Throughout the Results section, we report the percentage of samples containing seabirds or target taxa, representing the predation incidence of a population of seals.

Both in the Results where we did not detect significant difference for penguins using eDNA and morphological remains, page 10, paragraph 2; and in Discussion, the sensitivity of our methods are compared to other studies, page 12 paragraph 2, and discussed throughout, also regarding haplotypes.

We report a range of predation incidence throughout the MS, in the Abstract, Results and Discussion, for example page 12, paragraph 2 and the Results, page 10 and carrying over into the second paragraph of page 11.

For DNA filtering and sequence quality control, we explain extensively the criteria for inclusion and provide a detailed supplement to reproduce this work. For analyses, only one sample was explicitly excluded from statistical analyses, noted at page 8, paragraph 3. This sample was processed but was the only sample we obtained from one of the same locations.

Methods, page 8, paragraph 3, and page 9, paragraph 1, where we describe our model fitting process, the model architecture and data format.