



Fig S6. We did not find significant differences between 17-20 dpf hybrid craniofacial samples and samples sequenced on other dates for A) median percent GC content across reads, B) number of normalized read counts, or C) number of raw fastq reads. the proportion of duplicate reads for each sample (Pairwise Welch two sample t test; $P < 0.0001 = ****$, $*** = 0.001$, $** = 0.01$, $* = 0.05$).