## Output folder structure Results of searching for protein's found homolohous.tsv homologues with blastp assemblies list.tsv Proteins' RefSeg assemblies upstream\_sequences.fa annotated ORFs Upstream sequences fasta file ☐ locus\_id\_1.tsv ⊞ locus\_id\_2.tsv tsy tables with information about annotated ORFs in each locus. nt/aa\_orfs\_fasta\_files conserved\_ORFs\_set\_1.fa Fasta files with conserved ORFs conserved ORFs set 2.fa sequences nt/aa\_msa\_fasta\_files Multiple sequence alignments conserved\_ORFs\_set\_1.fa fasta files for conserved ORFs conserved ORFs set 2.fa Summary statistics for conserved results summary.tsv **ORFs** annotation\_visualisation conserved\_ORFs\_set\_1.pdf Visualisation of loci' annotation conserved\_ORFs\_set\_2.pdf nt/aa\_msa\_visualisation MSA visualisation conserved\_ORFs\_set\_1.pdf conserved\_ORFs\_set\_2.pdf Sequence logo visualisation nt/aa seglogo visualisation ₩ conserved\_ORFs\_set\_1.pdf Conserved\_ORFs\_set\_2.pdf