

Alu consensus sequence for 8 Alu subfamilies lineage specific to Cebus monkey [Cebus_imitator-1.0]

>subfamily1_Ceb

-GCCGGGCGCGGTGGCTACGCCTGTAATCCCAGCACTTGGGAGGCCGAGGCAGGGAT-CAC--
GAGGTCAAGAGATCGAGACCA-TCCCTGGCAACATGGTAAACCCCGTCTACTAAAA-ATACAAAAA---
TTAGCTGGGATGGTGGCACGCGCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATTGCTGAACCCA
GGAGGCAGGTTGCAGTGAGCCGAGATTGCCACTGCACTCCAGCCTGG-----CGACA-
GAGCGAGACTCCGTCTCA-----

>subfamily6_Ceb

GGCCGGGCGCGGTGGCTACGCCTGTAATCCCAGCACTTGGGAGGCCGAGGCAGGGAT-CAC--
GAGGTCAAGAGATCGAGACCA-TTCTCGCC---ATGATGAAACCCCGTCTACTAAAA-ATACAAAAA---
TTAGCCGGGTGTGGTGGCACGCGCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATTGCTGAACCCA
AGGAGGCAGGTTGCAGTGAGCCGAGATTGCCACTGCACTCCAGCCTGG-----CGACA-
GAGCAAGACTCCGTCTCA-----

>subfamily7_Ceb

-GCCGGGCGCGGTGGCTACGCCTGTAATCCCAGCACTTGGGAGGCCGAGGCAGGGAT-CAC--
GAGGTCAAGAGATCGAGACCA-TCCCTGGCAACATGGTAAACCCCGTCTACTAAAA-ATACAAAAA---
TTAGCTGGGATGGTGGCACGCGCTGTGGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATTGCTGAACCCA
AGGAGGCAGGTTGCAGTGAGCCGAGATTGCCACTGCACTCCAGCCTGG-----TGACA-
GAGCAAGACTCCGTCTCA-----

>subfamily3_Ceb

-GCCGGGCGCGGTGGCTACGCCTATAATCCCAGCACTTGGGAGGCCGAGGCAGGGAT-CAC--
GAGGTCAAGAGATCGAGACCA-TCCCTGGTCAACAAGGTAAACCTCGTCTACTAAAA-ATACAAAAA---
TTAGCTGGGATGATGGTGGCATGCCGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATTGCTGAACCCA
GGAGGCAGGTTGCCATTGCACTCCAGCCTGGG-----
CAACACAGCGAAACTCCGTCTCA-----

>subfamily5_Ceb

-GCCGGGCGCGGTGGCTCAAGCCTGTAATCCCAGCACTTGGGAGGCCGAGGCAGGGAT-CAC--
GAGGTCAAGAGATCGAGACCA-TCCCTGGTCAACAAGGTAAACCTCGTCTACTAAAA-ATACAAAAA---
TTAGCTGGGATGATGGTGGCATGCCGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATTGCTGAACCCA
GGAGGCAGGTTGCCATTGCACTCCAGCCTGGG-----
CAACACAGCGAAACTCCGTCTCA-----

>subfamily4_Ceb

-GCCGGGCGCGGTGGCTACGCCTGTAATCCCAGCACTTGGGAGGCCGAGGCAGGGAT-CAC--
GAGGTCAAGAGATCGAGACCA-TCCCTGGTCAAATGGTAAACCCCGTCTACTAAAA-ATACA-A-
AAAATTAGCTGGGATGGTGGCGCTGTAGTCCCAGCTACTCAGGAGGCTGAGGCAGGAGAATTGCTGA

ACCCAGGAGGC GGAGGTT GCGGT GAGCC GAGAT CGCGC CATT GC ACT CCAGC CTGGG-----
TAACAAGAGCGAAACTCCGACTCA-----

>subfamily2_Ceb

-GCCGGGCGCGGTGGCTCAAGCCTGTAATCCCAGCACTGTGGGAGGCCGAGGC GGTTGGAT-CAC--
GAGGTCAAGAGATCGAGACCA-TCCTGGTCAACATGGT GAAACCCG TCTACTAAAA-ATACA-AAAA-
TTAGCTGGGATGGTGGCGCGTGCCTGTAATCCCAGCGACTCAGGAGGCTGGGCAGGAGAATTGCCTGAACCC
GGGAGGC GGAGGTT GCGGT GAGCC GAGAT CGCGC CATT GC ACT CCAGC CTGGG-----
TAACAAGAGCGAAACTCCGCTCA-----

>subfamily8_Ceb

-GCCGGGCGCGGTGGCTCAAGCCTGTAATCCCAGCACTGTGGGAGGCCGAGGC GGTTGGAT-CAC--
GAGGTCAAGAGATCGAGACCA-TCCTGGTCAACATGGT GAAACCCG TCTACTAAAA-ATACA-AAAA-
TTAGCTGGGATGGTGGCGCGTGCCTGTAATCCCAGCGACTCAGGAGGCTGGGCAGGAGAATTGCCTGAACCC
GGGAGGC GGAGGTT GCGGT GAGCC GAGAT CGCGC CATT GC ACT CCAGC CTGGG-----
TAACAAGAGGGAAACTCCGCTCA-----