

Assemblies for primary comparison with BUSCO [1]

Name	Sources	N50, Kb
Minke whale (<i>Balaenoptera acutorostrata scammoni</i>)	http://www.ncbi.nlm.nih.gov/assembly/GCF_000493695.1/	Scaffold N50 12,843.7 Kb Contig N50 22.7 Kb
Bowhead whale (<i>Balaena mysticetus</i>)	http://alfred.liv.ac.uk/downloads/bowhead_whale/bowhead_whale_proteins.zip http://alfred.liv.ac.uk/downloads/bowhead_whale/bowhead_whale_scaffolds.zip	Scaffolds N50 877 Kb Contigs N50 34.8 Kb
Antarctic minke whale (<i>Balaenoptera bonaerensis</i>)	http://www.ncbi.nlm.nih.gov/assembly/GCA_000978805.1	Scaffold N50 20 Kb Contig N50 8 Kb

References

1. Simão FA, Waterhouse RM, Ioannidis P, Kriventseva EV, Zdobnov EM. BUSCO: assessing genome assembly and annotation completeness with single-copy orthologs. *Bioinformatics*. 2015;31:3210-12.