Tables\_Wild\_2024

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## Tables

**Table S1**. Basal treatment ANOVA model output.

| term | sumsq | df | statistic | p.value |
| --- | --- | --- | --- | --- |
| temp | 0.002 | 1 | 0.10 | 0.752 |
| hormone | 0.088 | 2 | 1.92 | 0.154 |
| scale(Juvenile3\_Age) | 0.017 | 1 | 0.75 | 0.390 |
| **sex** | **0.363** | **1** | **15.78** | **0.000** |
| **juv3\_mass\_g** | **0.114** | **1** | **4.94** | **0.029** |
| juv3\_oroboros | 0.012 | 1 | 0.52 | 0.472 |
| juv3\_oroboros:chamber | 0.006 | 1 | 0.27 | 0.604 |
| Residuals | 1.726 | 75 |  |  |

**Table S2**. ADP treatment ANOVA model output.

| term | sumsq | df | statistic | p.value |
| --- | --- | --- | --- | --- |
| temp | 1.77 | 1 | 1.39 | 0.242 |
| hormone | 5.24 | 2 | 2.06 | 0.135 |
| **sex** | **9.93** | **1** | **7.82** | **0.007** |
| scale(Juvenile3\_Age) | 0.28 | 1 | 0.22 | 0.641 |
| **juv3\_mass\_g** | **5.77** | **1** | **4.54** | **0.036** |
| juv3\_oroboros | 0.00 | 1 | 0.00 | 0.995 |
| juv3\_oroboros:chamber | 0.16 | 1 | 0.13 | 0.722 |
| Residuals | 95.32 | 75 |  |  |

**Table S3**. Oligo treatment ANOVA model output.

| term | sumsq | df | statistic | p.value |
| --- | --- | --- | --- | --- |
| temp | 0.011 | 1 | 0.64 | 0.425 |
| hormone | 0.044 | 2 | 1.31 | 0.277 |
| **sex** | **0.214** | **1** | **12.70** | **0.001** |
| scale(Juvenile3\_Age) | 0.019 | 1 | 1.12 | 0.293 |
| juv3\_mass\_g | 0.064 | 1 | 3.82 | 0.054 |
| juv3\_oroboros | 0.002 | 1 | 0.14 | 0.711 |
| juv3\_oroboros:chamber | 0.003 | 1 | 0.20 | 0.654 |
| Residuals | 1.263 | 75 |  |  |

**Table S4**. RCR treatment ANOVA model output.

| term | sumsq | df | statistic | p.value |
| --- | --- | --- | --- | --- |
| temp | 26.72 | 1 | 3.140 | 0.080 |
| hormone | 14.28 | 2 | 0.839 | 0.436 |
| sex | 6.22 | 1 | 0.731 | 0.395 |
| scale(Juvenile3\_Age) | 26.23 | 1 | 3.082 | 0.083 |
| juv3\_mass\_g | 2.92 | 1 | 0.343 | 0.560 |
| juv3\_oroboros | 0.32 | 1 | 0.037 | 0.848 |
| juv3\_oroboros:chamber | 5.96 | 1 | 0.700 | 0.405 |
| Residuals | 638.26 | 75 |  |  |

**Table S5**. Mass and SVL basal ANOVA model output.

| Test | term | sumsq | df | statistic | p.value |
| --- | --- | --- | --- | --- | --- |
| Mass | basal\_corrected\_pmol | 0 | 1 | 0.61 | 0.437 |
| **sex** | **0** | **1** | **20.29** | **0.000** |
| **log(CORT\_Final\_Hormone\_ng\_mL)** | **0** | **1** | **5.03** | **0.028** |
| log(T4\_corrected\_ng\_mL) | 0 | 1 | 0.19 | 0.667 |
| Residuals | 0 | 67 |  |  |
| SVL | basal\_corrected\_pmol | 0 | 1 | 0.61 | 0.437 |
| **sex** | **0** | **1** | **20.29** | **0.000** |
| **log(CORT\_Final\_Hormone\_ng\_mL)** | **0** | **1** | **5.03** | **0.028** |
| log(T4\_corrected\_ng\_mL) | 0 | 1 | 0.19 | 0.667 |
| Residuals | 0 | 67 |  |  |

**Table S6** Mass and SVL ADP ANOVA model output.

| Test | term | sumsq | df | statistic | p.value |
| --- | --- | --- | --- | --- | --- |
| Mass | adp\_corrected\_pmol | 0.000 | 1 | 1.615 | 0.208 |
| **sex** | **0.000** | **1** | **21.928** | **0.000** |
| **log(CORT\_Final\_Hormone\_ng\_mL)** | **0.000** | **1** | **4.602** | **0.036** |
| log(T4\_corrected\_ng\_mL) | 0.000 | 1 | 0.085 | 0.771 |
| Residuals | 0.000 | 67 |  |  |
| SVL | adp\_corrected\_pmol | 0.000 | 1 | 2.149 | 0.147 |
| log(CORT\_Final\_Hormone\_ng\_mL) | 0.000 | 1 | 2.011 | 0.161 |
| log(T4\_corrected\_ng\_mL) | 0.000 | 1 | 0.030 | 0.864 |
| **sex** | **0.000** | **1** | **7.455** | **0.008** |
| Residuals | 0.002 | 67 |  |  |

**Table S7** Mass and SVL Oligo ANOVA model output.

| Test | term | sumsq | df | statistic | p.value |
| --- | --- | --- | --- | --- | --- |
| Mass | oligo\_corrected\_pmol | 0.000 | 1 | 0.447 | 0.506 |
| **log(CORT\_Final\_Hormone\_ng\_mL)** | **0.000** | **1** | **5.583** | **0.021** |
| log(T4\_corrected\_ng\_mL) | 0.000 | 1 | 0.163 | 0.688 |
| **sex** | **0.000** | **1** | **19.972** | **0.000** |
| Residuals | 0.000 | 67 |  |  |
| SVL | oligo\_corrected\_pmol | 0.000 | 1 | 0.482 | 0.490 |
| **sex** | **0.000** | **1** | **6.265** | **0.015** |
| log(CORT\_Final\_Hormone\_ng\_mL) | 0.000 | 1 | 2.750 | 0.102 |
| log(T4\_corrected\_ng\_mL) | 0.000 | 1 | 0.002 | 0.968 |
| Residuals | 0.002 | 67 |  |  |

**Table S8** Mass and SVL RCR ANOVA model output.

| Test | term | sumsq | df | statistic | p.value |
| --- | --- | --- | --- | --- | --- |
| Mass | RCR | 0.000 | 1 | 0.19 | 0.661 |
| **sex** | **0.000** | **1** | **19.77** | **0.000** |
| **log(CORT\_Final\_Hormone\_ng\_mL)** | **0.000** | **1** | **5.25** | **0.025** |
| log(T4\_corrected\_ng\_mL) | 0.000 | 1 | 0.10 | 0.751 |
| Residuals | 0.000 | 67 |  |  |
| SVL | RCR | 0.000 | 1 | 0.80 | 0.374 |
| **sex** | **0.000** | **1** | **5.68** | **0.020** |
| log(CORT\_Final\_Hormone\_ng\_mL) | 0.000 | 1 | 2.36 | 0.129 |
| log(T4\_corrected\_ng\_mL) | 0.000 | 1 | 0.04 | 0.843 |
| Residuals | 0.002 | 67 |  |  |