

Matched pairs of 5' and 3' integration sites that come from the same koala.

>P_3rU_6s8_cluster_446_cluster_937__C_5fU_S8.120509_S8.334494

GAATTAGAAATCGAGGAGATGCCCATCAATTGGGGAATGGCTGAACAAGTCATG
GTATAAGGAATTATGATGTGATGTTGTCTTCATGATCCCATTTAGAGTTTTCTTG
GCAAAGA

>P_3rU_cluster_564_S_3fS_cluster_29_cluster_977_cluster_4_cluster_n1_cluster_2
8__C_5fU_2S3_S_5rS_cluster_7_C_5rS_S3.746665_S4.29352_S_S9.36057_S1.30
664_S5.12084_S10.22389

GAACCCACAAAAGACAGAATGAAACAAATATCCAGCCCAATACAGCCTGGATG
ATTGATCAGAAGGGTGTATCGCGC CAGGCTGGGAGCACAG

>S_3rU_S17.112853_P_3rU_8S2__P_5fS_cluster_35

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>P_3rU_S8.18869_S8.19758_S8.24709__P_5fU_cluster_54_C_5fU_2S8

AGTGGCTTGCCCCGGGGCACACAGCTAGTATGTATCGGAGGCTGGATTTGAAC
TCAGGT

>P_3rS_cluster_175.maffT1s4-6s8__P_5fU_cluster_54_C_5fU_2S8

AGGAAACTGAGGCAAAGTTAAGTGATTTGCCGGGTCACACAGCTAGTATGTATC
GGAGGCTGGATTTGAACTCAGGT

>P_3rU_S2.133819_S2.154135__P_5fU_S2.2639_C_5fU_S2.23072_S2.547057

GACAGCATTTTCCATCCTGCGGCCTCTGGAGATGTCTTAGATCCTTGATTGCT
GA

>P_3rS_S10.119113_7S2_P_3fU_S2.18246_S_3fS_cluster_68_S_3rU_S15.6385__
P_5fU_S2.2639_C_5fU_S2.23072_S2.547057

TATGCTTCAATCTACATTCAAACCTCCGTAGTTCTTTCTTTGGATGTGGATAGCATT
TTTCATCATGAGGCCTTTGGAGATGTCTTAGATCCT TGTATTGCTGA

>P_3rS_cluster_31_cluster_252_P_3fS_S2.18623_S2.14939_S_3rU_2S17__P_5fU
_S2.2639_C_5fU_S2.23072_S2.547057

CTTTCCGATTTGACAGCATTTTCCATCCTGAGGCCTCTGGAGATGTCTTAGATC
CTTGATTGCTGA

>P_3rU_2S8_P_3fU_7S8__C_5fU_3S8

CCGATCTTTGCCTGAGGTTGCACACTTGGTCATATTTGAGCTCAGGCAACTGGG
GTTAAGTGAAGTGGCCAGAGTCACAAGTCTGAGGT CGGATTTGAA

>P_3rU_S8.12674_S8.39898__C_5fU_S8.950860

GACACCTCGGTGTGTCTCAGTTTCCTCATCTATAATATGAGCTGGAAAAGGAAAT
GGCACACTACTCTAGTATCTT TGCCAAGAAAACCC

>P_3rU_7S3__C_5fU_S3.480683

TATTGTGTCATGTAACCATATAGTATCTCATTGTAAACAAATCATTACATGCACAC
ATTCCCACCAATGCATGCTGGACTTCCTGACAAAGTACAACATGCTCACCTGCC
AACACTTGCTTGTTAAGACCTGTCAGTGGACTTGACCACTGATGGGTTATAGCTT
GCATTT

>P_3rU_cluster_231__C_5fU_2S8

ATTCAGCCTCAGACACTTTCCAGCTGTGTGACCCTGGGCAAGTCAGTTAACCCC
GTCTGCCTCAGTTTCTCCATCCATAAAATGAGCTGGAGAAAGAAATGGCAAACC
ACTCCAGGATCTTTGCCAAGAAATCCCCAAATGGGG

>P_3rU_S8.36760__C_5fU_S8.958388

CATATCAAATGACTTGTCCACAGTCACACAGC

>P_3rU_S8.36760__C_5rU_4S8

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>P_3rU_cluster_446_cluster_937__P_5rU_3S8_C_5rU_S8.15867

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GTATATGAATGTAATGGAATACTATC GTGCTATAAGAAAGA

>P_3rS_8S3_S8.47689__S_5rU_cluster_541_C_5rU_S8.50529

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TC

>P_3rU_4S8_P_3fU_S8.70215__C_5rU_S8.1022487

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>P_3rU_6s3_P_3fS_3S2_2S9_S8.5461_S5.1115_2S4_3S3_S_3fS_cluster_29_cluster_977_cluster_4_cluster_n1_cluster_28__S_5rS_clsuter_cp1_C_5fU_2S7_P_5fU_cluster_38_C_5rS_S7.10526_2S10_2S3_S4.17612_S5.4886_S9.1880_S1.11056_2S8_S2.123759

GAACCCACAAAAAGACAGAATGAAACAAATATCCAGCCCAATACAGCCTGGATG
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AGCTG

>P_3rU_6s3_P_3fS_3S2_2S9_S8.5461_S5.1115_2S4_3S3_S_3fS_cluster_29_cluster_977_cluster_4_cluster_n1_cluster_28__S_5rU_S20.31638_C_5rU_S3.50689

GAACCCACAAAAAGACAGAATGAAACAAATATCCAGCCCAATACAGCCTGGATG
CTTGATCGGA

>P_3rS_cluster_31_cluster_252_P_3fU_3S2_S_3rS_2S17_S15.6385_S_3fS_cluster_68__S_5rU_S18.27629_C_5fU_S2.1235481_C_5rU_S2.466916

GACAGCATTTTCCATCCTGAGGCCTCTGGAGATGTCTTAGATCCTTGTATTGCTG
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>P_3rS_S5.47458_6S8__S_5rS_cluster_43_C_5fU_S8.854973

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GCTGATGAGTAGCAGAAGCAAGAAGAGAAACAAAATCTTCTGATTCCCAGGTTC
CTGCCAC

>P_3rU_cluster_231__C_5rU_S8.503819

ATTCAGCCTCAGACACTTTCCAGCTGTGTGACCCTGGGCAAGTCAGTTAACCCC
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>P_3rU_S8.12674_S8.39898__C_5rU_S8.911698

GACACCTCGGTGTGTCTCAGTTTCCTCATCTATAAAATGAGCTGGAGAAGGAAA
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>P_5fU_cluster_54.maffT3s8_C_5fU_S8.30495_S8.561570__P_3fU_S8.88040_S8.3563

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GCCACT

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CCACT

>P_5fU_S2.26104_C_5fU_S2.82531__P_3fU_S2.2953

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GCATATGTGATG

>C_5fU_S2.410158__S_3fU_S18.5221_P_3fU_S2.310

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CATCAGTG

>C_5fU_S8.589094__C_3fU_S8.917292_P_3fU_S8.49463

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>C_5fU_S3.480683__P_3fU_S3.28583_S8.82168_S3.90397_S3.64168_S_3fU_S20
.10443

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>C_5fU_S8.92543_S8.302794__P_3fU_S8.42372

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AGGGTCATACTAGGAAGTGTCTGAGGCCAGATTTGAATCCAAGAAGATAAG
TCCTCCTGACTCCGGGTTTGGCAGTCTGTCCACTATGACA

>C_5fU_S3.102282_S3.954929__S_3fU_cluster_195_P_3fU_S3.39389

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>C_5fU_S8.242952_S8.249062__S_3fU_S17.14523_C_3fU_S8.70423_C_3rU_cluster_339

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TCACTTACCTCGGTAGTGTACTA

>P_5rU_cluster_19_C_5rU_S8.10722__S_3fU_S17.14523_C_3fU_S8.70423_C_3rU_cluster_339

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>S_5rU_cluster_645.maffT2s20_C_5rU_S3.16848__S_3fU_S20.29912

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>C_5rU_S8.108028__P_3fU_S8.380

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>C_5rU_S8.503819__P_3fU_S8.42372

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