

Matched pairs of 5' and 3' integration sites that come from different koalas.

>P_3rS_S2.74718_S3.57590_11S8__C_5fU_3S8

TATTTGAGCTCAGGCAACTGGGGTTAAGTGACTTGCCCAGAGTCACAAGTCTGA
GGTCGGATTTGAA

>P_3rU_4S8_P_3fU_S8.70215__C_5fU_S4.684182_5fU_S4.1661030

TTAAAGACAGCAATCTTCTGTCTATTCTTCAAGATTGGTTTGCCATTTCTTCTCC
AGCTCATTTTATGGAT

>P_3rU_4S8_P_3fU_S8.70215__C_5fU_S4.1683056

TTAAAGACAGCAATCTTCTGTCTATTCTTCAAGATTGGTTTCTCATCTGTAAAAT
GGGGATAATAACA

>P_3rU_S3.26225__C_5fU_S8.589094

AATTGAGGAACTATGCCCAAAGGGCTATAAACTGCATACCCTTTGATCCAGCA
ATACCACTTCTTGG G

>P_3rU_S8.97760__C_5fU_S3.480683

ACTGAGCCATATAACCATATAGTATCTCATTGTAAACAAATCATTACATGCACAC
ATTCCCACCAATGCATGCTGGACTTCCTGACAAAGTACAACATGCTCACCTGCC
AACACTTGCTTGTTAAGACCTGTCAGTGGACTTGACCACTGATGGGTTATAGCTT
GCATTT

>P_3rS_S5.105847_5S8__C_5fU_S4.1481035

TGAATGAACATTTTCTCTACTCCGCCATCTTGGCTCCACCCCCC

>C_3rU_S8.1018030_P_3rS_S4.143028_6S3_S2.99154_17S8_P_3fU_2S8_S_3fU
_S20.58049__P_5rU_S9.30145

GGCTGATTCGGA CT CAGGTGAGTCTTCCA ACTCCAGGGCTGGCACTCTATCCAT
TCCCCACCTACCTGCCCTCCCCACATTCCTTC AAACCCTCTGTC

>P_3rS_S3.41285_S4.156475_S5.66477_S7.113364_S8.73375_S9.5274_36S2__S
_5rU_S18.20949_C_5rU_S2.111871

AGGAGGTTGAACCAGATGACCTCTGGGGTCTCTTTTAGCC

>S_3rU_S17.112853_P_3rU_8S2__S_5rU_S23.5584_2S20_C_5rS_3S3_S7.4344
GTAAATTGGTCTGAGGCTGGATTTGAACTCAGGTC

>P_3rU_4S8_P_3fU_S8.70215__C_5rU_S4.1173169
TTAAAGACAGCAATCTTCTGTCTATTCTTCAAGATTGGTTTGTCAATTCCTTCTCC
AGCTCATG

>P_3rS_S2.74718_S3.57590_11S8__C_5rU_S9.964451
TATTTGAGCTCAGGCAACTGGGGTTAAGTGACTIONGCCAGATCGGAAGAGCGT

>C_5fU_S2.547057__P_3fU_S4.4833
TCAGCAATACAAGGATCTAAGACATCTCCA

>C_5fU_S8.1066112_S8.1022169_S8.1055816__S_3fU_S20.2502
TTCAAATCCGACCTCAGACTTGTGACTCTGGGCAAGTCACTTAACCCCAGTTGC
CTCAGATCCAATTCACAT

>C_5fU_S4.684182_5fU_S4.1661030__S_3fU_S17.64559_P_3fU_S8.56518
ATCCATAAAATGAGCTGGAGAAGGAAATGGCAACTIONAGTTTCCCAGTCTATTTCA
TCCGGAGTTGT

>C_5fU_S4.1683056__S_3fU_S17.64559_P_3fU_S8.56518
TGTTATTATCCCCATTTTACAGATGAGGAACTIONAGTTTCCCAGTCTATTTTCATCC
GGAGTTGT

>C_5fU_S2.729164__S_3fU_4S20_P_3fU_S3.74019
CTTTATTCACCTGTGCCACCTAGATGCCCTGAAGAATATATTTTAGGCATATAAA
TGTGTAT

>C_5fU_S4.1481035__P_3fU_S8.33956
GGGGGGTGGAGCCAAGATGGCGGAGTAGAAGATGAGGAAAAGGCTCACAGAA
GG

>S_5rU_cluster_620_C_5rU_2S8_P_5rU_S8.78052__P_3fU_S3.22152
AGAGGTCCAGGCAAAGAGCTATGATCCCCGGTTTCTGCTTTCCTTCTAGTTAAA
TCGGA

>C_5rU_S2.906334__S_3fU_4S20_P_3fU_S3.74019

GGTCACCCAGCTAGTAAATATCTGAGGCCAGATTTGAGTCTTTCTGACTTCAGG
CCCTGCACTTTATTCAGTGTGCCACCTAGATGCCCTGAAGAATATATTTTAGGC
ATATAAATGTGTAT

>C_5rU_2S3__P_3fS_5S3_S2.110884_S8.68922

GGCCGGTGCTCTATTCAGTGTGCCACCTAGATGAGATCG

>C_5rU_S4.1173169__S_3fU_S17.64559_P_3fU_S8.56518

CATGAGCTGGAGAAGGAAATGACAACTAGTTTCCCAGTCTATTTTCATCCGGAG
TTGT

>S_5rU_S23.5584_2S20_C_5rS_3S3_S7.4344__P_3fS_8s2_1s5

GACCTGAGTTCAAATCCAGCCTCAGACAAATTCCAAGAAAAAGTTAGCTCTTTCC
CCTTCCTCCCCCTC CTGTGCCAT

>P_5rU_S4.11115__S_3fU_S20.10939_P_3fU_S3.74062

TTCAAGCAGAAGACGGCATAACGAGATAGAGGCGGTGACTGGAGTTCAGACGTG
TGCTTTGCCGATCTGAAGACACAGTAGTCAATGACTATAGTAGTCTTC