

Short Note

Ecotype Origin of an Entangled Killer Whale (*Orcinus orca*) Identified with Remnant mtDNA

Supplemental Table

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Supplemental Table 1. A variable sites table of the killer whale (*Orcinus orca*) control region haplotype alignments. The entangled killer whale's sequence is highlighted in yellow as the basis for comparison. Variable nucleotides are highlighted in blue. A colon (:) indicates a nucleotide insertion that is not present in the reference; a period (.) indicates the nucleotide is identical to the reference.

Genbank Code/Haplotype	Corresponding KW ecotype	6	133	147	276	284	304	392	409	458	503	505	542	626
OR661229 HMSC	Transient/Bigg's	:	T	A	G	T	A	C	G	T	T	C	C	T
GU187162.1 ENPTSEA2	Transient/Bigg's	:
DQ399075.1 NT1	Transient/Bigg's	:	.	G	A
DQ399076.1 NT2	Transient/Bigg's	:	C	G
DQ399080.1 GAT2	Transient/Bigg's	:	.	G	.	.	.	T
DQ399081.1 GAT	Transient/Bigg's	:	.	G
DQ399082.1 AT1	Transient/Bigg's	:	.	G	.	.	G	.	A
DQ399077.1 SR	Resident (Southern)	A	.	G	.	.	.	T	A	C	C	.	T	.
DQ399074.1 NEWR	Resident (Other)	:	.	G	.	C	.	T	A	C	C	.	T	.
DQ399078.1 NR	Resident (Northern)	A	.	G	.	.	.	T	A	C	C	T	T	.
DQ399079.1 OFF	Offshore	:	.	G	.	.	.	T	A	C	C	.	T	C