



Supplementary Figure 3: Expression of carp LHR and FSHR mRNA in various tissues as determined by real-time PCR. Each tissue represents a sample from three mature males and three mature females. The relative abundance of the mRNA was normalized to the amount of elongation factor 1 α (EF-1 α) by the comparative threshold cycle method (Δ CT). To receive fold expression each value was divided by the values of the hind brain tissue.