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481 ctggtcgtca accgcttcgc aactcccaag atcATTCTCG ATTCGACGAG CAAGGAGTGC
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601 TCGTACTAC GATGTGCAG CCCACTCCG GGCCGAGGAG GAGAGTGTCC TGCCTCGGTT
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781 ACCGCTGAGC AAAGGCCAGA GCTTGTCCAG GCAACCTGCC CTGCAGTCAA CTGGGTCGAG
841 TCAGTTGCTC CTGGAGATCG GGCCGCGTTT TAACTTCTCC ACGCCGTA CTCCGAAC TG
901 CGTGAACATA TTCCAGAATC TCGGGTACTC AGAGGTGCGT CGCATGGAAA CCTCCACCCG
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1021 GTCGGTGAC CGCATGACC AGTGCTTGT CACCGAGGAG AATACCCCA AGCGAGCTT
1081 TGACGAGCAG CTACCTGAGC GCCAGGCCAA CTGGCATTTC GTGCCGTTT TGGAGGAGGG
1141 TAGGGCGGCA CTGGAGCGGA TTAATCAGGA GCTGGGCTTA GCCTTCAAC ACTACGATTT
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1321 GGTGATCGAC GGCGTGGAGC AGCCAAGTC GCTGATTGCG ATGATCATGG ACACGCAGGC
1381 CCACACGAAC CCGAACAACA CCATTAAGTT CAGCGACAAC AGCAGTGCCA TGGTGGGATT
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1801 CGTTTGCGCC CCCACTCAG GTGCTCATTG AGGCGAGCAA TGGCGCTCC GACTACGGAA
1861 ATAAGTTCGG CGAGCCAGTG ATCTCTGGTT TTGCCCTCT CTATGGACTG AACAGTGCTG
1921 CCG

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FIGURE S1.—Sequenced region of *ade2^Δ*. Triangle = EY05576 insertion site as reported in Genbank accession BZ748943.1, Upper case = mRNA sequence, Lower case = intergenic sequence and introns, Lower case and underlined = partial sequence of a P-element inverted repeat, Red font = *ade2* CDS, Yellow highlight = sequence deleted in *ade2^Δ*.