

References – Supplementary information

1. Berry TE, Osterrieder SK, Murray DC, Coghlan ML, Richardson AJ, Grealley AK, et al. DNA metabarcoding for diet analysis and biodiversity: A case study using the endangered Australian sea lion (*Neophoca cinerea*). *Ecol Evol.* 2017;7(14):5435-53.
2. Deagle BE, Gales NJ, Evans K, Jarman SN, Robinson S, Trebilco R, et al. Studying Seabird Diet through Genetic Analysis of Faeces: A Case Study on Macaroni Penguins (*Eudyptes chrysolophus*). *PLoS ONE.* 2007;2(9):e831.
3. Pochon X, Bott NJ, Smith KF, Wood SA. Evaluating Detection Limits of Next-Generation Sequencing for the Surveillance and Monitoring of International Marine Pests. *PLOS ONE.* 2013;8(9):e73935.
4. Davies Claire H, Armstrong Amelia J, Baird M, Coman F, Edgar S, Gaughan D, et al. Over 75 years of zooplankton data from Australia. *Ecology.* 2014;95(11):3229-.
5. ALA. Atlas of Living Australia website <http://www.ala.org.au> 2016
6. Thompson H. Pelagic tunicates of Australia. 1948.
7. Anderson M, Gorley RN, Clarke RK. *Permanova+ for Primer: Guide to Software and Statistical Methods*: Primer-E Limited; 2008.
8. Roberts DW. *labdsv: Ordination and Multivariate Analysis for Ecology.* 2016;R package version 1.8-0.
9. Australian Bureau of Meteorology. Australian climate variability & change - Time series graphs at: <http://www.bom.gov.au/climate/change/> (data accessed in December 2018).
10. Kears M, Moir R, Wilson A, Stones-Havas S, Cheung M, Sturrock S, et al. Geneious Basic: An integrated and extendable desktop software platform for the organization and analysis of sequence data. *Bioinformatics.* 2012;28(12):1647-9.

11. Peters KJ, Ophelkeller K, Bott NJ, Deagle BE, Jarman SN, Goldsworthy SD. Fine-scale diet of the Australian sea lion (*Neophoca cinerea*) using DNA-based analysis of faeces. *Marine Ecology*. 2014:n/a-n/a.