**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).