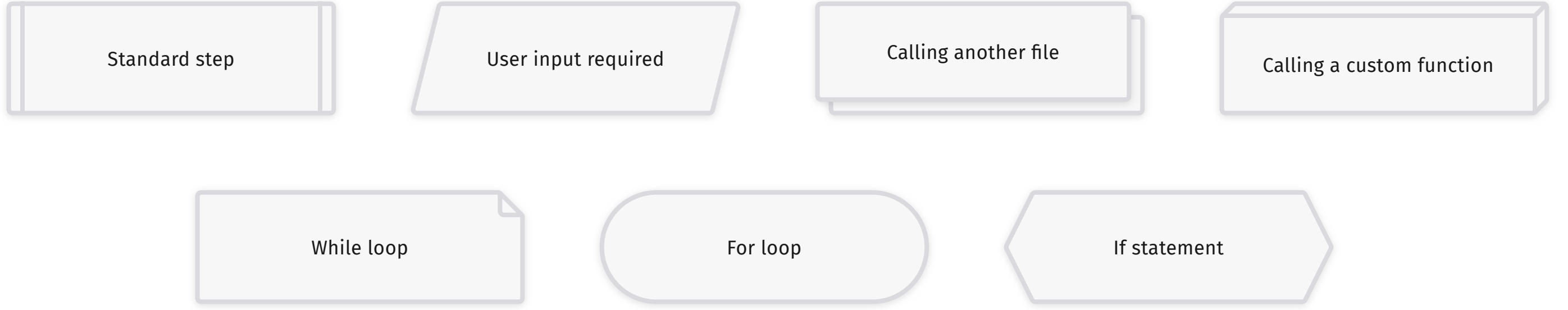
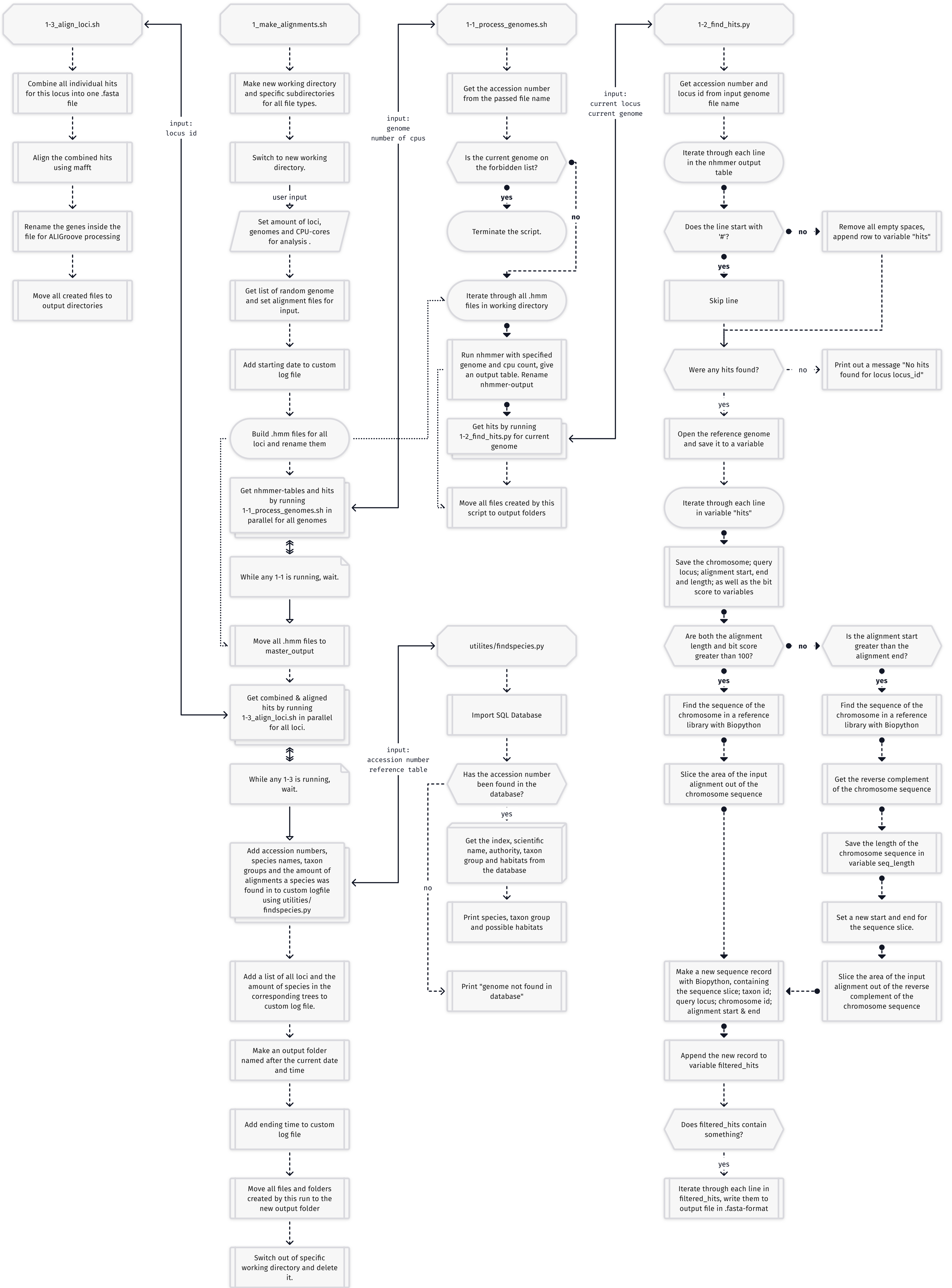


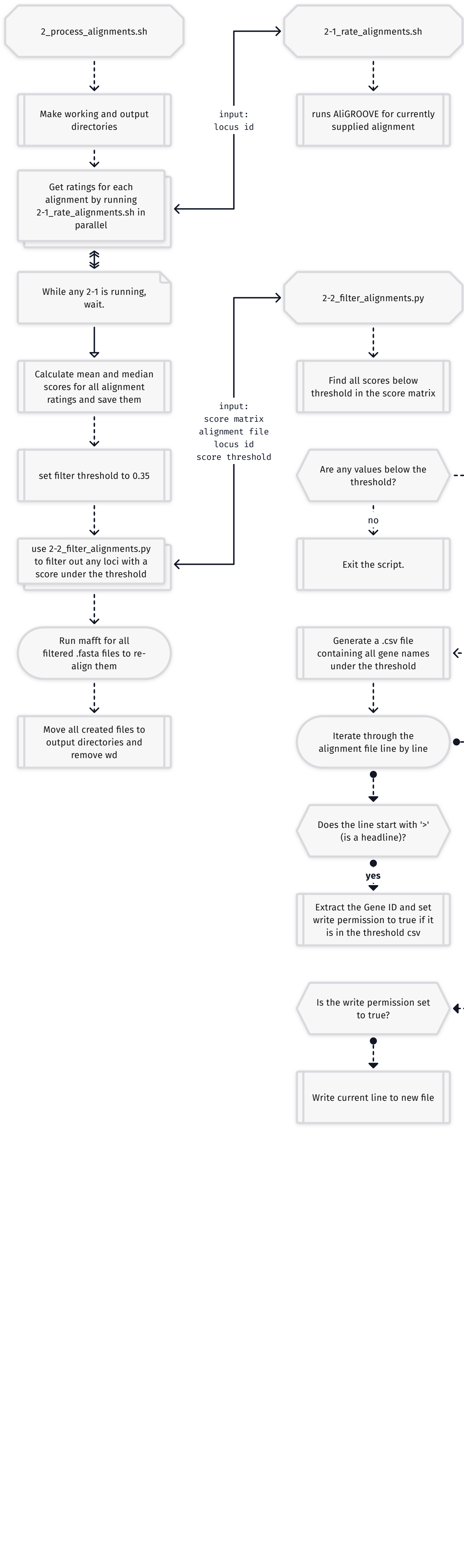
Cryptochrome evolution across fish species - genotree pipeline



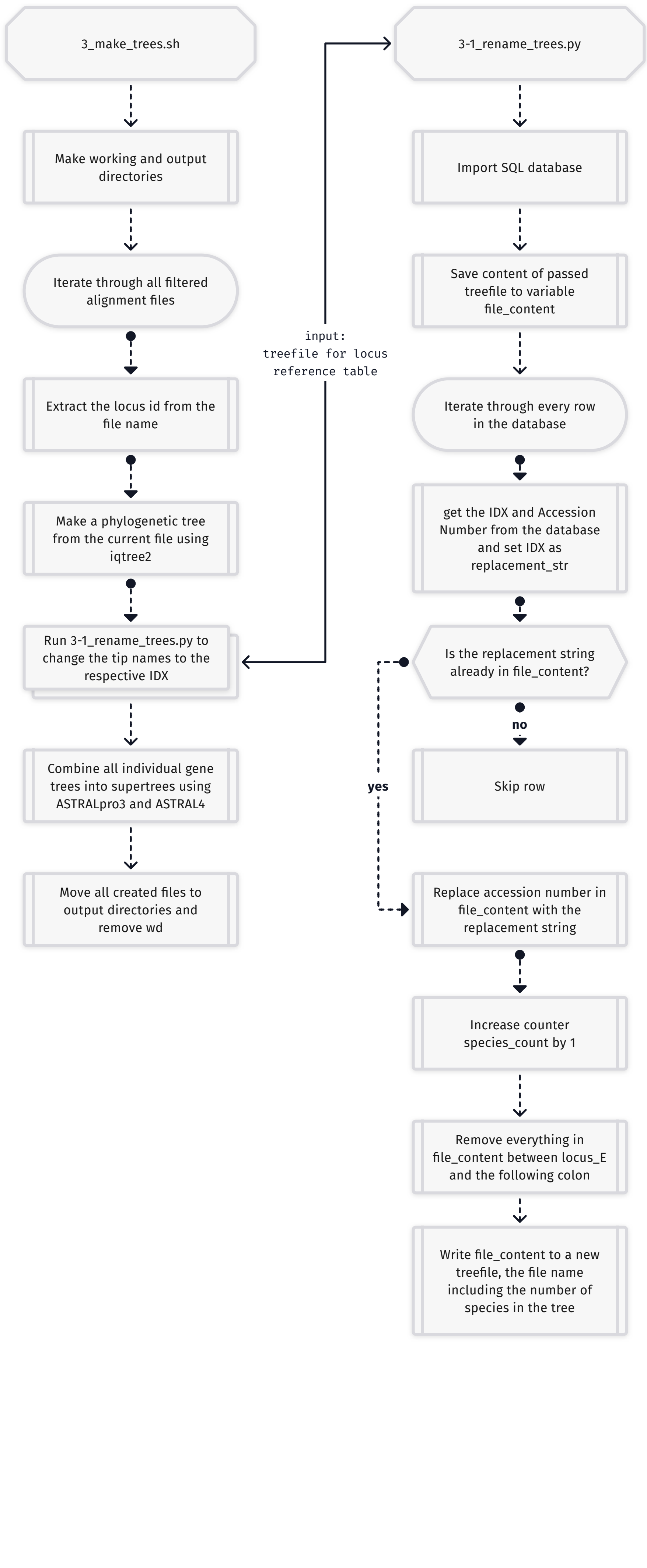
Section 1 - create alignments from raw genome and exon files



Section 2 - rate alignments and filter them for exons with bad scores



Section 3 - generate individual gene trees and combine them into a supertree



Section 4 - export generated phylogenetic trees as PDF files. Compare PhyloTrees using dendrograms

