



Figure. Expression of a subgroup of lncRNAs in fast skeletal muscle from adults and juveniles sequenced by RNAseq. Boxplot showing differences in RNAseq reads mapped against the gilthead sea bream genome (as DESEQ2 counts) between juvenile (n=3) and mature (n=3) gilthead sea bream's fast skeletal muscle. Statistical differences between groups are indicated as follow: *p<0.05, **p<0.01 and ***p<0.001.