	0.0004	Fratio=4.89
Male Type I		0.0049	0.0035	p=0.4511
Male Type II	**Mid/Di/Cbl**	0.0284	0.0189	dF=2, 11
Female		0.0131	0.0123	Fratio=0.8565
Male Type I		0.0004	0.0001	p=0.0021
Male Type II	**Vocal Hindbrain – SC**	0.0086*[2]	0.0037	dF=2, 10
Female		0.0004	0.0001	Fratio=12.12
Male Type I		0.0001	0.00002	p=0.0023
Male Type II	**Vocal Muscle**	0.0022*[2]	0.0003	dF=2, 11
Female		0.0007	0.0005	Fratio=11.10
Male Type I		0.0013	0.0012	p=0.1742
Male Type II	**Liver**	0.0024	0.001	dF=2, 12
Female		0.0144	0.014	Fratio=2.03
Male Type I	**Testis**	0.0078	0.0077	p=0.0623
Male Type II		0.0071	0.0014	dF=1, 7
[1]II>I, [2]II>F and I				Fratio=4.91

Supplemental Table 4: Mineralocorticoid receptor (MR) mRNA levels for tissues sampled in the three midshipman reproductive morphs. Mean normalized values (see Fig. 3 legend) and standard errors are listed for mRNA levels according to midshipman reproductive morph (column 1) and 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).