

Massively parallel sequencing-enabled mixture analysis of mitochondrial DNA samples

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<https://doi.org/10.1007/s00414-018-1799-3> [Journal metrics at Scopus](#) [Article at Scopus](#) [Journal metrics at WOS](#) [Article at WOS](#)

Monitoring of the toxic dinoflagellate *Alexandrium catenella* in Osaka Bay, Japan using a massively parallel sequencing (MPS)-based technique

Nagai, Satoshi; Chen, Hungyen; Kawakami, Yoko; Yamamoto, Keigo; **Sildever, Sirje** Harmful Algae 2019 / art. 101660, 13 p. : ill
<https://doi.org/10.1016/j.hal.2019.101660>

Parsing apart the contributors of mitochondrial DNA mixtures with massively parallel sequencing data

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