

PFC	Dist to 3' gene	Length	Sequence
TruAB _{EF} 61	2,733	59	TGGAAATGCGATCTACTGCAGATGTTATTTGATATC CATTACAAAGAGTTCCTTCATT GCATCCTGTGAGGTTCCGAGGAATCCGCCCTGATG GCTGCACATCCTCCCAGTTGCTCCAGTAACTTGCC ATAAAACGCGGACTCGTCTGAAGCATTGGAGAGCA GCGCAATAAAGCGTCTGCGACCAAGGTTATTAATG
TruAB _{EF} 65 _{acde}	2,263	163	TGACTGAGAGGTTGAAAAAC
TruAB _{EF} 66	2,029	31	CCACTGGGTGGCGCCCCTCAGATCAATGTCA GGGAGGAGTCTATAGCCGCTAGTGTTTAAATTGGC TGCAGGCTACAGTCAGAAGCGTGAGTATCAGTAATT ATTCAGCAATGTTTTGCACAAGAAATGTCAGCCAGA
TruAB _{EF} 75 _{abcd}	147	112	AAGG
TruAB _{FG} 80	1,492	28	AAGCAGGATGTTTTGTTTTTGTATTTTG GCCGGATGCGTCTGCCTCGGCAAGAAAAGGGACAC AAAAGCTGAACTTTATTGGTGATATTTAGCTTAGAC
TruAB _{FG} 81 _{ab}	806	97	AGCCAGGTTGTAACCTAAAGAAAAAC
TruAB _{FG} 86	859	14	CAAGGCCTCGGTTA AAATCATAACCTTTATAGCTTATAAAAAGTCATATT TTCCCAATAAATGCGCGGAATTTGCTATTAATAAAT ATGGTGGTGGGGGGAAGGGGGATGGTGGCCATTG AGTTGAGCAGCAGGCTGATTTATTATTATTGTTCTG TCCCACGTCACGTGTTGGGGCGCCCTATCCAGTC TTGGGCAAGGTTCAACTTGCTGGACGCACTGGTCCC
TruAB _{FG} 87	306	245	CTAGATGGTGCAGAAAGGCACGCAATGAA
TruAB _{FG} 90	864	28	CAGGACAAGGCCTCGGTTATGATCATTG
TruAB _{GN} 109	890	13	TGGCAAATCATCT
TruAB _{GN} 112	4,295	24	TATATTTGAGTATCATTGTTAATG CAATATGGTGTGATATATGATACAGCAGCCATAAAT
TruAB _{GN} 131	2,260	56	AATTTAGAACTTCTGTTA AAATGTGTTCTTAGGGCAGGGAGCTGTCAGACCTTT TGGCGTAAAAGATTGATCACACTCAGGGACCGAGGT
TruAB _{GN} 204 _{bcd}	1,027	80	CTTTGTTT ACAGACTTTCGTCTATAAATGAATTTCTCAATCATA
TruAB _{GN} 208	1,217	44	ATCGAGCC TGTAATAAAAGCAACCCCTACCCCGGAATGAGGC GTTACCTCTCCGACGTTCTCGATCAATCACAGGGGA CAGTGGCTTCTTTTGATAGAGGAAAAAAACCTCAA ATTGTCATTGGGCAGATTTAATCATGTGACAAGCAA CACGGTCTAATTCCAACCATCCTTGTGAAAGCAA
TruAB _{GN} 211 _{abce}	297	201	TAGTTTATGGCACAGGGGTCCC