

### The primary analysis with the BUSCO methodology [1]

Source	Complete BUSCOs (%)	Duplicated, (included in complete) (%)	Fragmented (%)	Missing, %
Grey whale/initial	24	0.4	24	50
Minke whale/proteins	97	44	1.5	0.6
Bowhead whale/genome	60	1.3	25	13
Bowhead whale/proteins	72	2	14	12
Antarctic minke whale	NA	NA	NA	NA

### 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.