**Supplemental Table S1.** Thirteen haplotypes of captive *Tursiops truncatus* defined by 35 variable sites within a 373bp fragment of the mitochondrial control region.

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| **Haplotype**  **Accession number** | **0**  **1**  **7** | **0**  **2**  **2** | **0**  **2**  **8** | **0**  **2**  **9** | **0**  **4**  **3** | **0**  **5**  **3** | **0**  **5**  **7** | **0**  **6**  **3** | **0**  **6**  **4** | **0**  **6**  **6** | **0**  **6**  **8** | **0**  **6**  **9** | **0**  **7**  **2** | **0**  **7**  **3** | **0**  **7**  **4** | **0**  **7**  **5** | **0**  **7**  **6** | **0**  **8**  **1** | **1**  **0**  **1** | **1**  **0**  **8** | **1**  **4**  **1** | **1**  **7**  **6** | **1**  **7**  **9** | **1**  **8**  **0** | **1**  **9**  **7** | **2**  **0**  **5** | **2**  **0**  **9** | **2**  **2**  **5** | **2**  **2**  **8** | **2**  **3**  **6** | **2**  **4**  **2** | **2**  **5**  **9** | **2**  **7**  **3** | **3**  **2**  **0** | **3**  **4**  **7** |
| H1 KX151147 | G | T | C | G | T | A | A | G | G | A | G | A | G | T | A | G | - | T | A | A | G | A | A | T | G | A | A | - | T | C | C | T | - | G | - |
| H2 KX151148 | . | . | . | . | . | . | . | A | . | G | A | . | A | A | T | A | G | G | G | . | A | . | . | . | A | . | . | A | C | T | T | . | G | . | - |
| H3 KX151149 | . | . | . | . | . | . | G | . | A | G | A | . | A | A | T | A | G | G | G | . | A | . | . | . | . | . | . | A | . | T | T | . | - | A | C |
| H4 KX151150 | . | . | . | . | . | . | . | . | . | G | A | . | A | A | T | A | G | G | G | . | A | . | . | . | A | . | . | A | C | T | T | . | G | . | - |
| H5 KX151151 | A | . | . | . | C | . | . | A | . | . | . | . | A | . | . | . | - | . | . | . | . | . | . | . | . | . | . | - | . | . | . | . | - | . | - |
| H6 KX151152 | . | . | . | . | . | . | . | . | A | G | A | . | A | A | T | A | G | G | G | . | A | . | . | . | . | . | . | A | . | T | T | . | - | A | - |
| H7 KX151153 | . | . | . | . | . | G | . | . | A | G | A | . | A | A | T | A | G | G | G | . | A | . | . | . | . | . | . | A | . | T | T | . | - | A | - |
| H8 KX151154 | . | C | . | . | . | . | . | . | A | G | A | . | A | A | T | A | G | G | G | . | A | . | . | . | . | . | . | A | . | T | T | . | - | A | - |
| H9 KX151155 | . | . | . | . | . | . | . | A | A | G | A | . | A | A | T | A | G | G | G | . | A | . | . | . | A | . | . | A | C | T | T | . | G | . | - |
| H10 KX151156 | . | C | . | . | . | . | . | . | A | G | A | T | A | A | T | A | G | G | G | . | A | . | . | . | . | . | . | A | . | T | T | . | - | A | - |
| H11 KX151157 | A | . | . | . | C | . | . | A | . | . | . | . | A | . | . | . | - | . | . | . | . | . | . | G | . | . | . | - | . | . | . | . | - | . | - |
| H12 KX151158 | . | . | . | . | C | . | . | . | . | . | . | . | A | . | . | C | T | . | . | . | . | . | . | . | . | . | . | - | . | . | . | . | - | . | - |
| H13 KX151159 | . | . | T | C | . | . | . | A | . | G | A | . | A | A | T | A | G | G | G | T | A | G | G | . | A | T | G | A | C | T | T | A | G | . | - |

(.) represents identity with haplotype H1 (GenBank accession no. KX151147), (-) represents a gap in the alignment (insertion/deletion).