TABLE 6. Nucleotide composition at variable sites in cytochrome b target region for 49 unique standard type and caviar test sequences

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AstG20904   : CCACCTACACCACTTCCTTAACCGCCCCGTCAACACGATACTCGCCTACTGTTCGAAATACCTACCCAACCAAACCCGT
AstG20907   : ....G.........................................................................
8378L1AU14  : ...........................G...........................T...............
AstG20909   : ...........................................................................A.
8304L2AU25  : ...........................................................................C
8508L67U32  : .......T...................................................................
8292L1AU2   : ..............C............................................................
8324L1AU8   : ...........................................................C...............
AbaG41168   : ......G...............TAT...A...........T..........C.......G.T...............
AbaG11324   : ...T..G...............TAT...A...........T..........C.......G.T...............
8325L1AU9   : T.....G...............TAT...A...........T..........C.......G.T...............
8513L81U35  : ....G.G...............TAT...A...........T..........C.......G.T...............
8662L20U38  : ......G...............TAT...A...........T..........C.......G.T...A.............
8513L11U34  : ....G.................TAT...A...........T..........C.......G.T...............
AguG20874   : ......G...............T.T...A...........T..........C.........T.......T.........
ApeG20903   : ......G...............T.T...A...........T..........C.........T.......T.........
8623L4U31   : ....G.G...............T.T...A...........T..........C.........T.......T.........
AguG20923   : ......G...............T.T...A...........T..........C.........T.G.....T.........
9309L2AU43  : ......G...............T.T...A...........T..........C.........T.......T.G.......
8663L6BU28  : .......T..............T.T...A...........T..........C.........T.......T.........
8609L2AU27  : ......G...............T.T...A...........T.....C....C.........T.......T.........
8487L2AU3   : ......G................T...A...........T..........C.........T.......T.........
9330L10U21  : ......G...............T.T...A...........T..........C.........T.......TT........
8519L2BU24  : ......G...............T.T...A...........T......T...C.........T.......TT........
9033L4BU30  : ....G.G...............T.T...A...........T..........C.........T.......TT........
HhuG20913   : .......T..T.......C.G.......A.T.........T.....C....C.......GT....T............C
9033L12AU3  : .......T..T.......C.G.......A.T.....A...T.....C....C.......GT....T............C
8346L2AU12  : .......T..T.......C.G.......A.T.........T.....C....C.......GT....T......G.....C
8417L1BU17  : .......T..T.......C.G.......A.T...T.....T.....C....C.......GT....T............C
8513L90U36  : ....G..T..T.......C.G.......A.T.........T.....C....C.......GT....T............C
HdaF40717   : ............T...T.....T....TA...........T...T.C....C...G.....TC..........G..T..
AshF40705   : ..G.........T.........T....T..........C.T...T.C........G......C.............T..
AtrF10246   : ..G.........T.........T....T.......T..C.T...T.C........G......C.............T..
G102011U7   : ..G...G.G...T........TT....TA...G.....C.T...T.C........G......C.........G...TAC
9123L1AU45  : ..T..A..GT..T.C...C.......TTTC.....T..C.T..A.TC..CCC..A.C.C..TC...TC.....GT..AC
PspGBM6491  : ..T..A..GT..T.C...C.......TTTC.....T..C.T..A.TC..CCC....C.C..TC...TC.....GT..AC
F40232U42   : ..T..G..GT..T.C...C.......TTTC.....T..C.T..A.TC..CCC....C.C..TC...TC.....GT..AC
PspD10570   : ..T..G..GT..T.C...C...T...TTTC.....T..C.T..A.TC..CCC....C.C..TC...TC.....GT..AC
SplF11159   : .......T..T.................AC.G......CG.CTC.......C...C......CC.........G..TAC
SsuH10433   : .......T..T.................AC.G......CG.CTC.......C...C..C...CC.........G..TAC
SalB40622   : ..G....T...................AC.G......CG.CTC.......C...C......CC.........G..TAC
AnuG41203   : ......G...............T....TA..G........T..........CCT....CG...................
B1AU13      : ......................T....TA..G........T..........CCT....CG...................
AfuE40805   : ......G....G.C.....GG...T...A...........T...T......C...G.....T...............
AoxF41302   : ......GG..T....T.C.........TA...........T.....C.TC.C...G.GC...C.....G.TGG..T...
AruG41158   : ..G...G.......C.......T....TA....T...C..T...........C.....CG...TT.............C
AruG20915   : .TG...G.......C.......T....TA....T...C..T...........C...G.CG...TT.............C
AruG20916   : .TG...G.......C.......T....TA....T...C..T...........C.....CG...TT.............C
AmeF10704   : ............T.C.T.....T..T.TA...G.T.....T...T.C....C...G.....TC................
AmiG41169   : ............T.C.T.....T..T.TA...G.T.....T...T.C....C...G.....TC................
AbrF10241   : ......G...............T.T...A...........T...T......C.....G.G.T...............
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
              3461111223334445667778999911111111111111111111111111111111122222222222222222222
                 0289170160587062587013601112222233333445556667778888899900011112223444445555
                                       5147036792458914039289157036792581470369028703679235 8
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