-888 GACGGCGAAGCACCGGCGGCCGCCGCTGCCACCGTCGCCTCGTTCGATTTGGGAAGGGC -849 GACGGCGAAGCACCGGCGGCCGCCGCTGCCACCGTCGCCTCGTTCGATTTGGGAAGGGC *****************************
-789 TCGCTCGGTAGGTTGGTTAGGGTTCGCTCGGCTGGAGGAGAAGAGATGCTTGGTTACTGG **********************************
-729 GCCAGAAAGCCCAAAAATATAATGGGCTGATGAAAAATTCGTCTGGAAAAAAACGCCATC *********************************
-669 GAAAAATAGCACCGATACGCGCTGTAGAGCGCAATCGCCGTATTTAGCGTGAATAGCGC *********** **********************
-609 CATAGCGAGTGAAATCTGCATAGTGTAGCGTTTAGTTTCCAGAAACGCTATAGC * ***** ************************
-555cgctatagcgcg <mark>ctatttttt</mark> ccttcatccatagagatgagcatatgtgtgt
-528 ATCTATGAGCGTCTT-TTATGTACTGTGAGAAAGAAAAAACCCAGACTAAGGAAGTGC
-502 ATCTATGAGCGTCTCCTTATGTACTGTGTGAGAAAAAAAA
-469 ATGCATCTGGCTCGGCGAGTGATTGTAAACAACTGGCTCACTGCATTGCATGCGTACATG
-442 ATGCATCTGGCTCGGCGAGTGATTGTAAACAACTGGCTCACTGCATTGCATG
-409 TGCGCAGTACACACGCAGCTGGTGGATCCAGCCAGCGGATTGTCTGTCTCGC
-382 TGCGCAGTACACACGCAGACGCAGCTGGTGGATCCAGCCAG
-314 -355 TGGCGGGAGAGGCACGCAGTACGCACCCAGATCATCACCCCA <mark>AT<i>ACGTGGC</i>A</mark> CGCCGGTCC
-333 IGGCGGGAGAGGCACGCAG IACGCACCCAGAICAICACCCAGAIAACGIGGCACGCGGTCC -322 TGGCGGGAGAGGCACGCAGTACGCACCCAGATCATCACCCCAACACGCGCACACGCCGGTCC
-295 TTCCAGAGGCCATGTGCCGGCTACGTGTCGCTTGACCTGTACATGCATG
-262 TTCCAGAGGCCATGTCCCGGCTACGTGTCGCTTGACCTGTACATGCATG
-222
-235 CATCAGCGATCGACACGTATGCGTAGGCTGCATGCACGCATGGCCCCAAGGAGGGTAGCC
-202
-175 CAAGTCCCAACCCCGTGACAAAACCCTCAGTTAATAATCCGCCGCGCTAGCCACCTGCT
-161TGACAAAACCCTCAGTTAATAAT <u>CCGCCGCGCT</u> AGCCACCTGCT
-115 TAGCGTAAGCCATATATACACCCAGCCATGCGTCATTTGTACAGGTCGTCGTTGGCTCTC
-115 TAGCGTAAGCCATATATACACCCAGCCATGCGTCATTTGTACAGGTCGTCGTTGGCTCTC **********************
+1 -55 GTCCAGAGAAAAGCAGGGGAAGACAAGGAGAAAAGAGCAGAGCAGAGGAG
-55 GTCCAGAGAAAAGCAGGGGAAGACAAGGAGAAAAGAGCAGAGCAGAGCAAACATGTC

+6 CCGGTTCGTTGATCCGCTGGTGGTGGGGCGGGTGATCGGCGAGGTGGTGGACATGTTCGT
+6 CCGGTTCGTTGATCCGCTGGTGGTGGGCGGGTGATCGGCGAGGTGGTGGACATGTTCGT

Figure S5 Comparison of the promoter sequences between Rio Blanco and NW97S186. Identical sequences are labeled by *. ABRE CEs are indicated by blue or yellow boxes if two ABRE CEs overlapped. RY repeats are indicated by green letters. Motif IIB is underlined. Two SNPs in the ABRE CEs at -222 and -314 positions are shown in red letters.

6 SI S. Liu et al.