

Table S5. PCR primers for dominant multiplex PCR markers

| Marker | Primer | Gene ^a | Oligonucleotide sequence | Chr. ^b | Nucleotide ^c |
|--------|----------|-------------------|-------------------------------------|-------------------|-------------------------|
| CHR1.3 | CHR1.3-F | <i>RFO2</i> | 5'-ttaaatacaggctggccctaatagccttt-3' | 1 | 7,969,890 |
| | CHR1.3-R | | 5'-gtggattttgcagtgctagttccctca-3' | 1 | 7,970,182 |
| CHR1.9 | CHR1.9-F | <i>RFO1</i> | 5'-gaaaaggagatgccgtttcgtccctac-3' | 1 | 30,425,560 |
| | CHR1.9-R | | 5'-caaagtcggtttcgattgggtttgt-3' | 1 | 30,426,060 |
| CHR3.3 | CHR3.3-F | <i>RFO3</i> | 5'-tgatggagtgaaaaagcatgtccattg-3' | 3 | 5,440,460 |
| | CHR3.3-R | | 5'-ctgaaagtataaccaaagcgcgagcaa-3' | 3 | 5,440,672 |
| CHR3.8 | CHR3.8-F | none | 5'-gtattggcgactccaaatcccattcac-3' | 3 | 22558313 |
| | CHR3.8-R | | 5'-gtcctgtgatagcctcaccgcttct-3' | 3 | 22558677 |

^a *RFO* QTL is linked to marker CHR*x.x*.

^b Abbreviation for chromosome number

^c Position of most 5' nucleotide in the Arabidopsis TAIR10 reference genome. Size of PCR product is the difference between nucleotide positions of the primer pair.