# $Start\ of\ Alignment$

		10	20	30	40	50	60 TGGGGĊCCGCT
75_Mmul	NNNNNNN	INNNNNNN	INNNNNNNNN	N N N A	TGGCAÁGTTAG	CTGCGÁGTTCT	GGGGCCCGCT
12_Ggor	NNNNNNN	INNNNNNN	INNNNNNNN	NNNTGGGAA'	TGAGGGTGCA	GTCGGAGCTTC	GAACATCTCTC
36_Nleu	NNNNNNN	INNNNNNN	INNNNNNNN	NNNTGGGAA	TGAGGGTGCA	GTCGGAGCTTC	GAACATCTCTC
47_Tsyr							CCCGCCCCCCT
06_Mmur	NNNNNNN	INNNNNNN	INNNNNNNNN	NNNC	TGGCAAGTTAT	<b>FAGCGCGTTAT</b>	CAGGCCCGCT
32_Pabe	NNNNNNN	INNNNNNN	INNNNNNNN	NNNTGAGAT	GGCAAGTTACA	AGCGCCTTCT	CCGGCCCGCTG
23_Cjac	NNNNNNN	INNNNNNN	INNNNNNNNN	NNNTGGGAA'	TGAGGGTGCA	GTCGGAGCTTC	GAACACCTCTC
32_Hsap	NNNNNNN	INNNNNNN	INNNNNNNNN	NNNTGGGAA'	TGAGGGTGCA	GTCGGAGCTTC	GAACATCTCTC
59_Ptro	NNNNNNN	INNNNNNN	INNNNNNNNN	N N N A	TGGCAAGTTAG	CAGCGGCTTCT	CCCGCCTGCT
95_0gar	NNNNNNN	INNNNNNN	INNNNNNNN	NNNTGGGAG	TGAGGATGCA	GTCGGAGCTTC	GAACAGCCCTC
Indel							

	70	80	90	100	110	120	130
75_Mmul	GCAGATTCGC	$CGG - \dot{-} C$	CCAGGÁCACCA	100 CAGCTGCNNNN	NNNNNNNNN	J N N N N N N N N N N	INNNN
12_Ggor	CGAGCTGCGG	CGGGAAG	<b>GCGGGGAAGGA</b>	TCGCAGCNNNN	NNNNNNNN	INNNNNNNNN	INNNN
36_Nleu	CGAGCTGCGG	CGGGAAG	GCGGGGAAGGA	TCGCAGCNNNN	NNNNNNNN	INNNNNNNNNN	INNNN
47_Tsyr				TAGCTGTNNNN			
06_Mmur				CAGCTGCNNNN			
32_Pabe				CAGCTGCNNNN			
23_Cjac	CGAGCTGCGG	<b>CGGGAA</b>	GCCGGGAAGGA	TCGCAGANNNN	NNNNNNNN	INNNNNNNNNN	INNNN
32_Hsap				TCGCAGCNNNN			
59_Ptro	GGAGATTCGC	TACGGGC	CCAGGACACCG	CAGCTGCNNNN	NNNNNNNN	INNNNNNNNNN	INNNN
95_0gar	CGAACTGCGG	CAGGAAC	CCGGAGAAGCT	GCTCAGCNNNN	NNNNNNNN	INNNNNNNNNN	NNNN

# Indel

	140 NNNNNNNNNNNNNNNNNN	150	160	170	180	190
75_Mmul	NNNNNNNNNNNNNNN	INNNNNNNNN	INNNNNNNNNN	NNNNNNNNN	NNNNNNNNN	NNNNNNNNN
12_Ggor	NNNNNNNNNNNNNN	INNNNNNNN	INNNNNNNNN	NNNNNNNN	NNNNNNNN	NNNNNNNNN
36_Nleu	NNNNNNNNNNNNNN	INNNNNNNN	INNNNNNNNNN	NNNNNNNNN	NNNNNNNNN	NNNNNNNNN
47_Tsyr	NNNNNNNNNNNNNN	INNNNNNNN	INNNNNNNNNN	NNNNNNNNN	NNNNNNNNN	NNNNNNNNN
06_Mmur	NNNNNNNNNNNNNN	INNNNNNNN	INNNNNNNNNN	NNNNNNNNN	NNNNNNNNN	NNNNNNNNN
32_Pabe	NNNNNNNNNNNNN	INNNNNNNN	INNNNNNNNNN	NNNNNNNN	NNNNNNNNN	NNNNNNNNN
23_Cjac	NNNNNNNNNNNNNN	INNNNNNNN	INNNNNNNNNN	NNNNNNNNN	NNNNNNNNN	NNNNNNNNN
32_Hsap	NNNNNNNNNNNNNN	INNNNNNNN	INNNNNNNNNN	NNNNNNNNN	NNNNNNNNN	NNNNNNNNN
59_Ptro	NNNNNNNNNNNNNN	INNNNNNNN	INNNNNNNNNN	NNNNNNNNN	NNNNNNNNN	NNNNNNNNN
95_0gar	NNNNNNNNNNNNN	INNNNNNNN	INNNNNNNNNN	NNNNNNNN	NNNNNNNNN	NNNNNNNNN

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220
                                       230
                                                240
       NNNNNNNCCCCAATTCGCTTTCTTTCCGCAGCCGCCGCTGCCCCGACCCCGGATCTGCATGTG
75 Mmul
       NNNNNNNCCCGGAGCCAACTTGTTTCCG---CCGCCGCTGCCCCGACCCCGGATCTGCATGTG
36_Nleu NNNNNNNCCCGGAGCCAACTTGTTTCCG---CCGCCGCTGCCCCGACCCCGGATCTGCATGTG
O6_Mmur NNNNNNNCCCCAATCCGCTTTCTCTCCGCAGCCGCCGCTGCCCCGACCCCGGATCTGCATGTG
32_Pabe NNNNNNNCCCCAATTCGCTTTCTTTCCGCAGCCGCCGCTGCCCCGACCCCGGATCTGCATGTG
       NNNNNNNCGTGGAGCCAACTTGTTCCCG---CCGCCGCTGCCCCGGCCCCAGATCTGCATGTG
23_Cjac
       NNNNNNNNCCCGGAGCCAACTTGTTTCCG---CCGCCGCTGCCCCGACCCCGGATCTGCATGTG
NNNNNNNNCCCCAATTCGCTTTCTTTCCGCAGCCGCCGCCCGGCCCCGGATCTGCATGTG
32_Hsap
59_Ptro
95_Ogar NNNNNNNCCTGGAACCAACTTGTTCCCG---CCGCCGCTGCCTCGACCCCGAATCTGCATGTG
                                    Indel
                         280
                                  290
                                           300
                                                     310
75_Mmul GAAGTACCTGGACATCCATTCCATGCACCAGCTGGAGAAGACTACCAGTGCTGAGGAGATGAGGC
12_Ggor GAAGTACCTGGACGTCCATTCCATGCACCAGCTGGAGAAGACCACCAATGCTGAGGAGATGAGGG
36_Nleu GAAGTACCTGGACATCCATTCCATGCACCAGCTGGAGAAGACCACCAATGCTGAGGAGATGAGGG
O6_Mmur GAAGTACCTGGACATCCATTCCATGCACCGGCTGGAGAAGACGCCAACGTTGAGGAGATGAGGG
32_Pabe GAAGTACCTGGACATCCATTCCATGCACCAGCTGGAGAAGACCACCAATGCTGAGGAGATGAGGG
23_Cjac GAAGTACCTGGACATCCACTCCATGCACCAGCTGGAGAAGACCACCAATGCTGAGGAGATGAGGG
       GAAGTACCTGGACGTCCATTCCATGCACCAGCTGGAGAGACCACCAATGCT---GAGATGAGGG
32_Hsap
      GAAGTACCTGGACGTCCATTCCATGCACCAGCTGGAGAAGACCACCAATGCTGAGGAGATGAGGG
59_Ptro
95_Ogar GAAGTACTTGGACATCCATTCTATGTACCGGCTGGAGAAGACCACCAACGTTGAGGAGATGAGGA
Indel
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360
                                               370
                   340
                            350
75_Mmul
       AGGTGTTGGCTGAGCTGCTGGAGCTAGGGTTCCTGAGCAGAGCCTGCGGGACGCCATCACCCTG
       AGGTGCTGGCTGAGCTGCTGGAGCTAGGGTGTCCTGAGCAGAGCCTGCGGGACGCCATCACCCTG
12_Ggor
36_Nleu AGGTGCTGGCTGAGCTGCTGGAGCTAGGGTGTCCTGAGCAGAACCTGCGAGACGCCATCACCCTG
O6_Mmur AGGTGCTGGCTGAGCTGTTAGAGCTAGGCCATCCTGAGCAGAACTTGCGGGATGCTATCACCCTG
32_Pabe AGGTGCTGGCTGAGCTGCTGGAGCTAGGGTGTCCTGAGCAGAGCCTGCGGGATGCCGTCACCCTG
      AGGTGCTGGCTGAGCTGTTGGAGCTAGGGCGTCCTGAGCAGAGCCTACGGGATGCCATCACCCTG
23_Cjac
      AGGTGCTGGCTGAGCTGCTGGAGCTAGGGTGTCCTGAGCAGAGCCTGAGGGACGCCATCACCCTG
32_Hsap
59_Ptro AGGTGCTGGCTGAGCTGCTGGAGCTAGGGTGTCCTGAGCAGAGCCTGCGGGACGCCATCACCCTG
95_Ogar AGGTGCTGGCTGAACTGTTGGAGCTAGGTTGTCCTGAGCAGAGCCTGCGAGATGCTATCACCCTG
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	400	410	420	430	440	450
75_Mmul	400 GACCTCTTCTGCCAC	CGCGCTCATC	TTCTGĊCGCCA	.GCAGĠGCTT(	CTCACŤGGAGC <i>i</i>	AGACTTCAGC
12_Ggor	GACCTCTTCTGCCAC	CGCGCTCATT	TTCTGCCGCCA	GCAGGGCTT	CTCACTGGAGC <i>I</i>	AGACGTCAGC
36_Nleu	GACCTCTTCTGCCAC	CGCGCTCATT	TTCTGCCGCCA	GCAGGGCTT	CTCACTGGAGC <i>I</i>	AGACATCAGC
47_Tsyr	NNNNNNNNNNNNN	INNNNNNNNN	NNNNNNNNNNN	NNNNNNNN	J N N N N N N N N N N N	INNNNNNNN
06_Mmur	GACCTCTTCTCCCAC	CGCACTCATC	TTCTGCCGCGA	GCAGGGCTT	CTCACTGGAGC <i>I</i>	AGACATCAAC
32_Pabe	GACCTCTTCTGCCAC	CGCGCTCATT	TTCTGCCGCCA	GCAGGGCTT	CTCACTGGAGCA	AGACGTCAGC
23_Cjac	GACCTCTTCTGCCAC	CATGCTCATC	TTCTGCCGCCA	GCAGGGCTT	CTCACTGGAGCA	AGACATCAGC
32_Hsap	GACCTCTTCTGCCAC	CGCGCTCATT	TTCTGCCGCCA	GCAGGGCTT	CTCACTGGAGCA	AGACGTCAGC
59_Ptro	GACCTCTTCTGCCAC	CGCGCTCATT	TTCTGCCGCCA	GCAGGGCTT	CTCACTGGAGCA	AGACGTCAGC
95_0gar	GACCTCTTTGCCCAT	GCACTCATG	TTCTGCCGCCA	ACAGGGCTT	CTCACTGGAGC <i>I</i>	AGACATCAAC

			480				
75_Mmul	AGCTTGTGCCC	TGCTCCAGGA	TCTTCACAAG	GCTTGTATTC	CAACCCCCTT	GGGCAACGTG	GAGĠ
12_Ggor	GGCTTGTGCCC	TGCTCCAGGA	TCTTCACAAG	GCTTGTATTC	CAACCCCCTT	GGGCAACGTG	GAGG
36_Nleu	GGCTTGTGCCC	TGCTCCAGGA	TCTTCACAAG	GCTTGTATTC	CAACCCCCTT	GGGCAACGTG	GAGG
47_Tsyr	NNNNNNNNN	NNNNNNNNN	NNNNNNNNN	NNNNNNNN	INNNNNNNNN	NNNNNNNNN	INNN
06_Mmur	GGCTTGTGCCA	TGCTCCAAGA	TCTTCACAAG	GCCTGTGTTC	CAACCCCCTT	GGGCAACGTG	GAGG
32_Pabe	GGCTTGTGCCC	TGCTCCAGGA	TCTTCACAAG	GCTTGTATTC	CAACCCCCTT	GGGCAACGTG	GAGG
23_Cjac	AGCTTGTGCCA						
32_Hsap	GGCTTGTGCCC	TGCTCCAGGA	TCTTCACAAG	GCTTGTATTC	CAACCCCCTT	GGGCAACGTG	GAGG
59_Ptro	AGCTTGTGCCC						
95_0gar	AGCTTGTGCCC	TGCTCCAAGA	TCTTCACAAG	GCCTGTGTTC	CAACCCCCTT	GGGCAACGTGC	AAGG

#### Indel

```
540
                                                      550
                                                                     560
                                                                                   570
           AGTGCTACCÁCTACTTCACCAGCGTTCTTTTTTGCCATGGAGTCAGGCGGCCTCCTTTCÁGCATCAGTGCTACCGCTACTTCACCAGTGTCCTTTTTTTGCCATGGAGTCAGGCGGCCTCCTTTCAGCATC
75_Mmul
12_Ggor
36_Nleu
           {\tt AGTGCTACCGCTACTTCACCAATGTCCTTTTTTGCCATGGAGTCAGGCGGCCTCCTTTCAGCATC}
           47_Tsyr
06_Mmur
           AATGCTACCGCTACTTTACCAGTGTTCTTTTTTGCCATGGAGTCAGGCGGCCCCCCTTCAGCATT
32_Pabe
           AGTGCTACCGCTACTTCACCAGTGTCCTTTTTTGCCATGGAGTCAGGCGGCCTCCTTTTAGCATC
           AGTGCTACCGCTACTTCACCAGTGTCCTTTTTTGCCATGGAGTCAGGCGGCCTCCTTTCAGCATC
AGTGCTACCGCTACTTCACCAGTGTCCTTTTTTGCCATGGAGTCAGGCGGCCTCCTTTCAGCATC
AGTGCTACCGCTACTTCACCAGTGTCCTTTTTTGCCATGGAGTCAGGCGGCCTCCTTTCAGCATC
AATGCTACCGCTACTTTACCAGTGTTCTTTTTTGCCATGGAGTCAGGCGGCCCCCCTTCAGCATT
23_Cjac
32_Hsap
59_Ptro
95_0gar
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	590	600	610	620	630	640	650
75_Mmul	GACCTCTTCA	A A G A Ġ G A A C <i>I</i>	ACTGCTGGCC	CTGGÁAGACT	ACGTGGTCAA	CACTTACTTCC	GCCÁ
12_Ggor	GACCTCTTCA	AAGAGGAACA	ACTGCTGGCC	TTGGAAGACT	ACGTGGTCAA	CACTTACTTCC	GCCA
36_Nleu	GACCTCTTCA	AAGAGGAACA	ACTGCTGGCC	TTGGAAGACT	ACGTGGTCAA	CACTTATTTCC	GCCA
47_Tsyr	GACCTCTTTGA	AGGAGGAAA	AGCTGCTGGCC	TTGGCAGACT	ATGTGGTCAA	CACCTACTTCC	GCCA
06_Mmur	GACCTATTTA	AGGAGGAACA	AGCTGCTGGCC	CTGGCAGACT	ATGTGGTCAA	CACCTACTTCC	GCCA
32_Pabe	GACCTCTTCA	AAGAGGAACA	AGCTGCTGGCC	TTGGAAGACT	ACGTGGTCAA	CACTTACTTTC	GCCA
23_Cjac						CACTTACTTCC	
32_Hsap						CACTTACTTCC	
59_Ptro	GACCTCTTCA	AAGAGCAACA	AGCTGCTGGCC	TTGGAAGACT	ACGTGGTCAA	CACTTACTTCC	GCCA
95_0gar	GACCTCTTTA	AAGAGGAACA	AGCTGCTGGCC	CTGGCAGACT	ATGTGGTCAA	CACCTATTTCC	GCCA
Ü							

	660	670	680	690	700	710
75_Mmul	CTTCAAACTĊTATAA	ATACĠTCTTC	ACACCCCAGG	TGCGĠCTGGA	TCTGTCTTTG	ACTTÁCATGG
12_Ggor	CTTCAAGCTCTATAA	ATACGTCTTC	ACACCCCAGG	TGCGGCTGGA	TCTGTCTTTG	ACTTACATGG
36_Nleu	CTTCAAGCTCTATAA	ATACGTCTTC	ACACCCCAGG	TGCGGCTGGA	TCTGTCTTTG	ACTTACATGG
47_Tsyr	CTTCAAGCTCTACAA	ATATGTCTTC	ACACCCCAGG	TGCGGCTGGA	TCTATCCTTG	ACTTACATAG
06_Mmur	CTTCAAGCTCTACAA	ATATGTCTTC	ACACCCCAGG	TACGGCTGGA	TCTATCTTTG	ACTTATATGG
32_Pabe	CTTCAAGCTCTATAA	ATACGTCTTC	ACACCCCAGG	TGCGGCTGGA	TCTGTCTTTG	ACTTACATGG
23_Cjac	CTTCAAGCTCTATAA	ATTCGTCTTC	ACATCCCAGG	TGCGGCTGGA	TCTATCTTTG	ACTTATATGG
32_Hsap	CTTCAAGCTCTATAA	ATACGTCTTC	ACACCCCAGG	TGCGGCTGGA	TCTGTCTTTG	ACTTACATGG
59_Ptro	CTTCAAGCTCTATAA	ATACGTCTTC	ACACCCCAGG	TGCGGCTGGA	TCTGTCTTTG	ACTTACATGG
95_0gar	CTTCAAGCTCTACAA	ATATGTCTTC	ACACCCCAGG	TGCGGTTGGA	TCTATCTTTG	ACTTACATGG

#### Indel

```
730
       740
          750
            760
               770
 75_Mmul
 12_Ggor
36_Nleu
 47_Tsyr
 06_Mmur
 32_Pabe
 23_Cjac
32_Hsap
 59_Ptro
95_0gar
 GGCTACAGTCACCCAAGCTCTGGCCAGAAGGTAAGATGGACAAAGAAGAAACCAAGGAAGTGGAG
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```
800
                                  810
                                             820
75_Mmul GAGCAGGCAGTTACGCCGCAG-----GAAGAGGAACTAGAGACAGTANNNNNNNNNNNCCCCCGCC
       GAGCAGGCAGTTACCCTGCAG-----AAAGAGGAACTAGAGACAGTGNNNNNNNNNGCCCCGCC
12_Ggor
36_NIeu GAGCAGGCAGTTACCCTGCAG-----GAAAAGGAACTAGAGACAGTGNNNNNNNNNGCCCCGCC
47_Tsyr GAGGAGACAGTCACTTCACAGGAGGAGGAGGAACTGGAGACAGTGNNNNNNNNNGCTCAGCC
O6_Mmur GAGCAGGCAGTCACCCCACGG-----GAAGAGGGGGCTGGAGACAGTGNNNNNNNNNGCCCAGCC
32_Pabe GAGCAGGCAGTTACCCCGCGG-----AAAGAGGAACTAGAGACGGTGNNNNNNNNGCCCCGCC
       GAGCATGCACTTACCCCCCAG-----GAAGGACTGGAGACAGTGNNNNNNNNNNNGCAGAGCC
23_Cjac
32_Hsap
       GAGCAGCCAGTTACCCCGCAG-----AAAGAGGAACTAGAGACAGTGNNNNNNNNNGCCCCACC
59_Ptro GAGCAGGCAGTTACCCCGCAG-----AAAGAGGAACTAGAGACAGTGNNNNNNNNNGCCCCGCC
95_Ogar GAGCAGGCAGTCACCCCACAC-----GAGGAGGAACTGGAAACGGTGNNNNNNNNNGCCCAGCC
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	850	860	870	880	890	900	910
75_Mmul	AGAGCCGGAGC	CAAĠCCACGT	CCACATCCTC	CGAĠCCTACA	TCAAGACCCA	AAATĠAACAAG	GAGĊ
12_Ggor	AGAGCCAGAGC	CAAGCCACGT	CCACGTCCTC	CGAGCCTACA	TCAAGACCCA	AAGTGAACAAA	GAGC
36_Nleu	AGAGCCAGAGC	CAAGCCACAT	CCACGTCCTC	CGAGCCTACA	TCAAGACCCA	AAGTGAACAAG	GAGC
47_Tsyr	TGAGCAAGAGC	CAAGCCAGAT	CCACATCCTC	CGAGCCTACA	TCAAGACCCA	AACTGAACCAG	GAGC
06_Mmur	AGAGCAAGAGC	CAAGCCAGAT	CCACATCCTC	CAAACGTATA	TCAAGACCCA	AACTGAACAAG	GAGC
32_Pabe	AGAGCCAGAGC	CAAGCCACGT	CCACATCCTC	CGAGCCTACA	TCAAGACCCA	AAGTGAACAAG	GAGC
23_Cjac	AGAGCCAAAGC	CCAGCCACAT	CCAAGTCCTC	CAAGCCTACA	TCAAGACCCA	AAGTGAACAAG	GAGC
32_Hsap	AGAGCCAGAGC						
59_Ptro	AGAGCCAGAGC	CAAGCCACGT	CCATGTCCTC	CGAGCCTACA	TCAAGACCCA	AAGTGAACAAA	GAGC
95_0gar	ACAGCAAGAGC	CAAATGAGGT	CCACATCCTC	CAAAACTATA	TCAAGACCCA	AACTGAACAAA	GAGC

#### Indel

```
930
                                    940
                                               950
                                                         960
       TGGAGGAGCTCCAGCAGCTGGTGGAAGAGCAGCTCAAGGCCAGCGAGGAAAGGCTCAGCAGCAAG
75_Mmul
       TGGAGCAGCTCCAGGGACTGGTGGAGGAGCGGCTCAAGGCCAGCAAGGAAAGGCTCAGCAAG
12_Ggor
36_Nleu TGGAGCAGCTCCAGCGGCTGGTGGAGGAGCGGCTCAAGGCCAGCGAGGAAAGGCTCAGCAGCAAG
47_Tsyr TGAAGCAGCTCCAACAGCTGGTGGAGGAGCAGCTCAAGGCCAGCGAAGAAAGGCTCAGCAGCAAG
       TGGGGCAGCTCCAACACCTGGTAGAGGAGAGGCTCAAGGCCAGTGAGGAAAGACTCAGCAGCAAG
06_Mmur
32_Pabe TGGAGCAGCTCCAGCAGCTGGTGGAGGAGCGGCTCAAGGCCAGCGAGGAAAGGCTCAGCAGCAAG
23_Cjac
       TGGATCAGCTCCAGCGGCTGGTGGAGGAGCAGCTCAAGGCCAGCGAGGAAAGGCTCAACAGCAAG
       TGGAGCAGCTCCAGGGGCTGGTGGAGGAGCGGCTCAAGGCCAGCGAGGAAAGGCTCAGCAAG
32_Hsap
59_Ptro TGGAGCAGCTCCAGGGGCTGGTGGAGGAGCGGCTCAAGGCCAGCGAGGAAAGGCTCAGCAGCAAG
95_Ogar TGGGGCAGCTCCAAGAACTGGTAGAGGAGAGGCCTCAAGGCCAGTGAGGAAAGACTCAACAGCAAA
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					1020	
75_Mmul	TTGACTGCACT	ΓAGAĠCGGCC	CTTCCAGCTA	CCTCCGGGT <i>I</i>	A A A G G Ċ A A G A G	CAAGACCAAG
12_Ggor	TTGACTGCACT	<pre>FAGAGCGGCC</pre>	CTTCCAGCTA	CCTCCGGGTA	AAAGGCAAGAG	CAAGACCAAG
36_Nleu	TTGACTGCACT	<pre>FAGAGCAGCC</pre>	CTTCCAGCTA	CCTCCGGGTA	AAAGCAAGAG	CAAGACCAAG
47_Tsyr	TTGACTGCACT	<b>TGGAGCAGCC</b>	CTTGCAGCTA	CATCCGGGC	AAAGGCAAGAA	CAAGACTAAG
06_Mmur	TTGACTGCACT	<b>TAGAGCGGCC</b>	CTTCCAGCTA	CCTCTGGCTA	AAAGGCAAGAA	CAAAACTAAG
32_Pabe	TTGACTGCACT	<b>TAGAGCGGCC</b>	CTTCCAGCTA	CCTCCGGGTA	AAAGGCAAGAG	CAAGACCAAG
23_Cjac	TTGACTGCACT	<b>TAGAGCGGCC</b>	CTTCCAGCTA	CCTCCCGGTA	AAAGGCAAGAG	CAAGACTAAG
32_Hsap	TTGACTGCACT	<b>TAGAGCGGCC</b>	CTTCCAGCTA	CCTCCGGGTA	AAAGGCAAGAG	CAAGACCAAG
59_Ptro	TTGACTGCACT	<b>TAGAGCAGCC</b>	CTTCCAGCTA	CCTCCGGGTA	AAAGGCAAGAG	CAAGACCAAG
95_0gar	TTGACTGCACT	<pre>FAGAGCGGCC</pre>	CTTCCAGCTA	CCTCCAGGC	AAAGCAAGAA	CAAGACTAAG