**Appendix C. List of cytochrome *b* haplotypes shared by several individuals**

**Casinycteris argynnis**

**– CA1**(24 individuals, 6 localities):

DRC 29: K13x301, K13x302, K13x303, K13x314, T2228, T2230, T2231, T2235, T2239.

DRC 30: K13x280, K13x281, K13x283, K13x297, T2225, T2227, T2232, T2245.

DRC 28: K13x163, K13x164, K13x172, K13x186.

DRC 23: T2259.

DRC 32: T2243

DRC36: NBC24558.

**– CA2**(7 individuals, 4 localities):

DRC 25: T2236, T2241, T2242.

DRC 26: T2260, T2261.

DRC 27: T2258.

DRC 28: K13x243.

**– CA3**(4 individuals):

DRC 31: T2244, T2246, T2247.

DRC 29: T2229.

**– CA4**(3 individuals, RC 19): T104, T143, T149.

**– CA5**(2 individuals, RC 20): 2209, T81.

**Scotonycteris**

**Cameroon**

**– SZ1** (7 individuals, 2 localities):

C 9: MNHN 2011–675, MNHN 2011–676, MNHN 2011–677, MNHN 2011–678, C07X178, C07X183.

C 10: CM 107997.

**– SZ2** (2 individuals, 2 localities):

C 12: ZMB54390

C 13: MNHN 2011–682

**– SZ3** (2 individuals, C 13): C07X53, C07X63.

**Western Equatorial Africa**

**– SZ4** (2 individuals, 2 localities):

EG 14: ZMB50004.

G 15: FMNH162156

**– SZ5** (2 individuals, 2 localities):

CAR 22: R08119.

RC 19: T77.

**– SZ6** (2 individuals, 2 localities):

G 17: 1337.

G 18: 1380.

**– SZ7** (2 individuals, G 16): FMNH167759, FMNH167761.

**Eastern DRC**

**– SZ8** (4 individuals, 3 localities):

DRC 26: T2262, T2263.

DRC 28: K13x203.

DRC 29: K13x312.

**– SZ9** (3 individuals, DRC 29): K13x311, T2237, T2238.

**– SZ10** (2 individuals, 2 localities):

DRC 34: SMF65010.

DRC 35: NBC24197.

**Upper Guinea**

**– SZ11** (3 individuals, L 1): 2012x545, 2012x546, 2012x547.

**– SZ12** (2 individuals, 2 localities):

IC 3: KB730.

IC 7: KBx675.