CORRECTION

Correction: A First Insight into the Genome of the Filter-Feeder Mussel *Mytilus galloprovincialis*

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A reference is omitted from the fourth and seventh sentences of the second paragraph under the subheading “Mitochondrial genomes” in the Results and Discussion section.

The fourth sentence should read: The largest number of reads (113,824) mapped onto an F mitochondrial genome from *M. edulis* (MeF, GenBank KM192128, Zbawicka et al., 2014, Fig 4A), slightly more (111 more) than the number of reads that mapped to MgF.

The seventh sentence should read: Surprisingly, a well-covered mapping against an M mitochondrial genome of *M. edulis* (MeM, Genbank KM192129, Zbawicka et al., 2014) was also observed (Fig 4B).

The reference is: Zbawicka M, Wenne R, Burzyński A. Mitogenomics of recombinant mitochondrial genomes of Baltic Sea *Mytilus* mussels. Mol Genet Genomics. 2014;289: 1275–1287. doi:[10.1007/s00438-014-0888-3](https://doi.org/10.1007/s00438-014-0888-3).

References

1. Murgarella M, Puiu D, Novoa B, Figueras A, Posada D, Canchaya C (2016) A First Insight into the Genome of the Filter-Feeder Mussel *Mytilus galloprovincialis*. PLoS ONE 11(3): e0151561. doi: [10.1371/journal.pone.0151561](https://doi.org/10.1371/journal.pone.0151561) PMID: [26977809](https://pubmed.ncbi.nlm.nih.gov/26977809/)
2. Zbawicka M, Wenne R, Burzyński A. Mitogenomics of recombinant mitochondrial genomes of Baltic Sea *Mytilus* mussels. Mol Genet Genomics. 2014; 289: 1275–1287. doi: [10.1007/s00438-014-0888-3](https://doi.org/10.1007/s00438-014-0888-3) PMID: [25079914](https://pubmed.ncbi.nlm.nih.gov/25079914/)

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