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BRE/TATA box

(sRk52) Pab(1103978-1104286) TCGTAGGTGGTAACAAATGGAACGTTTGGGCTCTTTTGTGTTTGCCTAATTTTAAATACTTTTACAGTATATTTGTGTTGGCTTGGCAGCTCTCCGATATGTGGGGT
 (sRk48) Pab(985726-986038) TTCATGGATAGCGACAAACGGAACTTTTTTGTCTTTTGGTTTCCATAATTTTAAATACTTTTACGGTATATTTATGTTGGTTGGTAGCTCCCGTATGTGGGGT
 Pab(909112-908804) TTTAGTTCCAAACTTTTAATTTGTTTACTACCTCTTTGGAGCTCTTATAGGAGCGCTTTTCTAAACAAACACTTCCCTCCACTTTCTTACCTTCTCTTACAGCTT
 Pab(1024101-1023793) TAGAATCTCGAAGTGCACATGCTAGGGGCTGTTTCATAGATGACAACGACGTTTGAGCTCTTTTATTGGATTGAGTGGTAGGCAGTTCTAGGTATGTGGGGT
 Pab(862648-862350) TTGCCCCCACTTTATGATACCTGGGACTGTTTCGTGGAGTAGCAACAAATGAAAAATGTTTACGGTGATTAGGTTGGCTTGGCAGTCTCG--ATGTGAGGT
 Pab(5328-5010) CCCTAATGATGTCACGTTTCCACGTTACACACATCCCAATTTCTCGAATATATGGAGGTCATCCTCGTCAATAAATGTTTGGGTTTGAGAGAAAAAGATATT
 Pho(369969-370267) CAATGCTAAGAAGGGCTGGTTTCTGAAAAATAATAAAGATTATTTGGATGAAATTTTAAAGCTGGTGAACTTTGTAGTGGTGGCAGTTCTCGGCATGTGGGGT
 Pho(374302-374573) TTCGTAATAAGTTAAAGTGTAGAGTTTGTGGGGTTTGTAGAACGAAATTTAAATGTTCTGTGGTGAATTTGGGGTGATGTGGCAGTCTCGGTATGTGGGGT
 Pho(543812-544132) ACTGCTCCTCGTAATACTGTATAAAACACCACTAGCgttagtatatgcaatcccaaccctatggctccacatggaatataagaactgaagaaatcaatacaggga
 (PfQ12) Pfu(1792887-1792576) atataacagatgttaggagcatttctaaagaaataaCTATTACTTAGCTTCTGCTTCCCTGGGAAAGAGCGAATTCAGCTTCTCCTTTGCTTACGGGTGCTT
 Pfu(1500218-1499923) ATGAGATTGAAGTGAAGGGCTGTTTCGTAATAGTTGAGATCGTAAGGTTTATGAGTCATACTTTATTAATGGAGTCTTCTACCTTGTGCCGATATGTGGGGT
 (PF1015) Pfu(1204258-1204554) TCATAGAAAAGGCTGAAAAGAGGACGCTGGGATGGGAGGTGGAGGATGAGGTGTTCTCTAAATAATTTAAAAAAGCGTAAGAGCTGCTATCTGTGTGGTGTAGTG
 Pfu(971398-971033) tttaggaaactccagcgcataatcgaatacagaaggcgaagttaagggtataaaagttatttctgtaaccccgcttatacttccaccctgtgcccgatatgtggggg
 (TK1842) Tko(1653785-1654163) TTTAGGAAGCTTCAAACATAATCGAATAACAAGGCCAAGCTGAAGGGTATGTTTGTGTAATCCTGCTTACACTTCGCTCGCTGTGTCCGTTATGTGGGAG
 (TSIB2041) Tsi(1826786-1826408) tttcacagctccagtcattatcagagtacagaaggcgaagtgaagggtggttttgttgatccggcttacacttccctccctgtgtccgatatgtggggg
 (TSIB1736) Tsi(1559055-1559433) TTTAGGAAGCTTCAGTCAATCATCGAGTACAAGGCCGAAGTTAAAGGTGTTGAGGTTGTTTTGTTGATCCAGCTTACACTTCTCCTCTGTGCCGGTATGTGGGGA

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(sRk52) Pab(1103978-1104286) TTG--TAAGCCCGAATGGGGACTGCCTACTGCCTGAAGATGTGGGGAGTTCCGTTCCCCC-CGAAAGCCAGCCGATGAAGACGAGAGGCTGGAAGGTATCCACTA
 (sRk48) Pab(985726-986038) TAAGTTAAGCCCGAATGGGGACTACC-----TGAAGATGTGGGGAGTTCCGTTCCCCC-CGAAAGCCAGCCGATGAAGGCGGGAGGCTGGAAGGTTATCCGTTA
 Pab(909112-908804) GTTTCATGGGTTTGTCTTACCAAGCAACTCTGAAGATGTGAGAAGTTCCGTTCCCCA---AAGGCCAGCCGATGATAGCAAGAGGCTGAAGATTATCCACTA
 Pab(1024101-1023793) TTG--TAAGCCCGAATGAAGACTGCTTA--GGCTAAGAGATGGGGAGTCTGTGTTCCCTTACGAAGGCCCAACCGAGGAATATGAAGGCTGGAAGGTTATCCGCTA
 Pab(862648-862350) TTG--TAAGCCCGAATGAGGACTGCC-----TGAAGATGTAGGGAGTTACCGTTCCCCC-GGAAGGCCAGCCGATGAAGATAGGAGGATGGAAGGTTATCCGCTA
 Pab(5328-5010) TTGTGCTCTAGGAGCATTCTGTTGAGAGACCTGAGAGGAGTGGGGAGTTCCGTTCCCCAC--GAAGGTGAGCCGATGAAGGAGAGTAGGTTGGAAGGTTATCCACTA
 Pho(369969-370267) TTC--CAAGCCCGAATGGGGACTAC-----TGAAGATGTGGGGAGTCAACATTCCCCC-CGAAGGTGAGCCGATGAAGACGGGAGGTTGGAAGCTACCCGTTA
 Pho(374302-374573) TT--TAAGCCCGAATAAGGACT-----TGTGGGGAGTCTCCGTTCCCC-----TATAATGAAAAGGGAGGTTGGAAGGTGAGCCGTTAA
 Pho(543812-544132) gaaaagagtacatatgcaagtttagAGGAGTACTTCGAATTGCCATGACGTTACCGTTCCCCC-CGAAGGTACCCGATGAAGAAGGGAGGTTGAAGGTGAGCCGTTGA
 (PfQ12) Pfu(1792887-1792576) ATAGCCATAAGTGCCTAGACCTCTTCTGGGACTtgagatgtggggagtcaccggttccccc-cgaagccagccaatgaagatgagaggctggaaggttgcccatgt
 Pfu(1500218-1499923) TTG-ATAAGCCCGAATGGGACCGGAT---CCGAAGATGTGGGGAGTCTCCGTTCCCCC-CGAAAGCCAGCCGATGAAATGGGAGGCTGGAAGGTTGGCCGTTAA
 (PF1015) Pfu(1204258-1204554) GTGATCTCGAGTATGTGGGGGTTTAAAGCCCAAGCGAAGTTGGAGAGTAACCGTTCCCCC-AGAAAGCCAGCCGATGAAGATGAAG-----CAGCCGTTT
 Pfu(971398-971033) taagttaagcccgatgggcacagggg/ \cctgaagatgtggggagtcaccggttccccc-cgaagccacccgatgaagaaggagggtggaaggtcagccgat
 (TK1842) Tko(1653785-1654163) AAAGTTAAGCCCGAATGGGCACAGGGT/ \CCCTGAAGATGTGGGGAGTTCCGTTCCCCC-CGAAAGCCCAAGATGAAGACGGGAGTGGGGAAGGTGCGCCGTTAG
 (TSIB2041) Tsi(1826786-1826408) aaaaataagcccgatgggcacagggg/ \ccctgaagatgtggggagtaaccggttccccc-cgaagccctccgatgaagacgggaggagggaaggttatccgcta
 (TSIB1736) Tsi(1559055-1559433) gaagttaagcccgatgggcacagggg/ \ccctgaagatgtggggagtttccggttccccc-cgaagccctccgatgaagatgggaggagggaagggcgaagccgctgg

Pfu(971398-971033) /TTTGAATGTC--AAATGTGGTTTGTAGGCTGATAGGGATGTTGTTGGAAGTTGGAATGTTGCTTGAAG\

Tko(1653785-1654163) /CTTGAAGTGTCTAAGTGTGGTTTGTAGGCCGATAGGGACGTTATTGGCTCTTGAATATCTCCCTTCGTG\

Tsi(1826786-1826408) /TTTGAAGTGTCTCAAAATGTGGTTTGAAGCCGACAGGGATGTTGTGCGCTCTTGAACATCCGCTTGAAGG\

Tsi(1559055-1559433) /TTTGAAGTGTGTTGAATGTGCGTTTGTAGGCCGACAGGGATGTTGTGCTCTTGAATGTTGCTTGAAG\

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(sRk52) Pab(1103978-1104286) CGAG-----AGTTCCTCCCTGCCACTCGTAGTGATGACCAGAACGGGCTCCTTCTATATACG
 (sRk48) Pab(985726-986038) CGAGTGGTGCAAAAGTTCATCCCTGCCACTCGTAACGGATAAACCAGAACGGTTTCTCATAAATCCTG
 Pab(909112-908804) CGAGTGGTGCAAAAGTTCCTCA-----AAAGTAGCGGATAACCAGAACGGTAACAGATCCTCAGGT
 Pab(1024101-1023793) CT-----GAAAGTTTCACCTGCTCCAAAGTAGGCATCTAACTAAAGATCGGATAATAATACCTG
 Pab(862648-862350) -CATTTGATTTCCCAAGTTCCA-----AAGTAGCGGATAACCAGAAATGGTTTAATTAACCGTTG
 Pab(5328-5010) TGGTGGTGCAAAAGTTCATCCCTGCCACCTGAGCGGATAACCAGAACGGTTATTATCATGGAAT
 Pho(369969-370267) CG-----AAA--ttaacactctacacact-ttacgggtaggcagaacaggatcttcttaacactc
 Pho(374302-374573) CGAC-----A-TTACG-----CAA--ATCACGGCTGACCAGAACGGTTACTGGGTATTTACA
 Pho(543812-544132) TGATATTCA-CAAACTTACGCAAA-----ATCACGGCTGACCAGAACGGTTGAGACTGGGATAA
 (PfQ12) Pfu(1792887-1792576) cgatatattacgaactttacacaa-----atcacggccaaccagaaacggcattcttatagcctt
 Pfu(1500218-1499923) -----TTACA-----CAA--ATACGGCCAACCAGAACGGTTAAAAATGGTGGGACTCCTTT
 (PF1015) Pfu(1204258-1204554) -TAATATTCAACAGCTTAA-----CAAAA-GTTTGGCTGATCAGAACGGATATTGAGTATTATCTC
 Pfu(971398-971033) cgaca-----ttaca-----caaa-tcacggctgaccagaacgggttactttatgcggcggc
 (TK1842) Tko(1653785-1654163) cgatg-----tttacgaactttacacaaattacggctgaCCAGAACGGAACCGGTATTAAGGA
 (TSIB2041) Tsi(1826786-1826408) cgata-----gtttcacatgtttt-aaaagtagcggataaccagaacgggtctgctgaggtaga
 (TSIB1736) Tsi(1559055-1559433) cgat-----gtttatgaactttacacaaattacggctgaaccagaacgggacgactatggtcgt