Fig. 1 Suppl. PCA with SNP data from genotype-by-sequencing of selection candidates from C0, C1, C2, C3. In the case of C1 selection candidates, these originated from one of five half-sib families (K142, K15, K204, K209, K65). UA: unassigned to half-sib family.
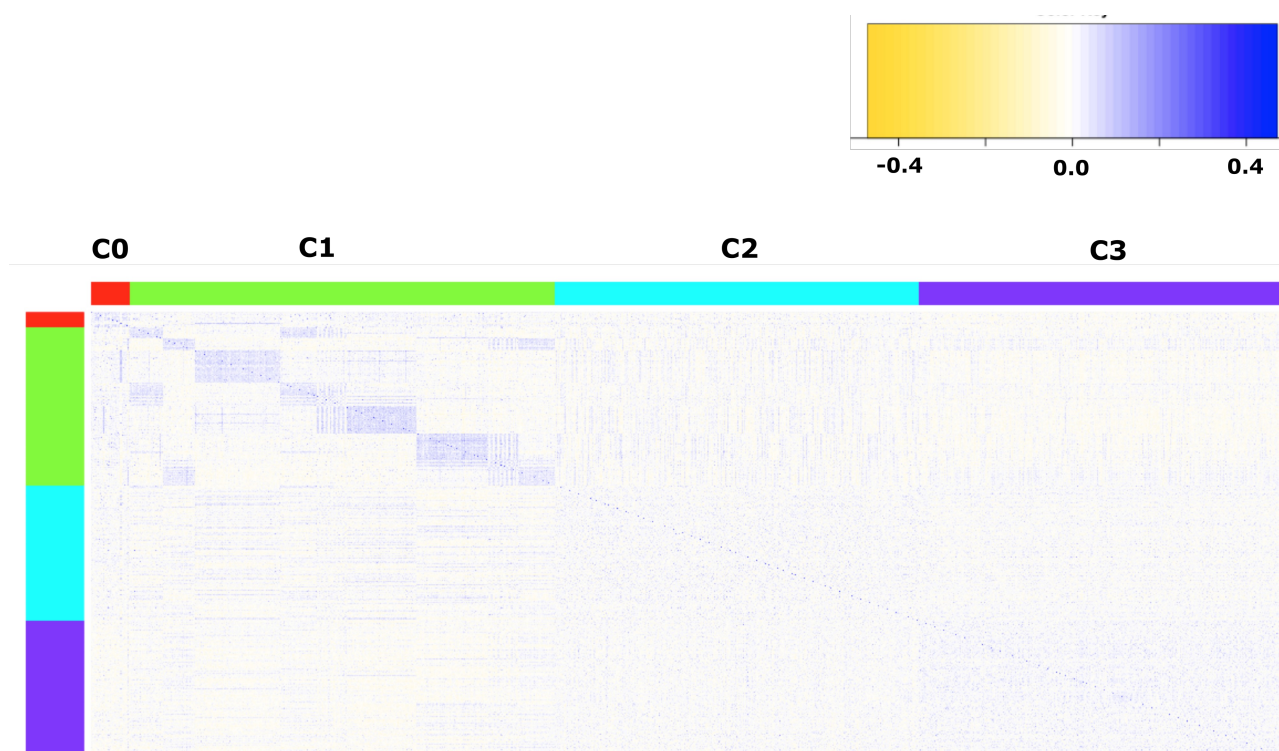
Fig. 2 Suppl. Genomic Relationship Matrix as heatmap. Plants are grouped according to cycle of selection from C_0 (training population) in red to C_3 in purple.