Supplementary Material

Genome-wide association study for crown rust and powdery mildew resistance in an oat collection of commercial varieties and landraces

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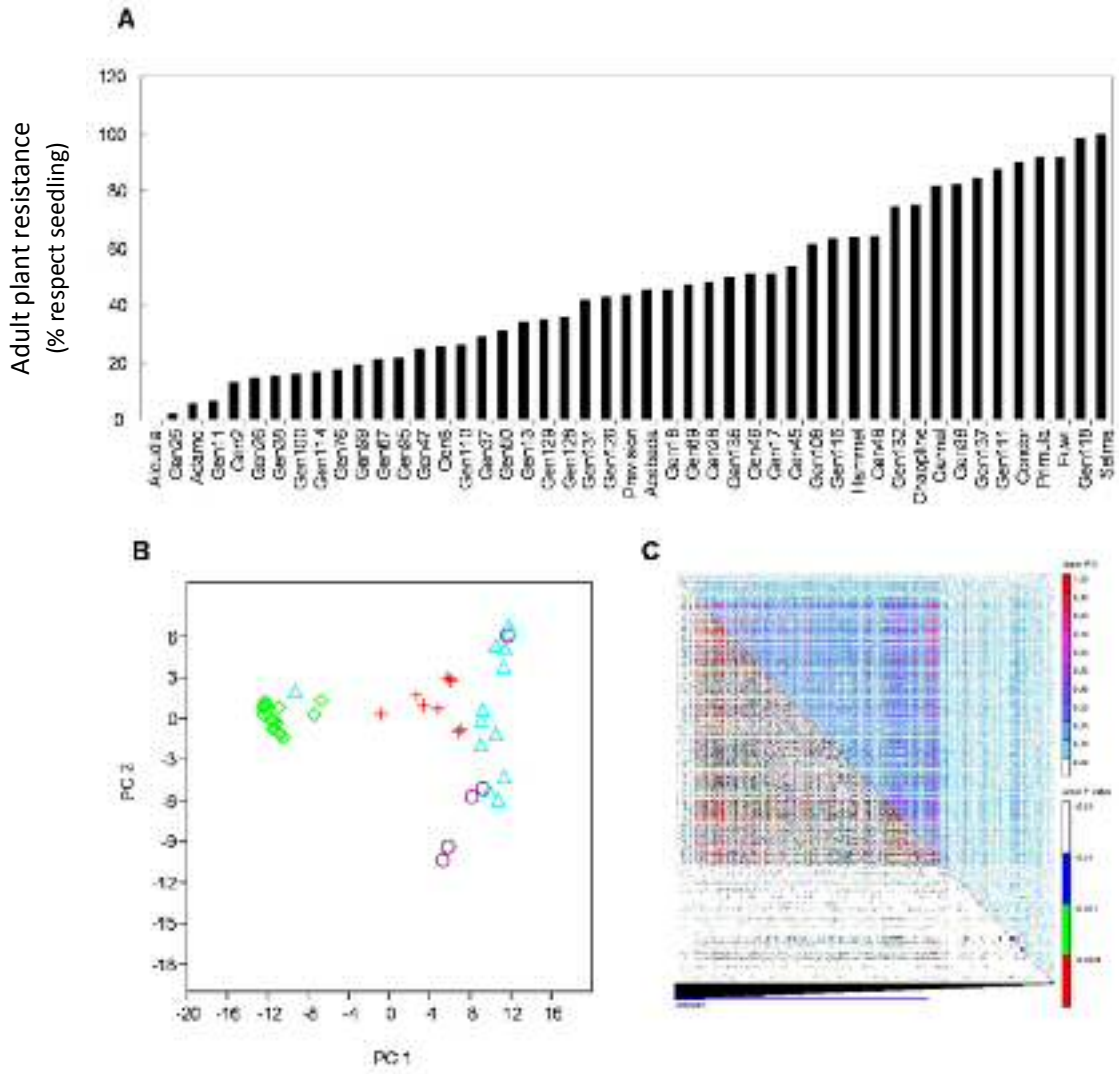
Supplementary Table 1

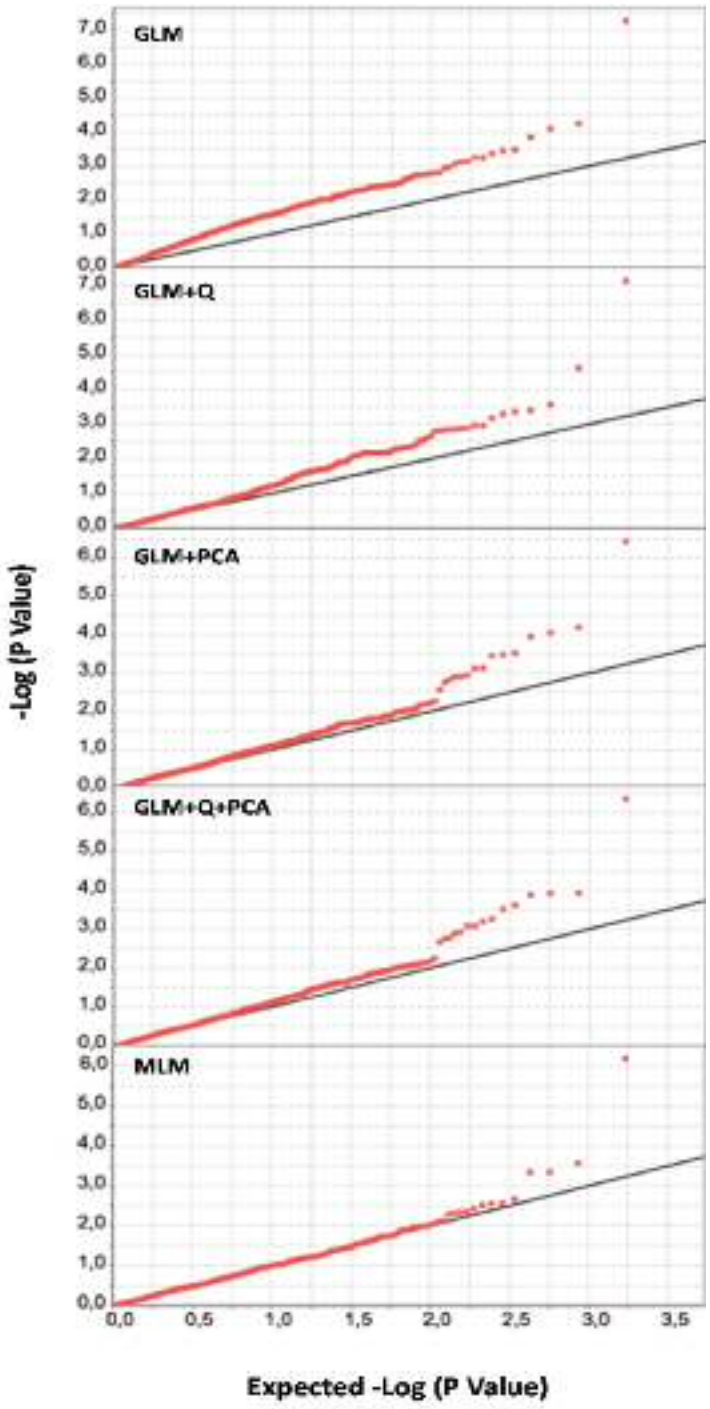
Marker	Accession Nº	Sequence
oPt-11795	FI157472	CTGGATTTGTTGTAAACTGATCATCAGTTGGCCGAGTTGA TGTTGGGCAGTTGCTGTCAAAGGAGACAGAAGCTTGAGC ACCTAGTTCCCATCTTCTCGTTGGGATTCTACCGTGCATTG TGGGTCTTCAGCAAAGGAGTATAGTATATCGTAGGCTATG TAAAACCTGCCCCCCTCGGTCCATTTGTCATGCCAGAAAG ATATTTTGGTACCATTTGCCGGGGAGCAGATAGTTGCGCT CTGCACCGTGGGTATCAATGCAGGTAGCGATTTCCAGAAG GGTGTGTCTTGCTGTGATGTTGTGTGTGGTATAGTGCC
oPt-15665	FI158620	CAGGCTAGTAACAAGCAACAAGGGAATATGAAGAAGGGA AAAGTTGTGGTGATTGAAGATCAAAACTCAGGGAGAGAG AATCAGTTAAGGTCAAAAGATAGTTACCTGGATGTGGTCT ACTTTACCTGTGGGGAACCTGGCCATAACAAGTCTCAGTT CCCCTCTGCTCCTTTTTTTTCATATGCAAGATGGTAAACCA CAAGGAAGACAAGTGTCCAGTGAGGAAGCTAGCTTTGCC TGCTGCCAAGCTTTATGGAAAAGCTGCTCAAGGGCTGGAA TTCTTCCATGTGGAGGTTCCTGAAAGCTACAACAATGATA TGGGAGCAAAAAATGTGGGGATTGTGTTCATTGAGGCTG GTGAAATCAACAAAGAGGAGCTGGCTCAGGAGTTTGTAG TTATCTACAAAACTACTTGGCCTTGGCAGATTAGGCAGCT AGATGATTGGTCCTTTCTTGTCAAGTTCCCTCCCCACCTCC CAGTAGAAGATGTAGTTGGTTATCCATGCTTTGGCCTAGT GAAAGATGGGGTTACTGTAAATGTGGAAGTCTAGGATGG GGAA
oPt-14317	FI159214	TGCAGGCAGAGGCAGTGTTTATCCTTGAGGGAGTCCATCC CGAGCAGCGGCATGGCAGCCTGGCCGTTCTGTCCCACTGC CACTGGTATTTCAGCTAGCATCCACAGGCGGCTGGTGTGG GGAGTGCTGGAAATACTAGTCATGTCACCGTCCAGTGTTA GTAGCATTTCTTGAAAGGGAAAAGTCAGCACACACACAC TCTTGGCGCCAGCTCCAGGATCCTCGCTTCTCCACAGGGG GGATGCTGTCGCCAGCTACTAGCTGTAGCCCAGCACTCAC TCACCACTACCCGGGAGTAGGAGTGCGTGATGTTTGCGGC GTGGTGAAAGGTCACATGGAATTTGCCCGATTTGGAGCCT GCATGCCATGCCACGGGGATCATGTGATCTTATTTTTTGT CTTGCTTTAATTTGAGGAGGAGTTGATGTTACATTACAAT AAGTCAATAATGATAATGATCCGGCCGCGTGGGTATCTTG TGATCCTGGTTGGTGGTTTGTTCCGTTTTGTTCGCATCATC CAAAAACTC

oPt-5014	FI159708	GTCTAGGGGTTAGCGGCTGCAAAGGTACTTACGTCCAAAT CCTTCAGTATGCAGTCGCTAAAGAGCTCAATGCACGCAAC GTGGAATCCAGCAAGGGAAATCTCGTTCACTGCTGCGGA AGAAAATCTGTTCGTACTACAAGCCCAATGTCTAGGGGAC TAGAACTGTATGATGCTGGAGGGCACTTGGCTCTTTAGGG AATGTGCTCTGATGGTTGAACCTTTTGATGGGGCAACAAC GGTGCAGTCATACAGAGAGGTTCCCAAGCATGGGTTCAA ATCCACAAATTACCATCACTATTTTGCAAGAAGCAGGTTC TTGATCAATTGGCAAGCAGAGTGGGAGAACTGATATCTAC TGATTTGACCCCTGTTCAGATGCGTACAGGTGTGTTTCATC GGGTACGCGTGAAGCTTAACTCTGCAAAACTGCTTATGCG CTTTGTGTCACTAGCTATTGAAGGAAGCCCGAGGATGTTT

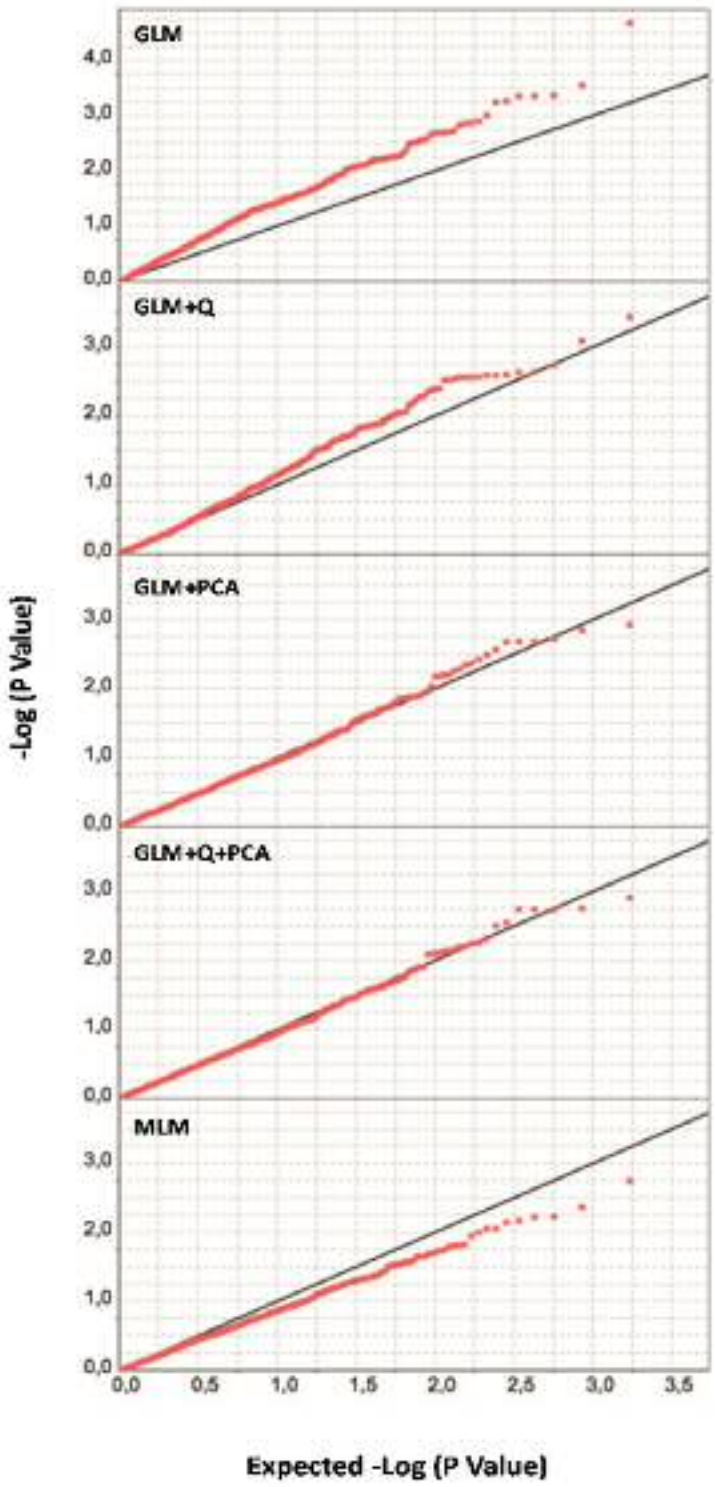
Supplementary Figures



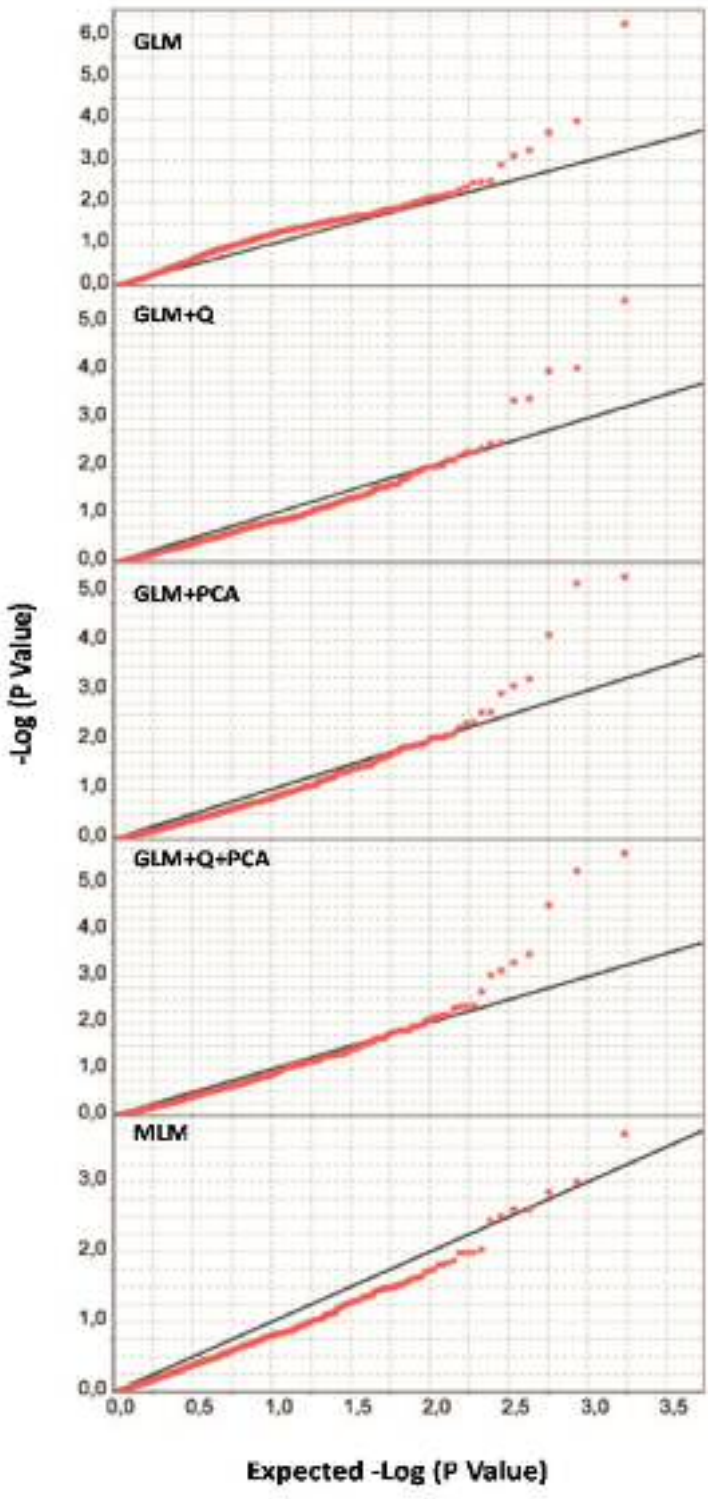
Supplementary Figure 1. **A.** Adult Plant resistance, measured as percentage of the seedling resistance in a subset of 54 oat accessions. **B.** Scatterplot of Principal Component Analysis scores of components 1 and 2 based on 1587 DArT and SSR markers used in this study in the oat subset. Represented are the genotypes belonging to cluster 1(red), cluster 2 (green), cluster 3 (violet) and cluster 4 (blue). **C.** Linkage disequilibrium matrix in the oat subset. Pair-wise LD values of polymorphic sites displaying r^2 above the diagonal and the corresponding *p*-values from rapid 1000 shuffle permutation test below the diagonal. Each cell represent the comparison of two pairs of marker sites with the color codes for the presence of significant LD. Colored bar code for the significance threshold levels in both diagonals is shown.



Supplementary Figure 2. Distribution of p values for the different models used in this study for the association between markers and rust resistance. Axes represented the expected p values versus the observed p values in the negative log10 scale where the solid line represent the null expectation (absence of type I errror).

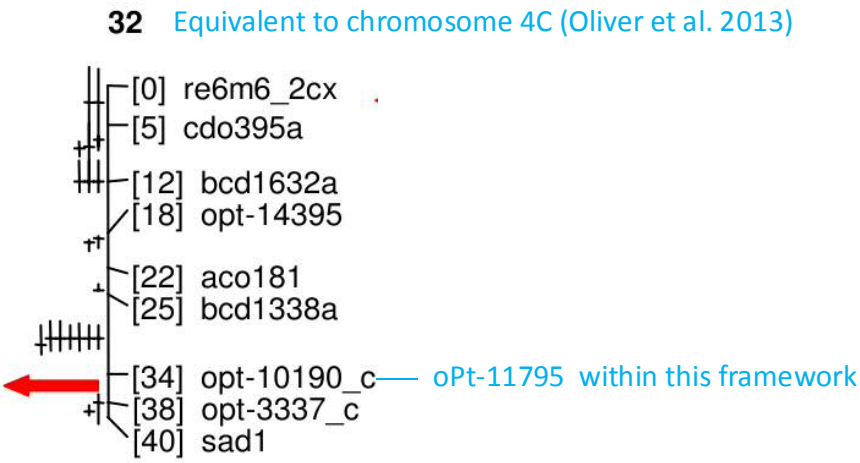


Supplementary Figure 3. Distribution of *p* values for the different models used in this study for the association between markers and powdery mildew resistance at seedling stage. Axes represented the expected *p* values versus the observed *p* values in the negative $\log_{10}$ scale where the solid line represent the null expectation (absence of type I errror).

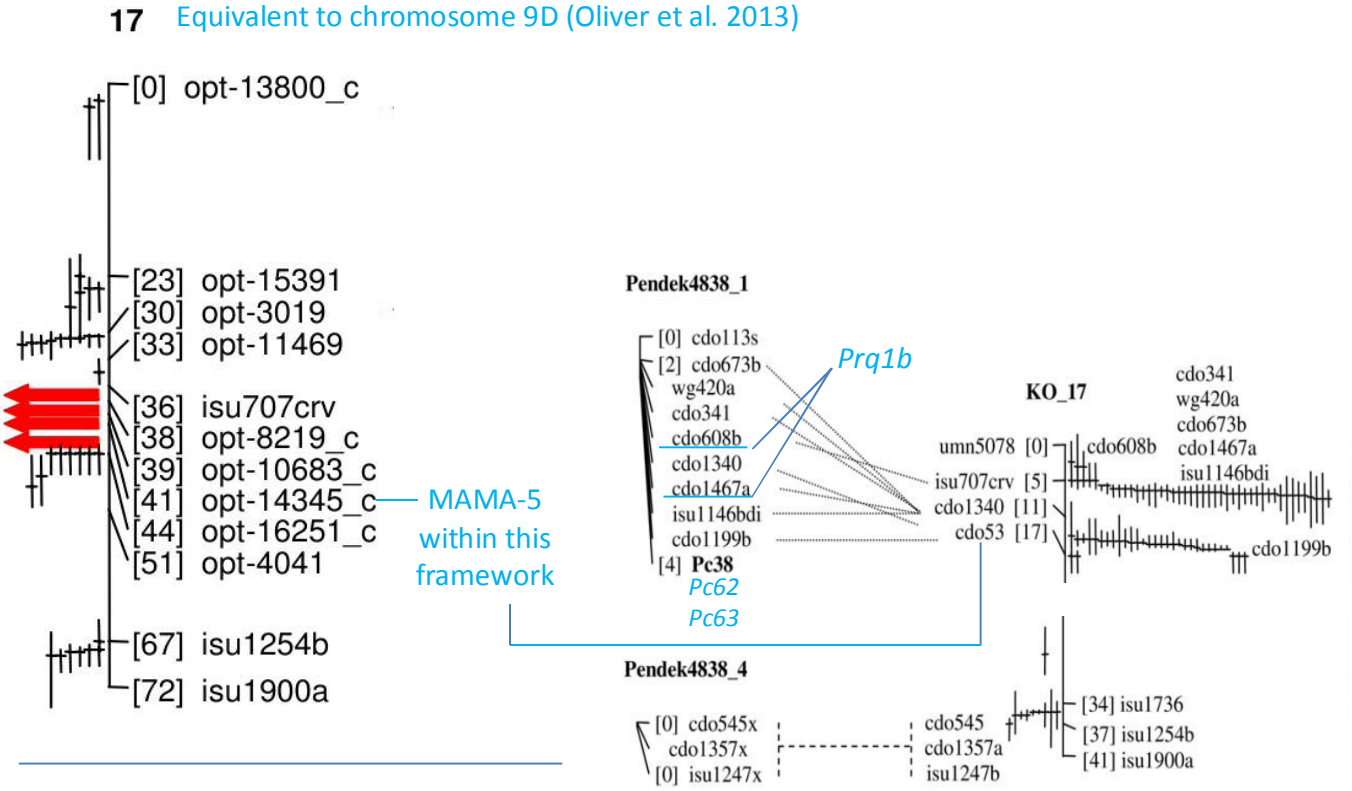


Supplementary Figure 4. Distribution of *p* values for the different models used in this study for the association between markers and powdery mildew resistance at adult plant stage. Axes represented the expected *p* values versus the observed *p* values in the negative $\log_{10}$ scale where the solid line represent the null expectation (absence of type I errror).

Marker oPt-11795



Marker MAMA5



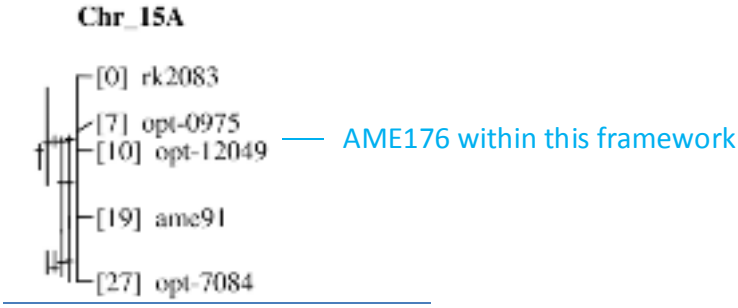
From Tinker et al., 2009

From Wight et al., 2004

Supplementary Figure 5. Chromosomal organization of the different markers highlighted in this study according to localisations of previously reported markers. Cont.

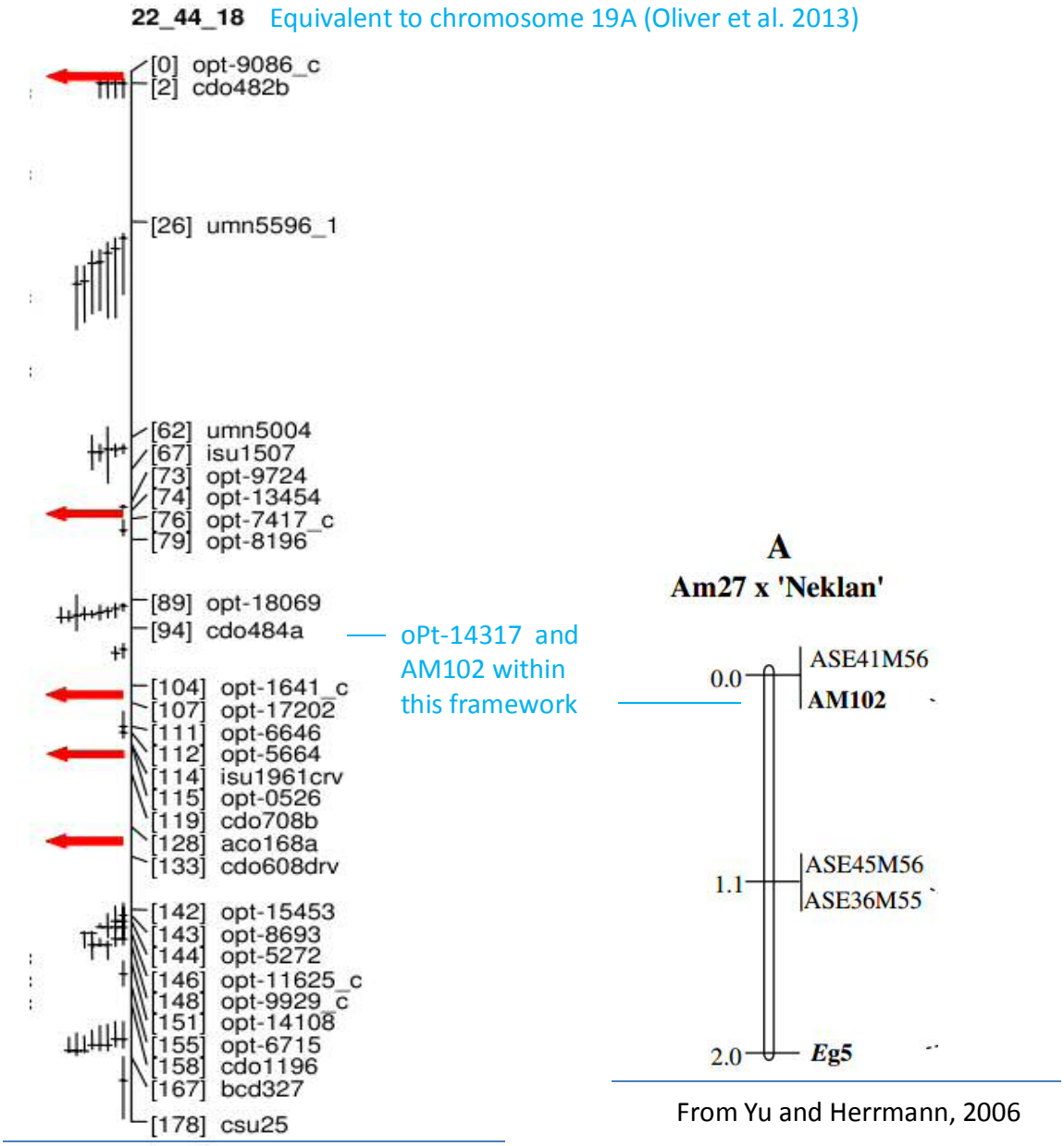
Marker AME176

Supplementary Figure 5. Cont



From He et al., 2013

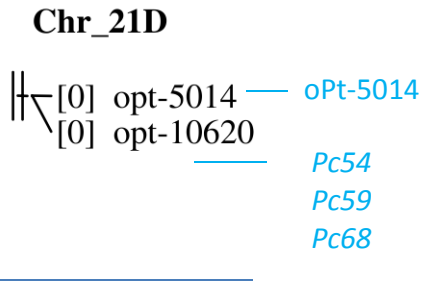
Marker oPt-14317



From Tinker et al., 2009

From Yu and Herrmann, 2006

Marker oPt-5014



From He et al., 2013

Supplementary Figure 5. Cont