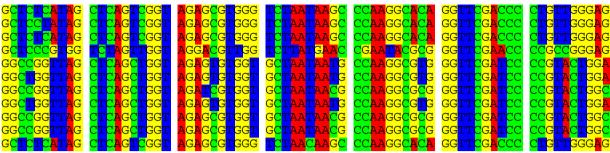
BayalaiB08-376_Baya_023_15_I GCTCTCATAG CTCAGTCGGT AGAGCGTGGG TCTAATAAGC CCAAGGCACA BayalaiB08-376_Baya_065_30_I GCTCTCATAG CTCAGTCGGT AGAGCGTGGG TCTAATAAGC CCAAGGCACA	GGTTCGACCC CTGTTGGGAG
BayalaiB08-376_Baya_065_30_I GCTCTCATAG CTCAGTCGGT AGAGCGTGGG TCTAATAAGC CCAAGGCACA	CCTTCCACC CTCTTCCCAC
	GGIICG CIGIIGGG
BayalaiB08-376_Baya_237_58_I GCTCTGTGG TCTAGTTGGT AGGACGCTGG TCTTATGAAC CGCATACGCCG	GGTTCGAGCC CCGCCAGGAG
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CfasciculataCfCl_CfaC1_18_58_I GCTCTCATAG CTCAGTCGGT AGAGCCTGGG TCTAATAAGC CCAAGGCACA	GGTTCGACCC CTGTTGGGAG
CfasciculataCfCl_CfaC1_25_73_I GCTCTCATAG CTCAGTCGGT AGAGCCTGGG TCTAATAAGC CCAAGGCACA	GGTTCGACCC CTGTTGGGAG
CfasciculataCfCl_CfaC1_25_75_I GCTCTCATAG CTCAGTCGGT AGAGCGTGGG TCTAATAAGC CCAAGGC <mark>ACA</mark>	GGTTCGACCC CTGTTGGGGTG
CfasciculataCfCl_CfaC1_29_114_I	GGTTCGACCC CTGTTGGGAG
EmonterogeiiLV88_EmoLV88_23_45_I GCTCCCGTGG TCTAGTTGGT AGGACGCTGG TCTTATGAAC CGGATACGCG	GGTTCGAGCC CCGCCGGGAG
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LbraziliensisMHOMBR75M2903_LbrM2903_23_52_I GCTCCCGTGG TCTAGTTGGT AGGACGCTGG TCTTATGAAC CGGATACGCG	GGTTCGAGCC CCGCCGGGAG
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	GGTTCGAGCC CCGCCGGGAG
LdonovaniBHU1220_CM002164.1_59_I GCTCTCATAG CTCAGTCGGT AGAGGCGTGGG TCTAATAAGC CCAAGGCACA	CCTTCCACCC CTGTTGGGAG
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LenriettiiLEM3045_LenLEM3045_23_42_I GCTCCCGTGG TCTAGTTGGT AGGACGCTGG TCTTATGAAC CGGATGCGCG	GGTTCGAGCC CCGCCGGGAG
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LinfantumJPCM5_LinJ.24_57_I GCTCTCATAG CTCAGTCGGT AGAGCGTGGG TCTAATAAGC CCAAGGCACA LinfantumJPCM5_LinJ.34_85_I GCTCTCATAG CTCAGTCGGT AGAGCGTGGG TCTAATAAGC CCAAGGCACA	GGTTCGACCC CTGTTGGGAG
	GGTTCGAGCC CCGCCGGGAG
LmajorFriedlin_LmjF.24_64_I GCTCTCATAG CTCAGTCGGT AGAGCGTGGG TCTAATAAGC CCAAGGCACA	GGTTCGACCC CTGTTGGGAG

LmajorFriedlin_LmjF.24_65_I	GCTCTCATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CCAA <mark>GG</mark> CACA	GGTTCGACCC	CTGTTGGGAG
LmajorFriedlin_LmjF.34_89_I	GCTCTCATAG	CTCAGTCGGT	AGAG <mark>CGTGGG</mark>	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GGTTCGACCC	CTGTTGGGAG
LmajorLV39c5_LmjLV39_23_48_I	GCTCCCGTGG	TCTAGTTGGT	AGGACGCTGG	TCTTATGAAC	CGGATGCGCG	GGTTCGAGCC	CCGCCGGGAG
LmajorLV39c5_LmjLV39_24_62_I	GCTCTCATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
LmajorLV39c5_LmjLV39_24_63_I	GCTCTCATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CCAA <mark>GG</mark> CACA	GGTTCGACCC	CTGTTGGGAG
LmajorLV39c5_LmjLV39_34_89_I	GCTCTCATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CCAA <mark>GG</mark> CACA	GGTTCGACCC	CTGTTGGGAG
LmajorSD75_LmjSD75_23_50_I	GCTCCCGTGG	TCTAGTTGGT	AGGACGCTGG	TCTTATGAAC	<mark>CGGATGC</mark> GCG		CCGCCGGGAG
LmajorSD75_LmjSD75_24_64_I	GCTCTCATAG	CTCAGTCGGT	AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
LmajorSD75_LmjSD75_24_65_I	GCTCTCATAG	CTCAGTCGGT	AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTT CGA CCC	CTGTTGGGAG
LmajorSD75_LmjSD75_34_89_I	GCTCTCATAG	CTCAGTCGGT	AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
LmexicanaMHOMGT2001U1103_LmxM.23_67_I	GCTCCCGTGG	TCTAGTTGGT	AGGACGCTGG	TCTTATGAAC	CGGATGCGCG		CCCCCCCCC
LmexicanaMHOMGT2001U1103_LmxM.24_81_I	GCTCTCATAG	CTCACTCGCT	AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
LmexicanaMHOMGT2001U1103_LmxM.24_82_I	GCTCTCATAG	CTCAGTCGGT	AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
LmexicanaMHOMGT2001U1103_LmxM.33_105_I	CCTCTCATAG	CTCACTCGGT	AGAGGGTGGG	TCTAATAAGC		GGTTCGACCC	CTCTTCCCAC
LpanamensisMHOMCOL81L13_LpaL13_20.1_39_I	CCTCTCATAG	CTCACTCCCT	AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTCTTCCCAC
LpanamensisMHOMCOL81L13_LpaL13_23_47_I	COTCCCCTCC	TCTACTTCCT	AGA GCCTCC	TCTTATGAAC	CGGATACGCG		CCCCCCCCC
LpanamensisMHOMCOL81L13_LpaL13_24_61_I	CCTCTCATAG	CTCACTCCCT	A GA GGG TGGG	TCTAATAAGC	CCAAGGCACA	CCTTCCACCC	CTCTTCCCAC
LpanamensisMHOMCOL81L13_LPAL13_SCAF000321_99_I	COTOTOATAC	CTCACTCCCT	ACACCCTCCC	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTCTTCCCAC
LpanamensisMHOMPA94PSC1_CP009389_36_I	COTOTOATAC	CTCACTCCCT	ACACCCTCCC	TOTANTANCO	CCAAGGCACA CCAAGGCACA	GGTT CGACCC	CTCTTCCCAC
LpanamensisMHOMPA94PSC1_CP009392_45_I	COTTCCCCTCC	TOTACTTCCT	ACCA CCCTCC	TCTTATGAAC	CGGATACGCG		CCCCCCCCAC
<u> </u>	COTTOTOTO	CTCACTCCCT	AGGACGCT GGG	TCTAATAAGC			CUGCUGGGAG
LpanamensisMHOMPA94PSC1_CP009393_59_I LpyrrhocorisH10_LpyrH10_03_14_I	COTTOCOCOTTOC	TOTACTOCCA	AGAGGGTGG		CCAAGGCACA	GGTT CGA CCC	CIGITIGGGAG
	COTTOTOTO	CTCACTCCCT	AGGACGCTGG	TCTTATGAAC TCTAATAAGC	CGGATACGCG CCAAGGCACA	GGTTCGAGCC	CTCTTTCCCAC
LpyrrhocorisH10_LpyrH10_05_30_I	COLCUCATAG	CICAGICGGI	AGAGGGGGGG			GGTTCGACCC	CIGIIIGGGAG
LpyrrhocorisH10_LpyrH10_11_71_I	CONCUCATAG	CICAGICGGI	AGAGGGGGGG	TCTAATAAGC	CC <mark>AAGGCA</mark> CA CCAAGGCACA	GGTTCGACCC	CTGTT GGGAG
LpyrrhocorisH10_LpyrH10_11_73_I	GCTCTCATAG	CTCAGTCGGT	AGAGGGTGGG	TCTAATAAGC		GGTTCGACCC	CTGTTGGGAG
LpyrrhocorisH10_LpyrH10_25_122_I	GCTCTCATAG	CTCAGTCGGT	AGAGGGTGGG	TCTAATAAGC		GGTT CGA CCC	CTGTTGGGAG
LseymouriATCC30220_Lsey_0029_8_I	GCTCTCATAG	CTCAGTCGGT	AGAGGGTGGG	TCTAATAAGC		GG <mark>TT</mark> CGACCC	CTGTTGGGAG
LseymouriATCC30220_Lsey_0029_10_I	GCTCTCATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GG <mark>TT</mark> CGACCC	CTGTTGGGAG
LseymouriATCC30220_Lsey_0039_13_I	GCTCTCATAG	CTCAGTCGGT	AGAGGGTGGG	TCTAATAAGC	CCAA <mark>GGCA</mark> CA	GGTT CGA CCC	CTGTTGGGAG
LseymouriATCC30220_Lsey_0053_19_I	GCTCCTGTGG	TCTAGTCGGT	AGGACGCTGG	TCTTATGAAC	CGGA <mark>TA</mark> CGCG		CCGCCAGGAG
LseymouriATCC30220_Lsey_0130_55_I	GCTCTCATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CCAA <mark>GG</mark> CACA	GGTTCGACCC	CTGTTGGGAG
LspMARLEM2494_LMARLEM2494_23_39_I	GCTCCCGTGG	TCTAGTTGGT	AGGACGCTGG	TCTTATGAAC	CGGATGCGCG		CCGCCGGGAG
LspMARLEM2494_LMARLEM2494_24_53_I	GCTCTCATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA		CTGTTGGGAG
LspMARLEM2494_LMARLEM2494_34_75_I	GCTCTCATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GG <mark>TT</mark> CGACCC	CTGTTGGGAG
LtarentolaeParrotTarII_LtaP23_37_I	GCTCCCGTGG	TCTAGTTGGT	AGGACGCTGG	TCTTATGAAC	CGG <mark>AT</mark> GCGCG	GG <mark>TT</mark> CGAGCC	CCGCCGGGAG
LtarentolaeParrotTarII_LtaP24_50_I	GCTCTCATAG	CTCAGTCGGT	AGAG <mark>CGT</mark> GGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GGTTCGACCC	CTGTTGGGAG
LtarentolaeParrotTarII_LtaP24_51_I	GCTCTCATAG	CTCAGTCGGT	AGAG <mark>CGTGGG</mark>	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GGTTCGACCC	CTGTTGGGAG
LtarentolaeParrotTarII_LtaP34_79_I	GCTCTCATAG	CTCAGTCGGT	AGAG <mark>CGT</mark> GGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GGTTCGACCC	CTGTTGGGAG
LtropicaL590_LtrL590_23_49_I	GCTCCCGTGG	T <mark>CTA</mark> GTTGGT	AGGACGCTGG	T <mark>CTTA</mark> TGAAC	CGG <mark>AT</mark> GCGCG		CCGCCGGGAG
LtropicaL590_LtrL590_24_63_I	GCTCTCATAG	CTCAGTCGGT	AGAG <mark>CGTGGG</mark>	TCTAATAAGC	CCAA <mark>GGCA</mark> CA	GG <mark>TT</mark> CGACCC	CTGTTGGGAG
LtropicaL590_LtrL590_24_64_I	GCTCTCATAG	CTCAGTCGGT	AGAG <mark>CGT</mark> GGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GG <mark>TT</mark> CGACCC	CTGTTGGGAG
LtropicaL590_LtrL590_34_93_I	GCTCTCATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GG <mark>TT</mark> CGACCC	CTGTTGGGAG
LturanicaLEM423_LtuLEM423_23_42_I	GCTCCCGTGG	TCTAGTTGGT	AGGACGCTGG	TCTTATGAAC			CCGCCGGGAG
LturanicaLEM423_LtuLEM423_24_57_I	GCTCTCATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CCAA <mark>GG</mark> CACA	GGTTCGACCC	CTGTTGGGAG
LturanicaLEM423_LtuLEM423_24_58_I	GCTCTCATAG	CTCAGTCGGT	AGAG <mark>CGTGGG</mark>	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GGTTCGACCC	CTGTTGGGAG
LturanicaLEM423_LTULEM423_SCAF000129_98_I	GCTCTCATAG	CTCAGTCGGT	AGAG <mark>CGTGGG</mark>	TCTAATAAGC	CCAA <mark>GGCA</mark> CA	GGTTCGACCC	CTGTTGGGAG
LturanicaLEM423_LTULEM423_SCAF000130_103_I	GCTCTCATAG	CTCAGTCGGT	AGAG <mark>CGTGGG</mark>	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
PconfusumCUL13_Pcon_scaffold00074_88_I	GCTCTCATAG GCTCTCATAG GCTCCTGTGG	TCTAGTTGGT	AGG <mark>ACGTT</mark> GG	TCTTATGAGC	CGAACGCGCG	GG <mark>TTCGA</mark> GCC	CCGCCAGGAG
TbruceigambienseDAL972_Tbg972_08_28_I	GCTCCTATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TbruceigambienseDAL972_Tbg972_08_33_I	GCTCCCGTGG	TCTAGTTGGT	AGGACGTTGG	TCTTATGAAC	CGAA <mark>TA</mark> CGCG	GG <mark>TT</mark> CGAACC	
TbruceiLister427_Tb427_08_v4_32_I	GCTCCTATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC			CTGTTGGGAG
TbruceiLister427_Tb427_08_v4_36_I	GCTCCTATAG	C <mark>TCA</mark> GTCGGT	AGAG <mark>CG</mark> TGGG	TCTAATAAGC	C <mark>CAAGG</mark> CACA CCAAGGCACA	GG <mark>TT</mark> CGACCC	CTGTTGGGAG
TbruceiLister427_Tb427_08_v4_40_I	GCTCCCGTGG	TCTAGTTGGT	AGGACGTTGG	TCTTATGAAC	CGAATACGCG	GGTTCGAACC	CCGCCGGGAG
TbruceiTREU927_Tb927_08_v5.1_31_I	GCTCCTATAG	CTCAGTCGGT	AGAGCGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TbruceiTREU927_Tb927_08_v5.1_35_I	GCTCCTATAG	CTCAGTCGGT	AGAG <mark>CGT</mark> GGG	TCTAATAAGC	CCAAGGCACA	<mark>GG</mark> TTC <mark>GA</mark> CCC GGTTCGACCC	CTGTTGGGAG
					30 31		

TbruceiTREU927_Tb927_08_v5.1_39_I	GCTCCCGTGG TCTAGTT	TGGT AGGACGTTGG	TCTTATGAAC	CGAATACGCG	GGTTCGAACC	CCGCCGGGAG
TcongolenseIL3000_T.congo_bin_contig_1714_25_I	GCTCCTATAG CTCAGTO	CGGT AGAGCGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcongolenseIL3000_T.congo.pschr.8_78_I	GCTCCTATAG CTCAGTO	CGGT AGAGCGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcongolenseIL3000_T.congo.pschr.8_81_I	GCTCCTATAG CTCAGTC	CGGT AGAGCGTGGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GG <mark>TT</mark> CG <mark>A</mark> CCC	CTGTTGGGAG
TcruziCLBrener_Tcruzi_11019_2_I	GCGAGCGTAG CTCAGCT	GGT AGAGCACACG	ACTGATAATC	G <mark>T</mark> GAGGCGAT	GGTTCAAGTC	CATTCGTTCG
TcruziCLBrenerEsmeraldo-like_TcChr22-S_9_I	GCTCCTGTGG TCTAGT	GGT AGGACGTTGG	TCTTATGAGC	CGAA <mark>TA</mark> CGCG	GG <mark>TT</mark> CGAGCC	CCGCCAGGAG
TcruziCLBrenerEsmeraldo-like_TcChr26-S_23_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CCAA <mark>GG</mark> CACA	GGTTCGACCC	CTGTTGGGAG
TcruziCLBrenerEsmeraldo-like_TcChr41-S_57_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziCLBrenerNon-Esmeraldo-like_TcChr22-P_10_I	GCTCCTGTGG TCTAGT	GGT AGGACGTTGG	TCTTATGAGC	CGAA <mark>TA</mark> CGCG	GGTTCGAGCC	CCGCCAGGAG
TcruziCLBrenerNon-Esmeraldo-like_TcChr26-P_25_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziCLBrenerNon-Esmeraldo-like TcChr41-P 54 I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruzicruziDm28c_MBSY01000058_2_I	GCTCTCATAG CTCAGTC	CGT AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruzicruziDm28c MBSY01000093 5 I	GCTCCTGTGG TCTAGT	GGT AGGACGTTGG	TCTTATGAGC	CGAA <mark>TA</mark> CGCG	GG <mark>TT</mark> CG <mark>A</mark> GCC	CCGCCAGGAG
TcruzicruziDm28c_MBSY01000240_26_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
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TcruzicruziDm28c_MBSY01000739_100_I	GCTCCTGTGG TCTAGT	GGT AGGACGTTGG	TCTTATGAGC	CGAA <mark>TA</mark> CGCG	GG <mark>TTCGA</mark> GCC	CCGCCAGGAG
TcruziDm28c_AYLP01000078_19_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
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TcruziEsmeraldo ANOX01013115.1 21 I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziEsmeraldo_ANOX01013135.1_23_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziEsmeraldo_KB205264.1_40_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziEsmeraldo KB205393.1 57 I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziEsmeraldo_KB205774.1_79_I	GCTCCTGTGG TCTAGT	GGT AGGACGTTGG	TCTTATGAGC	CGAA <mark>TA</mark> CGCG	GGTTCGAGCC	CCGCCAGGAG
TcruziJRc14_AODP01008205.1_6_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziJRc14 AODP01010036.1 8 I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziJRc14_KB222969.1_64_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziJRc14 KB223000.1 71 I	GCTCCTGTGG TCTAGT	GGT AGGACGTTGG	TCTTATGAGC	CGAA <mark>TA</mark> CGCG	GG <mark>TT</mark> CGAGCC	CCGCCAGGAG
TcruzimarinkelleiB7_TcMARK_CONTIG_2326_34_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruzimarinkelleiB7_TcMARK_CONTIG_3490_45_I	GCTCTCATAG CTCAGTC	GGT AGAGGGTGGG	TCTAATAAGC	CCAA <mark>GG</mark> CACA	GGTTCGACCC	CTGTTGGGAG
TcruzimarinkelleiB7_TcMARK_CONTIG_722_58_I	GCTCCTGTGG TCTAGT	GGT AGGACGTTGG	TCTTATGAGC	CGAA <mark>TA</mark> CGCG	GG <mark>TT</mark> CGAGCC	CCGCCAGGAG
TcruziSylvioX10-1-2012_ADWP02000205_1_I	GCTCCTGTGG TCTAGTT	GGT AGGACGTTGG	TCTTATGAGC	CGAATACGCG	GG <mark>TT</mark> CGAGCC	CCGCCAGGAG
TcruziSylvioX10-1-2012_ADWP02006284_21_I	GCTCTCATAG CTCAGTO	GGT AGAGCGTGGG	TCTAATAAGC	CCAA <mark>GG</mark> CACA	GGTTCGACCC	CTGTTGGGAG
TcruziSylvioX10-1-2012_ADWP02006285_24_I	GCTCTCATAG CTCAGTO	GGT AGAGCGTGGG	TCTAATAAGC	CCAA <mark>GG</mark> CACA	GGTTCGACCC	CTGTTGGGAG
TcruziSylvioX10-1-2012_ADWP02016913_64_I	GCTCTCATAG CTCAGTO	GGT AGAGCGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziSylvioX10-1_TcX10_chr21_29_I	GCTCTCATAG CTCAGTO	CGGT AGAGCGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTG-GAG
TcruziSylvioX10-1_TcX10_chr22_30_I	GCTCTCATAG CTCAGTC	GGT AGAGCGTGGG	TCTAATAAGC	CCAAGGCACA	GG <mark>TT</mark> CGACCC	CTGTTGGGAG
TcruziSylvioX10-1_TcX10_chr31_39_I	GCTCTCATAG CTCAGTO	GGT AGAGCGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziSylvioX10-1_TcX10_chr37_45_I	GCTCCTGTGG TCTAGTT	GGT AGGACGTTGG	TCTTATGAGC	CGAA <mark>TA</mark> CGCG	GG <mark>TT</mark> CG <mark>A</mark> G-C	CCGCCAGGAG
TcruziTulacl2_AQHO01024116.1_36_I	GCTCTCATAG CTCAGTO	GGT AGAGCGTGGG	TCTAATAAGC	CCAA <mark>GG</mark> CACA	GGTTCGACCC	CTGTTGGGAG
TcruziTulacl2_AQHO01027870.1_53_I	GCTCTCATAG CTCAGTO	GGT AGAGCGTGGG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TcruziTulacl2_AQH001034196.1_67_I	GCTCCTGTGG TCTAGTT	GGT AGGACGTTGG	TCTTATGAGC	CGAA <mark>TA</mark> CGCG	GGTTCGAGCC	CCGCCAGGAG
TcruziTulacl2_KB851533.1_117_I	GCTCCTGTGG TCTAGTT	r <mark>ggt aggacgtt</mark> gg	TCTTATGAGC	CGAA <mark>TA</mark> CGCG	GGTTCGAGCC	CCGCCAGGAG
TcruziTulacl2_KB852025.1_144_I		CGGT AGAGCGTGGG	TCTAATAAGC		GGTTCGACCC	CTGTTGGGAG
TcruziTulacl2_KB852086.1_146_I	GCTCTCATAG CTCAGTC	CGG <mark>T AGAGCGTGGG</mark>	TCTAATAAGC	CCAA <mark>GGCA</mark> CA	GGTTCGACCC	CTGTTGGGAG
TevansiSTIB805_STIB805_Chr08_31_I	GCTCCTATAG CTCAGTO	CGG <mark>T AGAGCGTGGG</mark>	TCTAATAAGC	CC <mark>AA</mark> GGCACA CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TevansiSTIB805_STIB805_Chr08_35_I	G <mark>CTCCTATA</mark> G CTCAGTC	CGG <mark>T AGAGCGTGGG</mark>	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GGTTCGACCC	CTGTTGGGAG
TevansiSTIB805_STIB805_Chr08_39_I	GCTCCCGTGG TCTAGT	GGT AGGACGTTGG	TCTTATGAAC	CGAA <mark>TA</mark> CGCG	GGTTCGAACC	CCGCCGGGAG
TgrayiANR4_JMRU01001200_74_I		CGG <mark>T AGAGCGTG</mark> GG	TCTAATAAGC	CCAAGGCACA	GGTTCGACCC	CTGTTGGGAG
TgrayiANR4_JMRU01001200_78_I	GCTCTCATAG CTCAGTC	CGG <mark>T AGAGCGT</mark> GGG	TCTAATAAGC	CC <mark>AA</mark> GGCACA	GGTTCGACCC	CTGTTGGGAG
TrangeliSC58_AUPL01002787_6_I	GCTCTCATAG CTCAGTC	<mark>CGGT AGAGCGTGGG</mark>	TCTAATAAGC	CC <mark>AA</mark> GGCACA CCAAGGCACA	GG <mark>TT</mark> CGACCC	CTGTTGGGAG
TtheileriEdinburgh_NBCO01000013_65_I	GCTCCCGTGG TCTAGT	<mark>rggt a</mark> gg <mark>acgtt</mark> gg	TCTTATGAAC	CGAATACGCG	GG <mark>TT</mark> CGAGCC	CCGCCGGGAG
TtheileriEdinburgh_NBCO01000015_71_I	GCTCCCATAG CTCAGTC	C <mark>GGT AGAGCGTG</mark> GG	TCTAATAAGC	CC <mark>AAGGCA</mark> CA	GG <mark>TT</mark> CG <mark>A</mark> CCC	CTGTTGGGAG
TtheileriEdinburgh_NBCO01000015_72_I	GCTCC <mark>TATA</mark> G CTCAGTC	<mark>CGGT AGAGCGT</mark> GGG		CC <mark>AA</mark> GGCACA	GG <mark>TT</mark> CG <mark>A</mark> CCC	CTGTTGGGAG
TtheileriEdinburgh_NBCO01000021_87_I	GCTCCTATAG CTCAGTC	CGG <mark>T AGAGCGT</mark> GGG	TCTAATAAGC	CC <mark>AA</mark> GGC <mark>A</mark> CA	GG <mark>TT</mark> CG <mark>A</mark> CCC	CTGTTGGGAG





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LmajorLV39c5_LmjLV39_34_89_I C
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