



Homoeologous duplicated regions are involved in quantitative resistance of *Brassica napus* to stem canker

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Additional file 1-Table S1: Extent of linkage disequilibrium on each linkage group of *B. napus*

Linkage group	Efficient size of population (Ne)	Extent of LD until the critical value of $r^2=0.2$ (c in cM)
A01	84,21	1,19
A02	39,9	2,51
A03	103,38	0,97
A04	118,1	0,85
A05	70,95	1,41
A06	37,75	2,65
A07	99,55	1
A08	42,2	2,37
A09	82,8	1,21
A10	168,41	0,59
C01	98,97	1,01
C02	58,67	1,7
C03	107,16	0,93
C04	238,31	0,42
C05	55,39	1,81
C06	49,86	2,01
C07	77,28	1,29
C08	131,22	0,76
C09	98,84	1,01
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A genome	73,78	1,36
C genome	89,85	1,11
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Whole genome	78,26	1,28