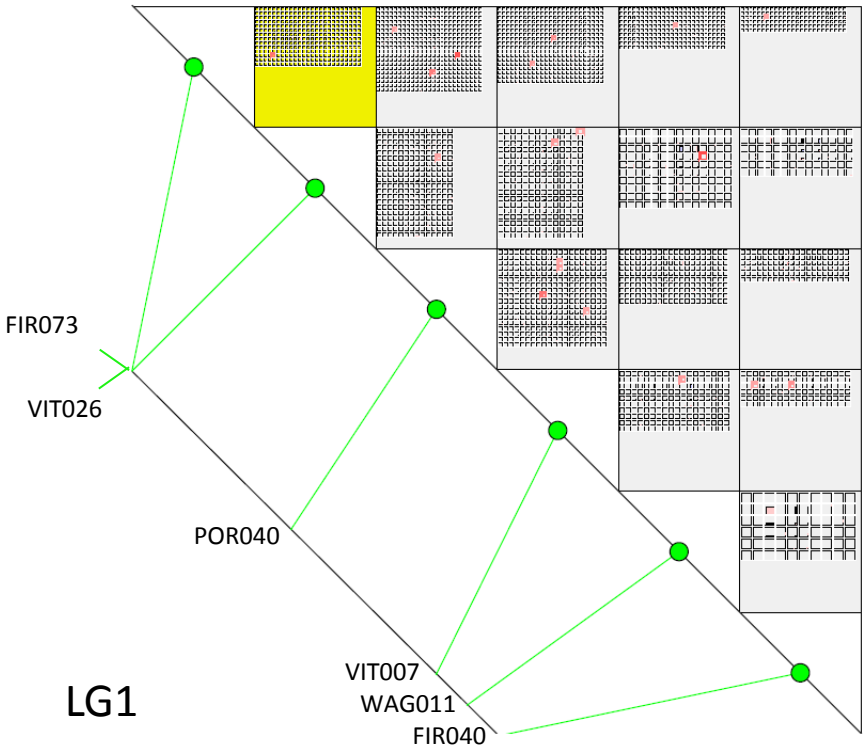
Figure S6-1: LD plots for all LGs from the two hybridizing oak species targeted in this study. Significant inter-allelic associations after Yates correction are indicated by red (D'>0) and blue (D'<0) dots within the small squares; their sizes being proportional to the D' value. Shades of yellow to white in the large squares indicate physical distances between loci (dark yellow < 100 kb; white > 20 Mb).

A – *Quercus faginea*



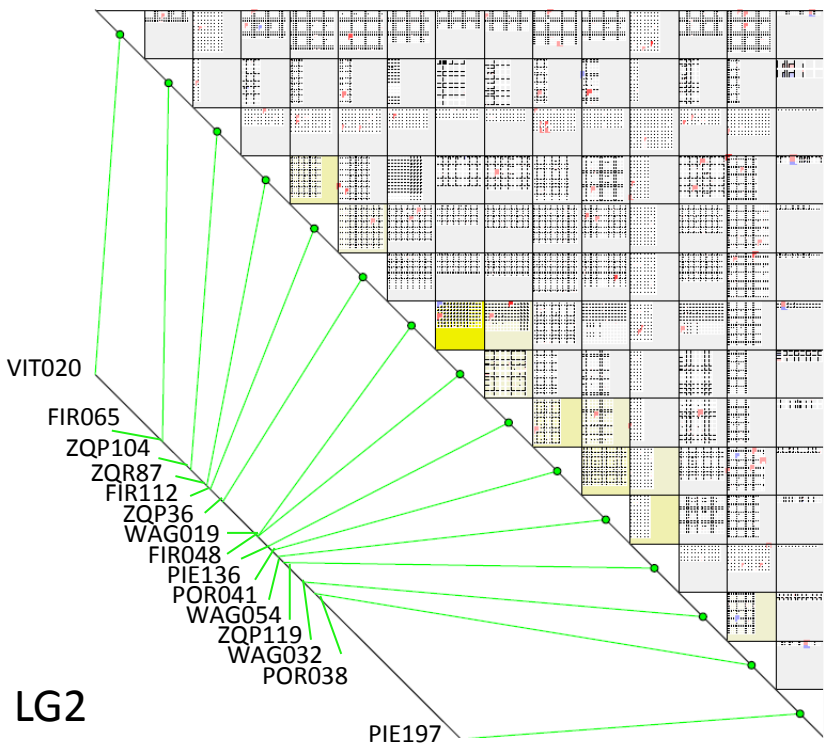
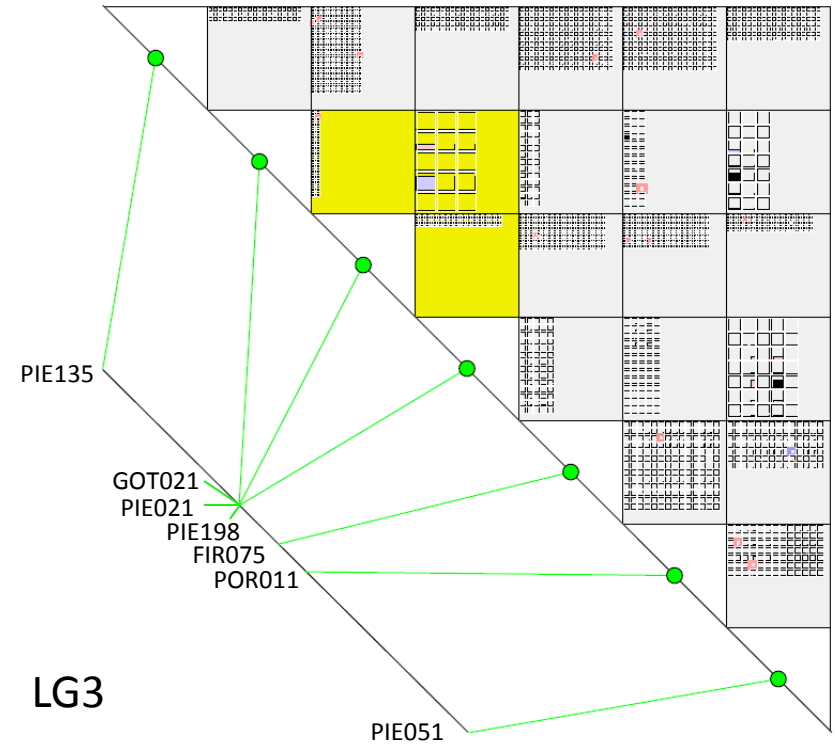


Figure S8-1: (cont.)



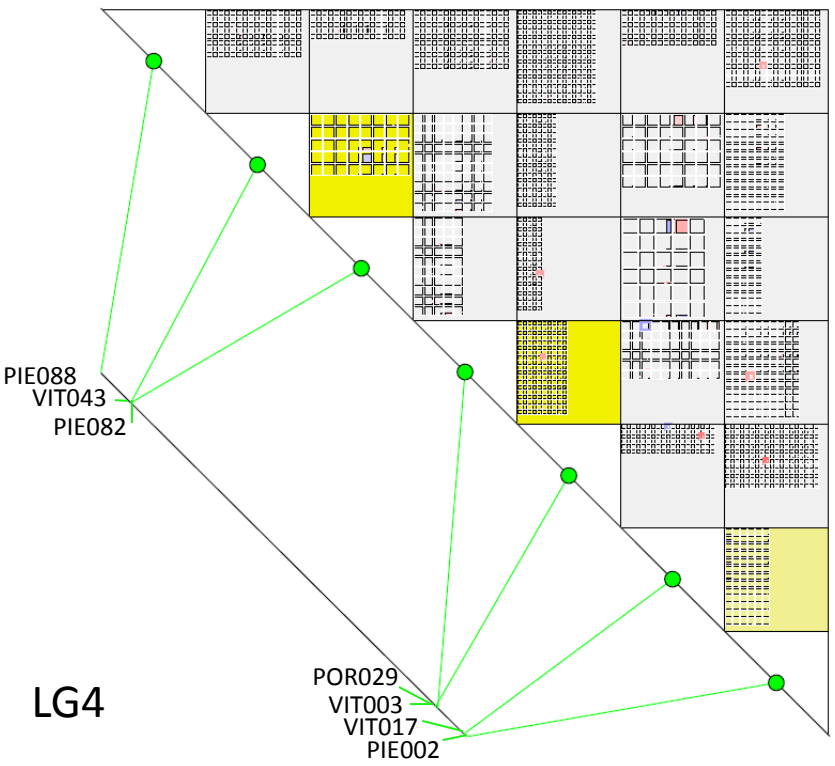
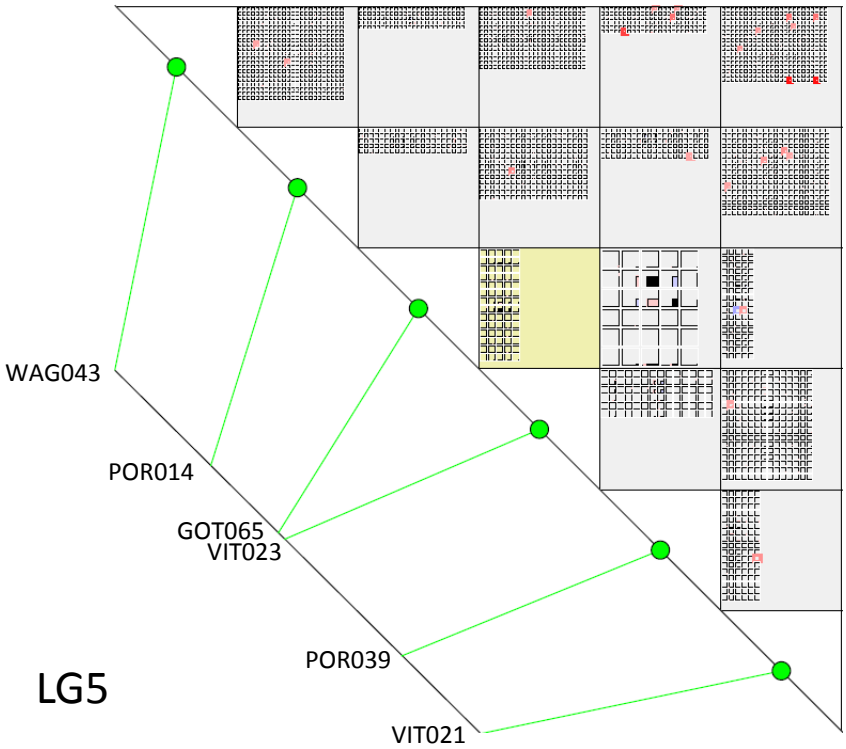


Figure S6-1: (cont.)



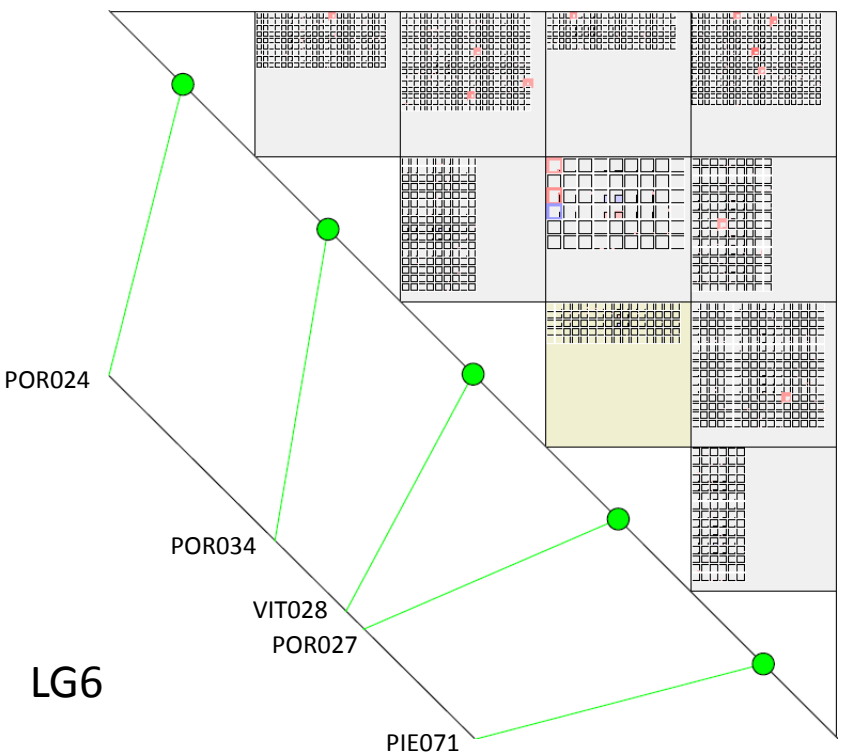
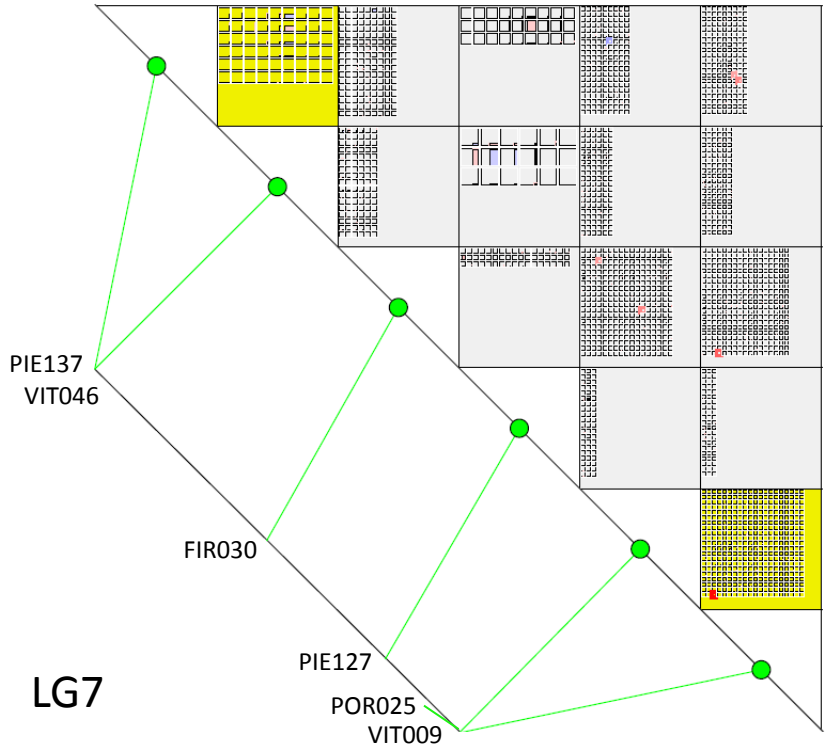


Figure S6-1: (cont.)



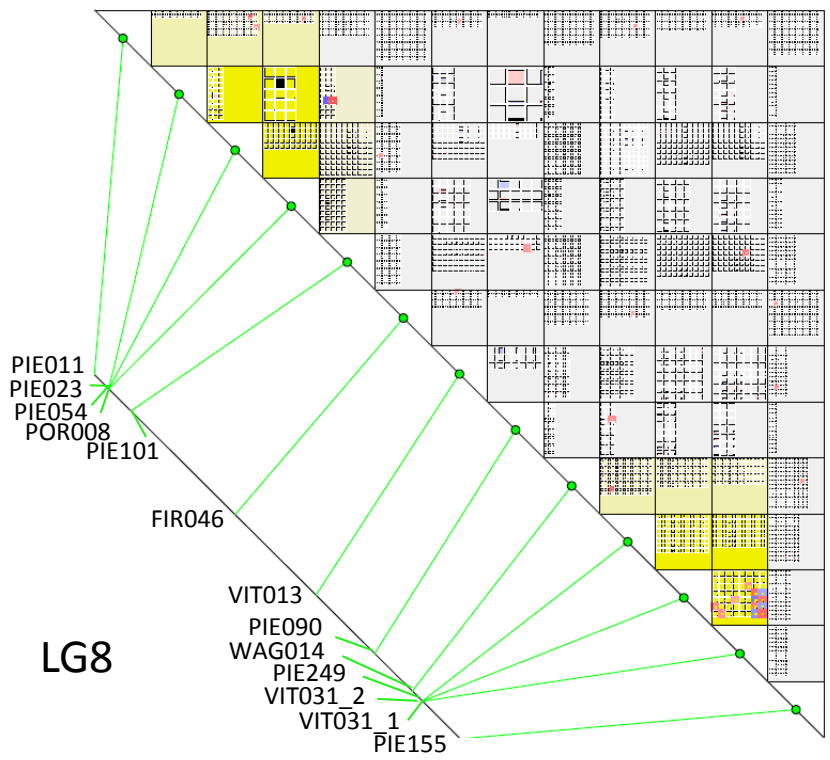
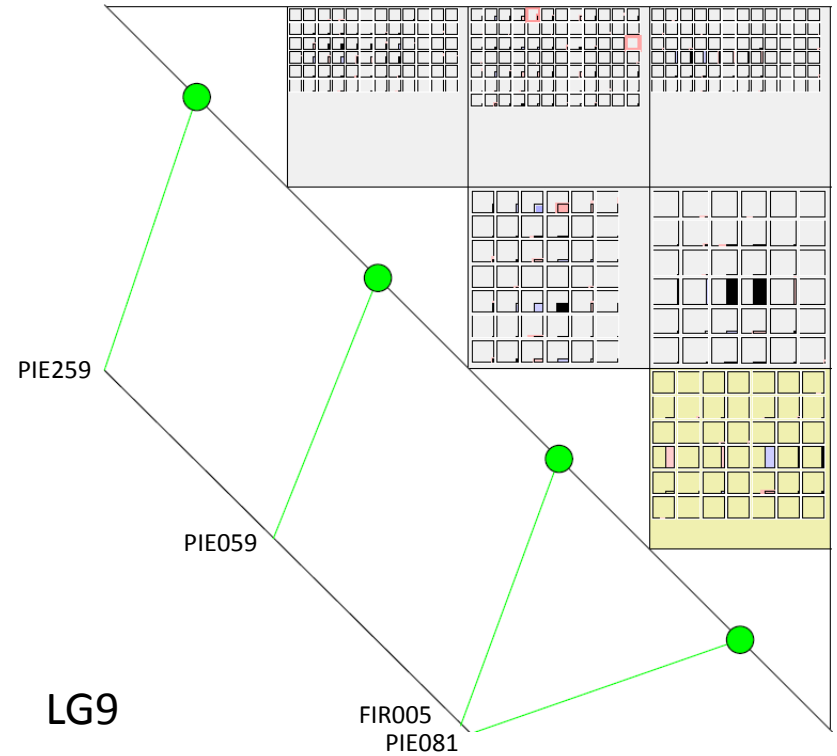


Figure S6-1: (cont.)



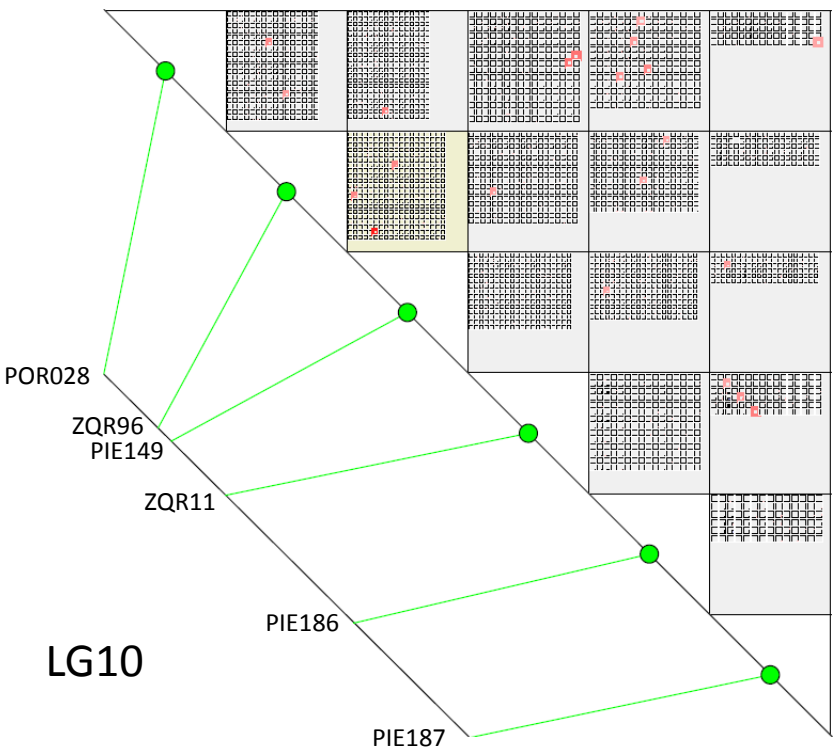
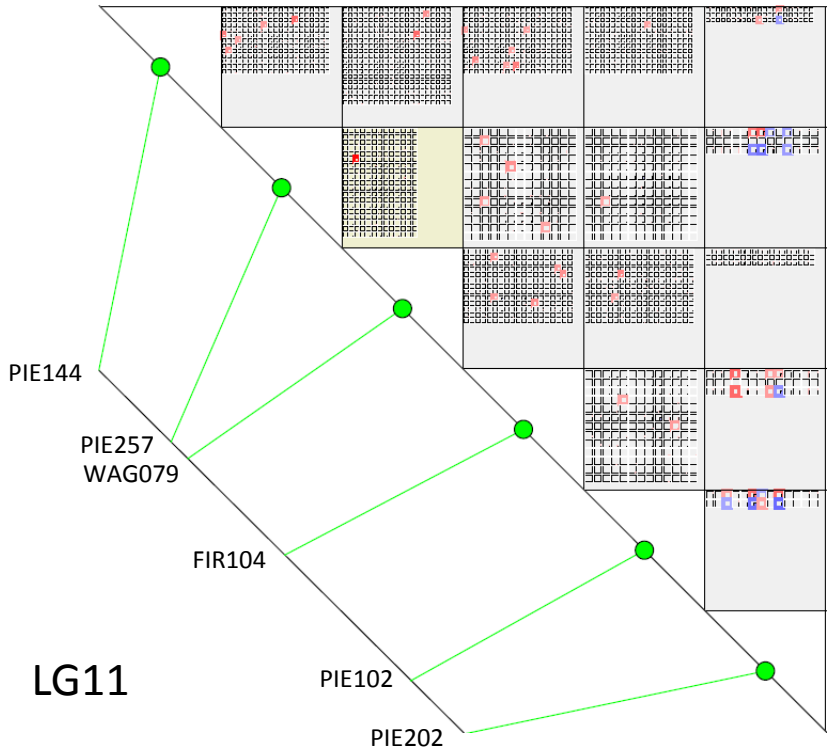


Figure S6-1: (cont.)



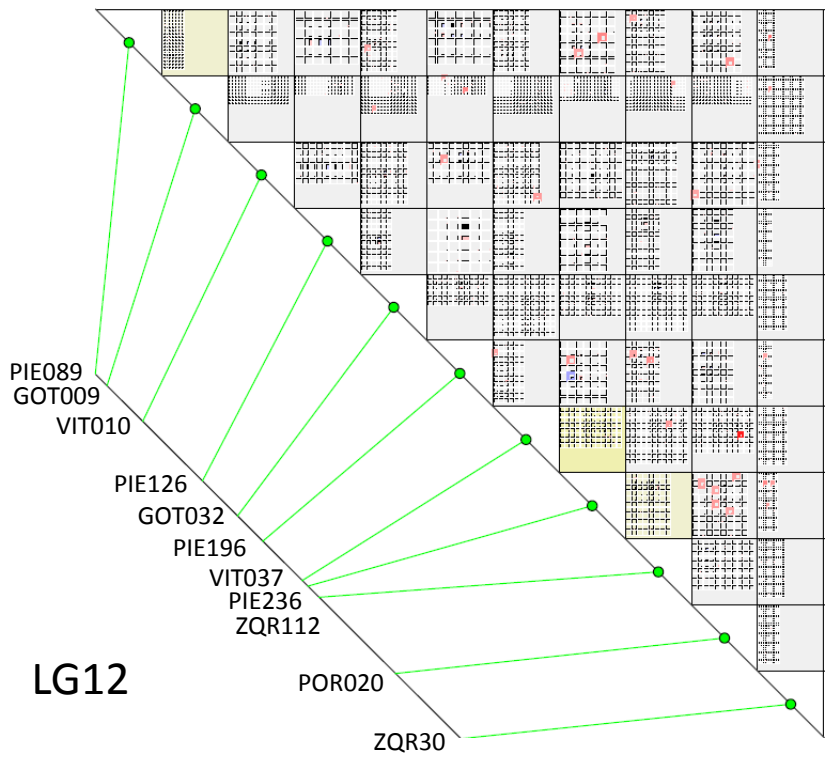


Figure S6-1: (cont.)

B – *Quercus pyrenaica*

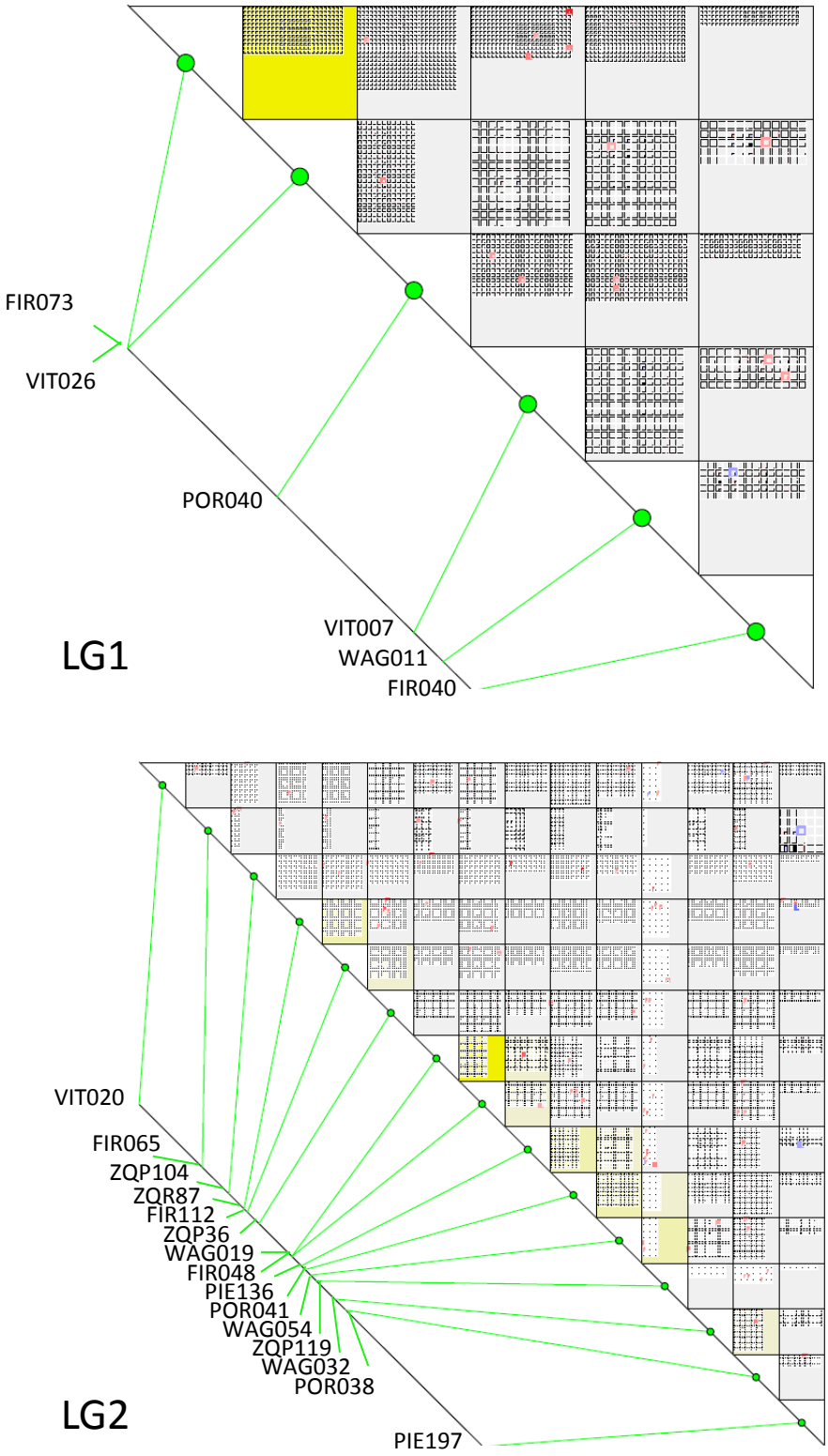
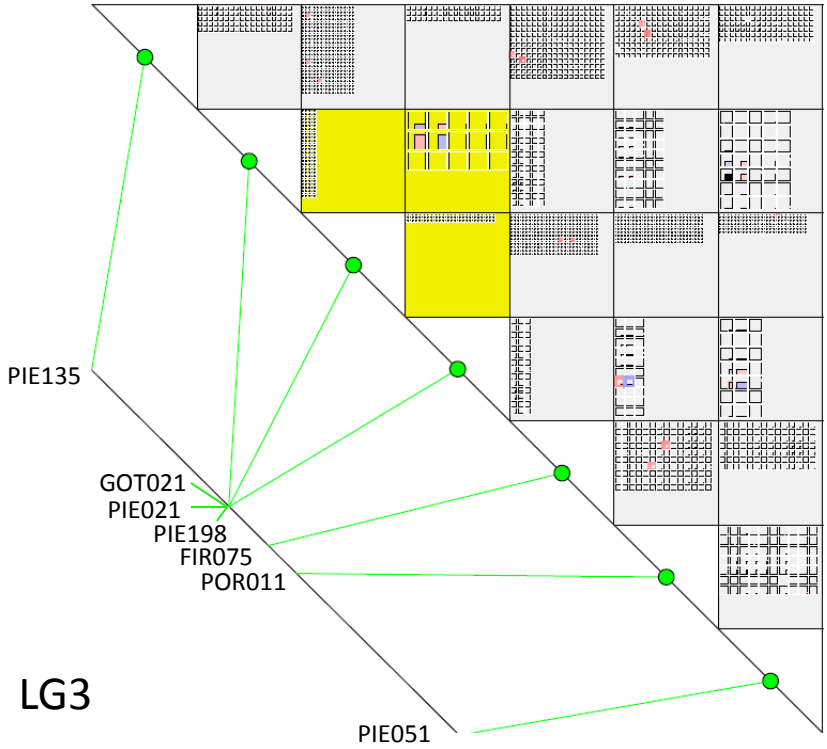


Figure S6-1: (cont.)



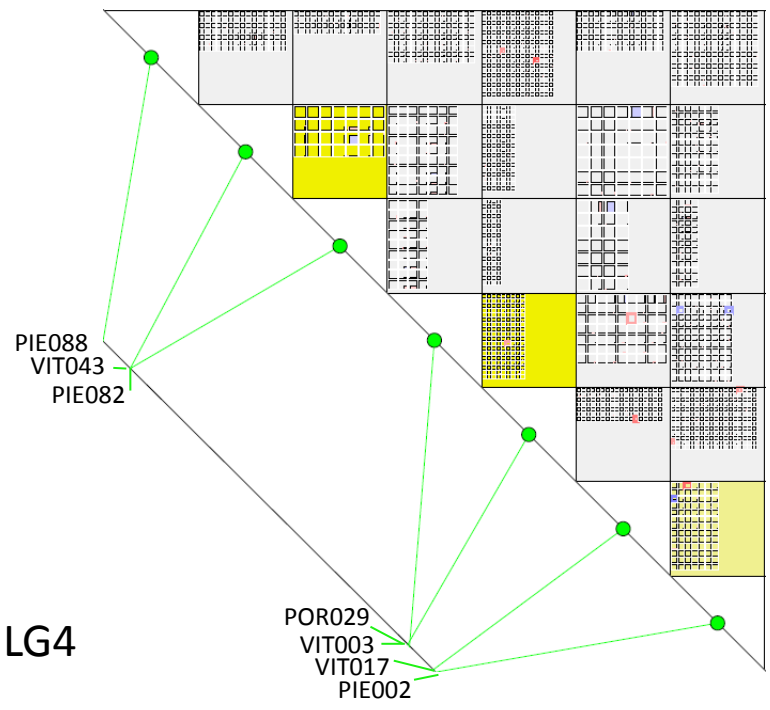
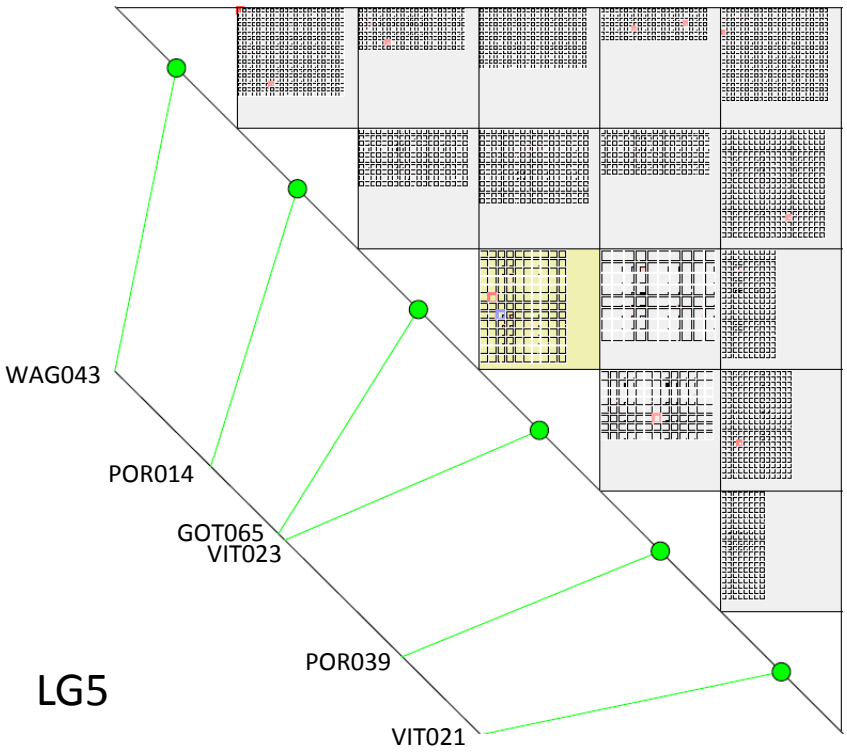


Figure S6-1: (cont.)



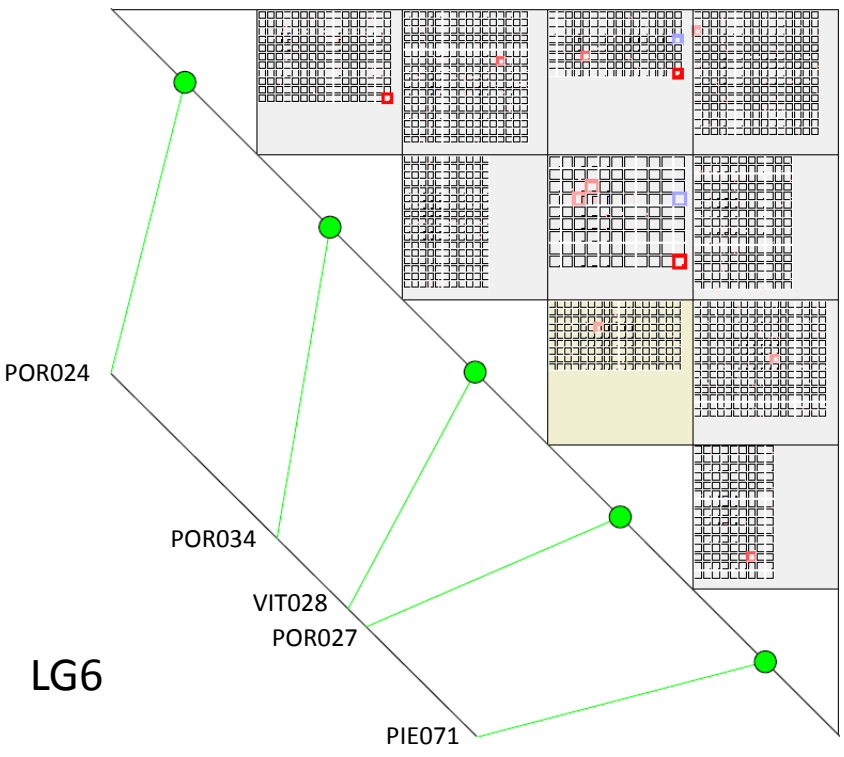
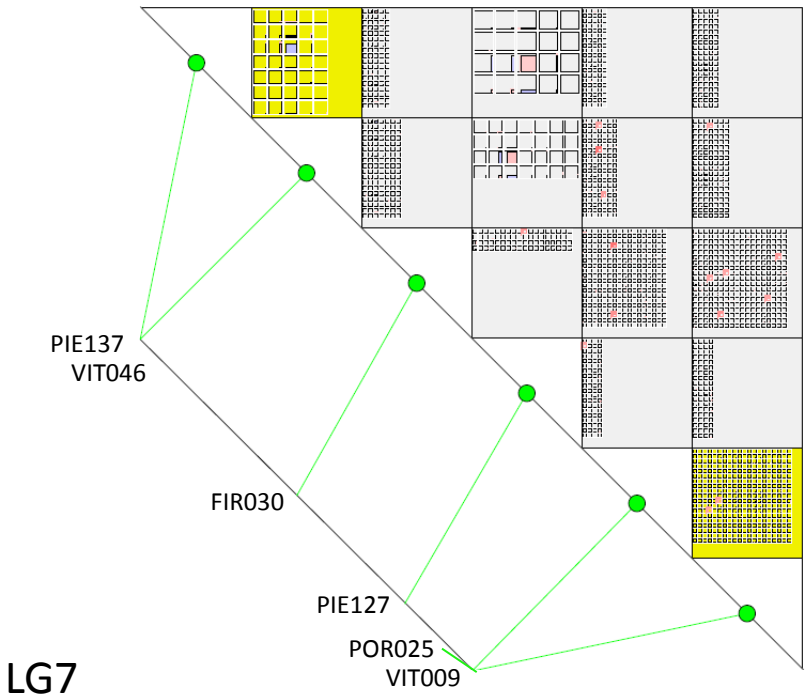


Figure S6-1: (cont.)



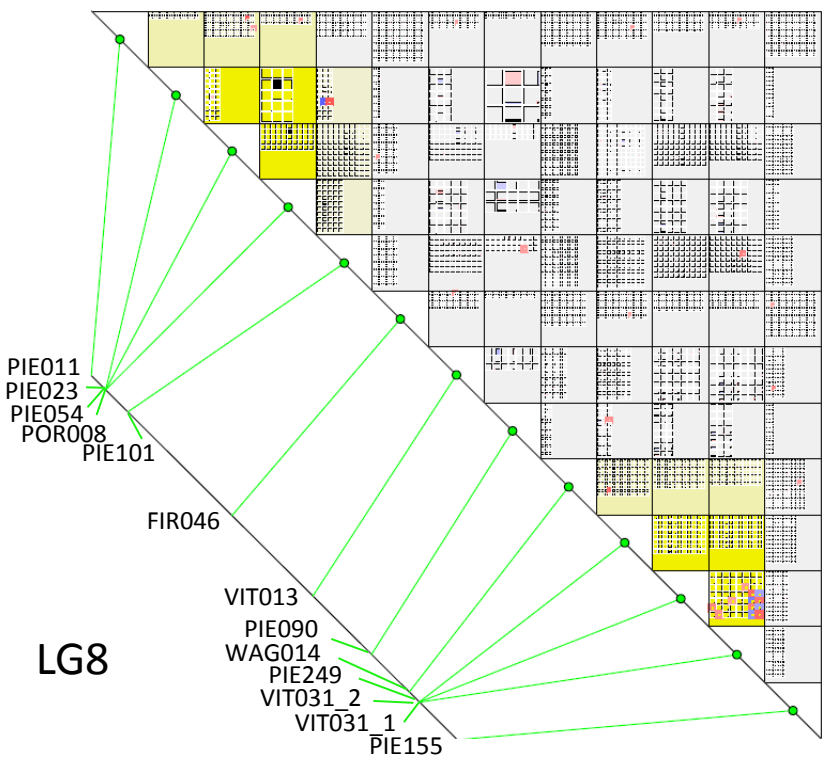
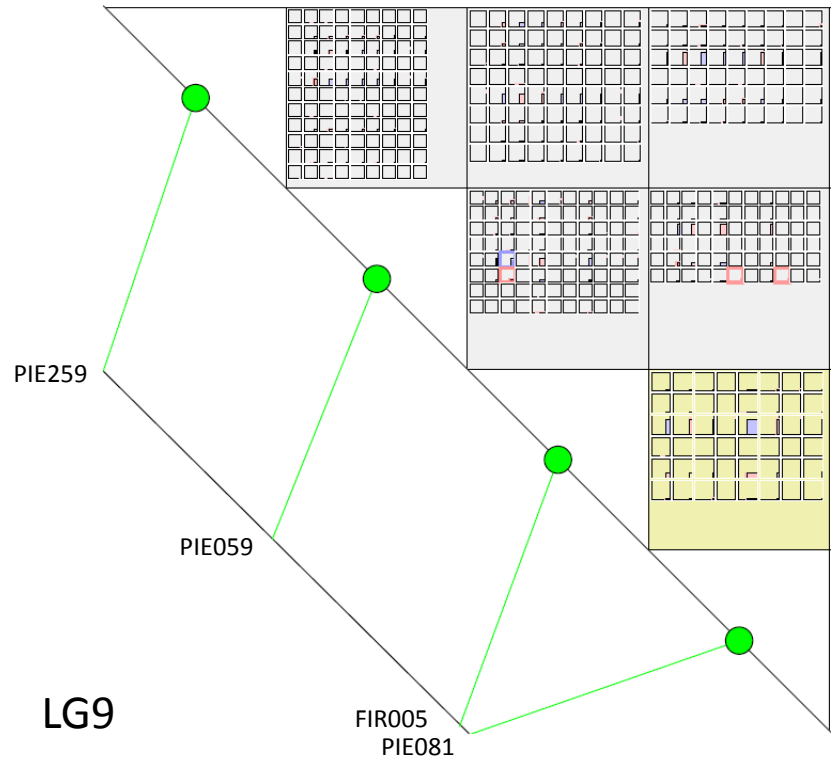


Figure S6-1: (cont.)



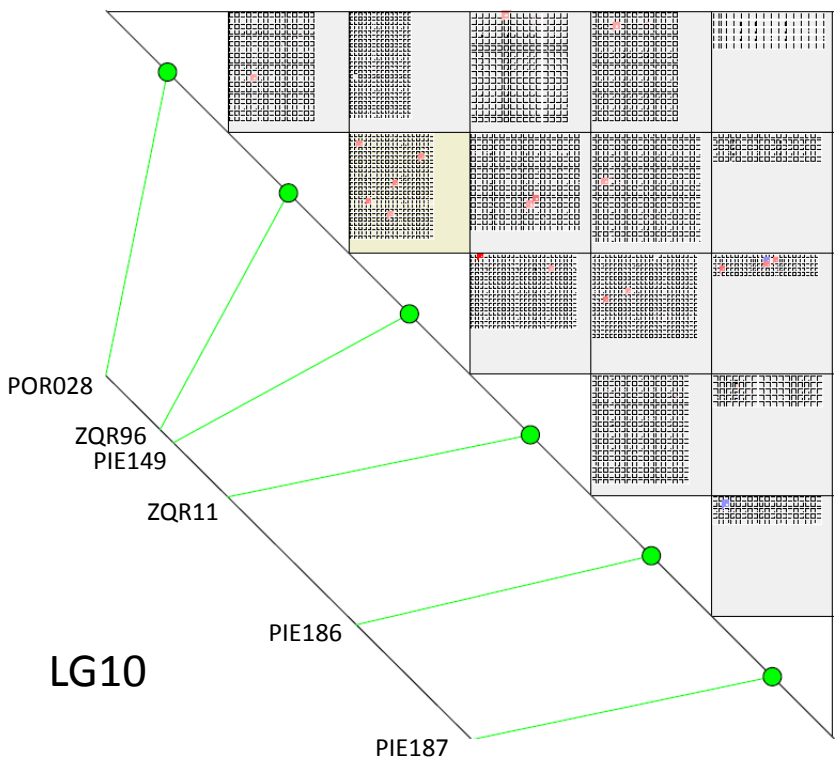
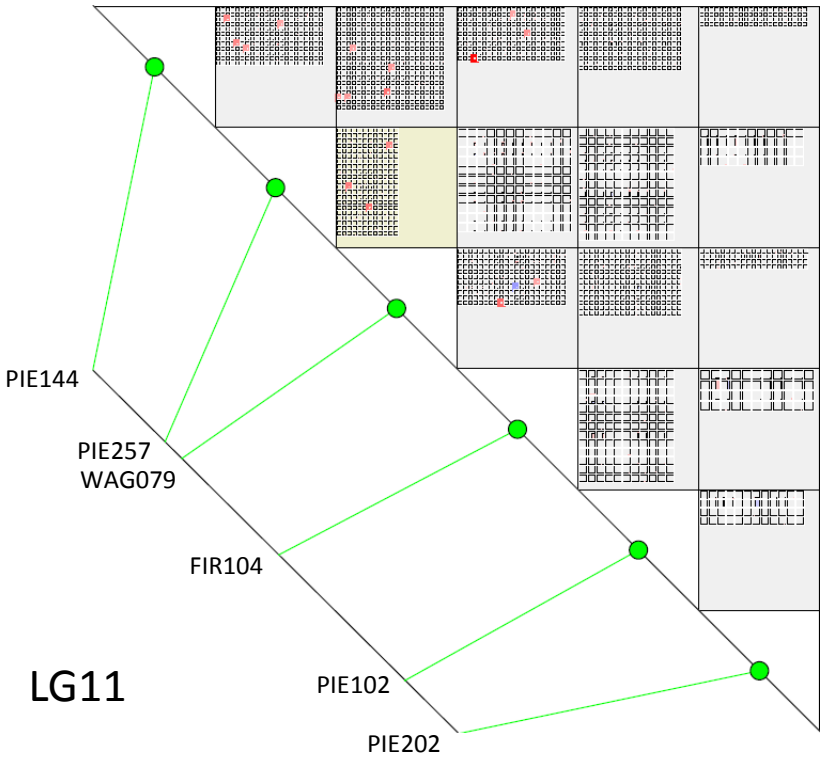


Figure S6-1: (cont.)



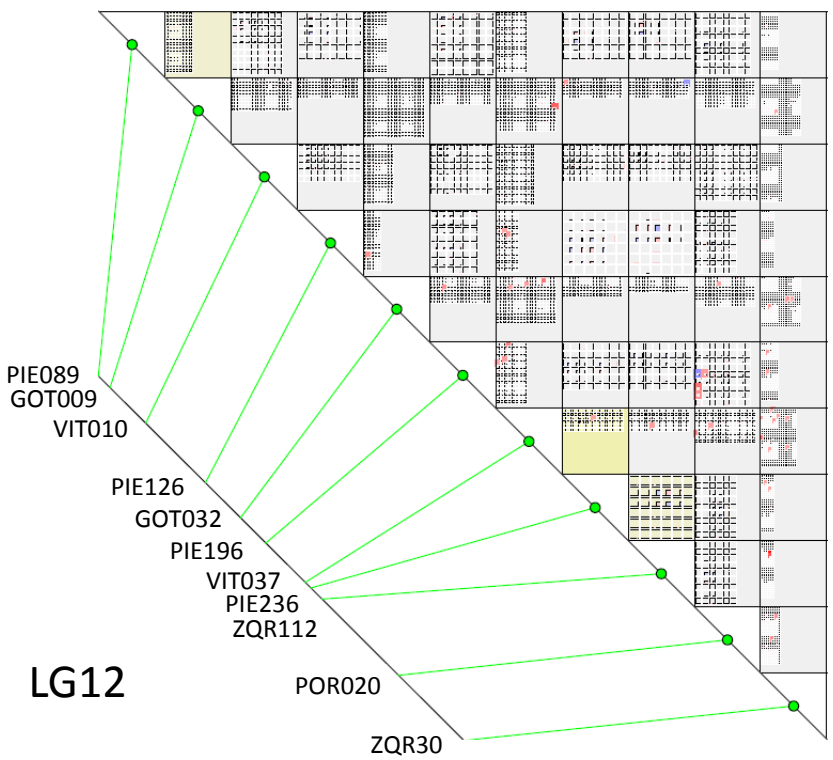


Figure S6-2:: LD plots for all LGs from the two hybridizing oak species targeted in this study. Significant inter-allelic associations without Yates correction are indicated by red (D'>0) and blue (D'<0) dots within the small squares; their sizes being proportional to the D' value. Shades of yellow to white in the large squares indicate physical distances between loci (dark yellow < 100 kb; white > 20 Mb).

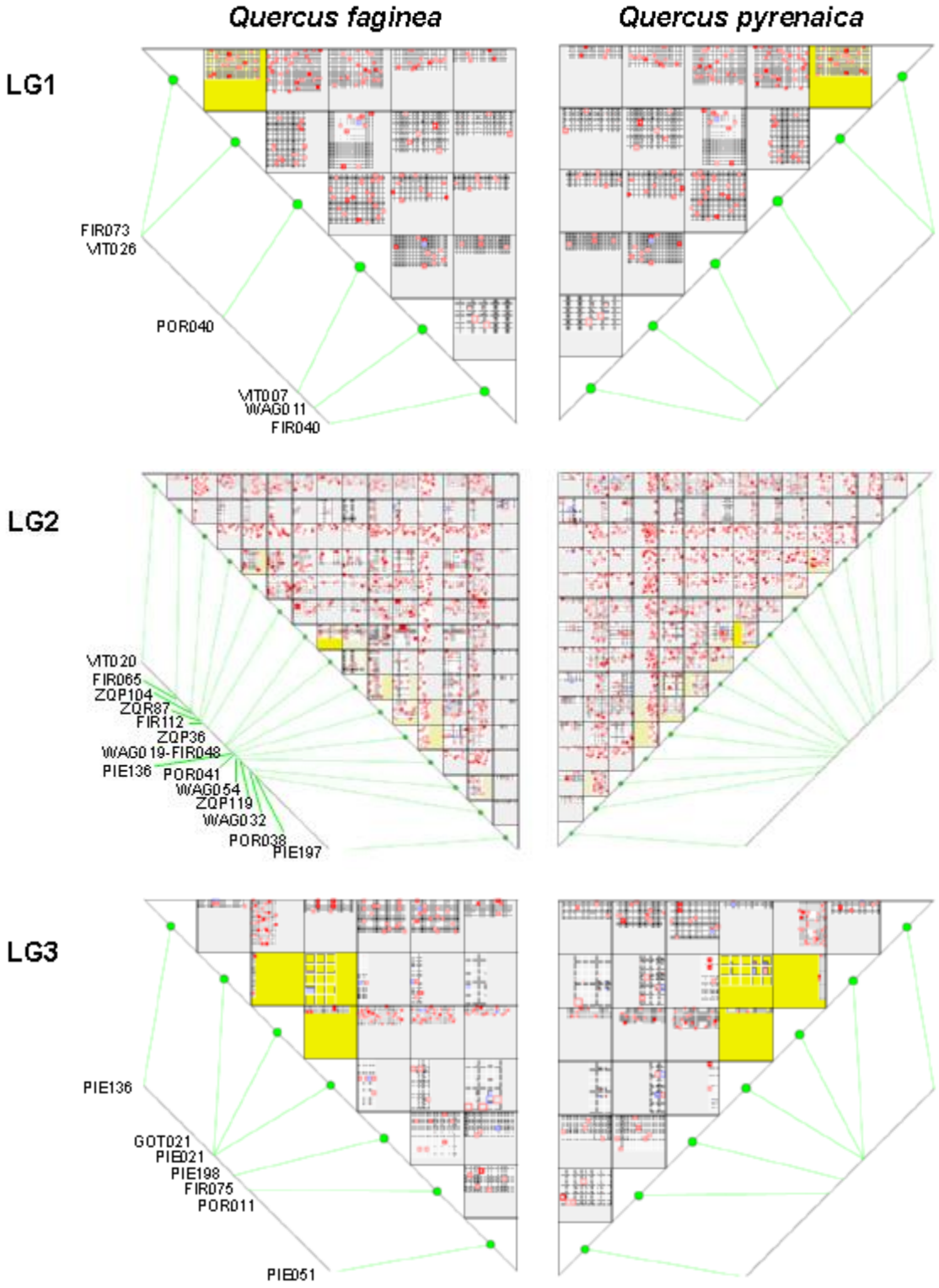


Figure S6-2: (cont.)

Figure S6-2: (cont.)

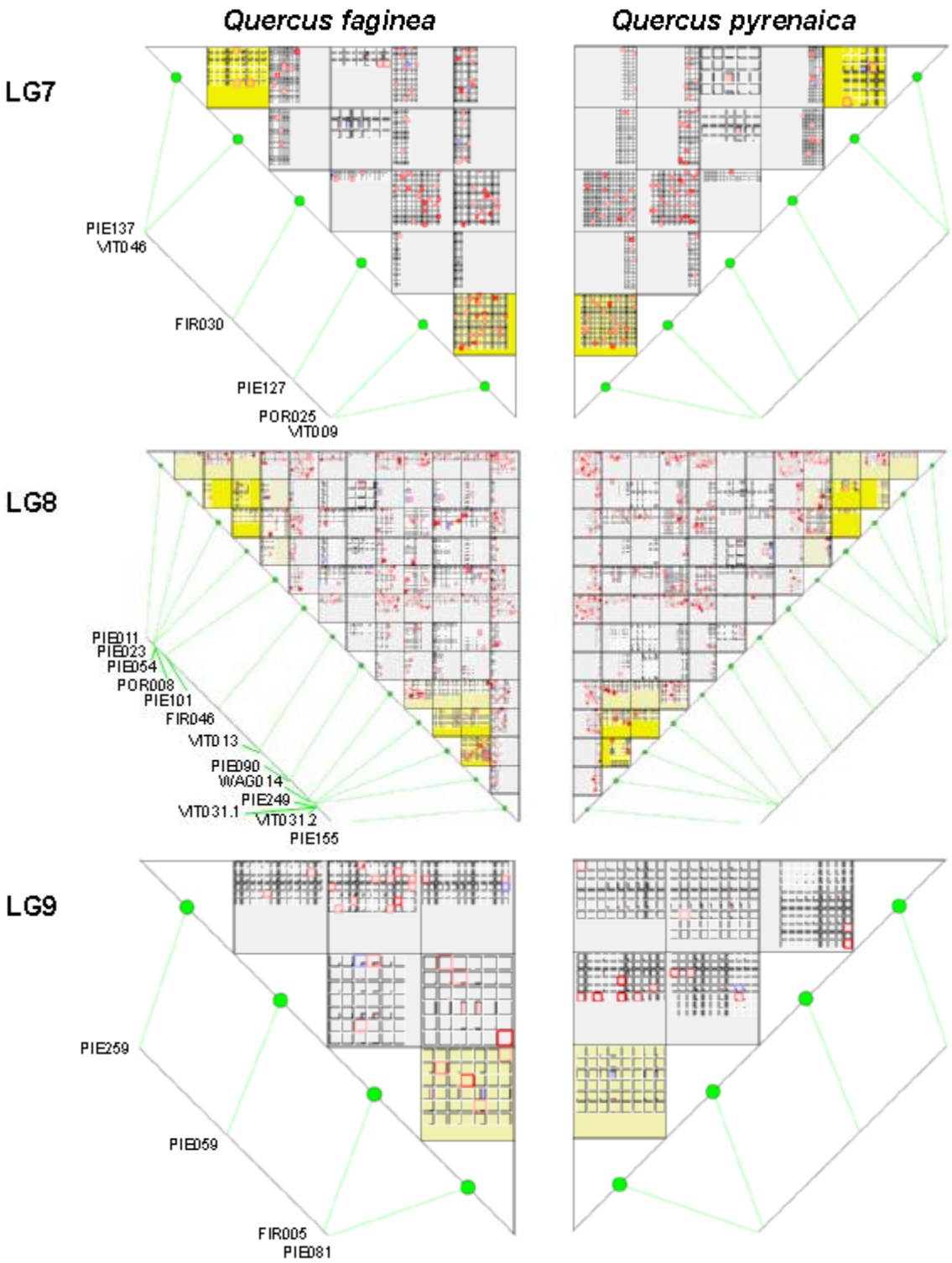


Figure S6-2: (cont.)