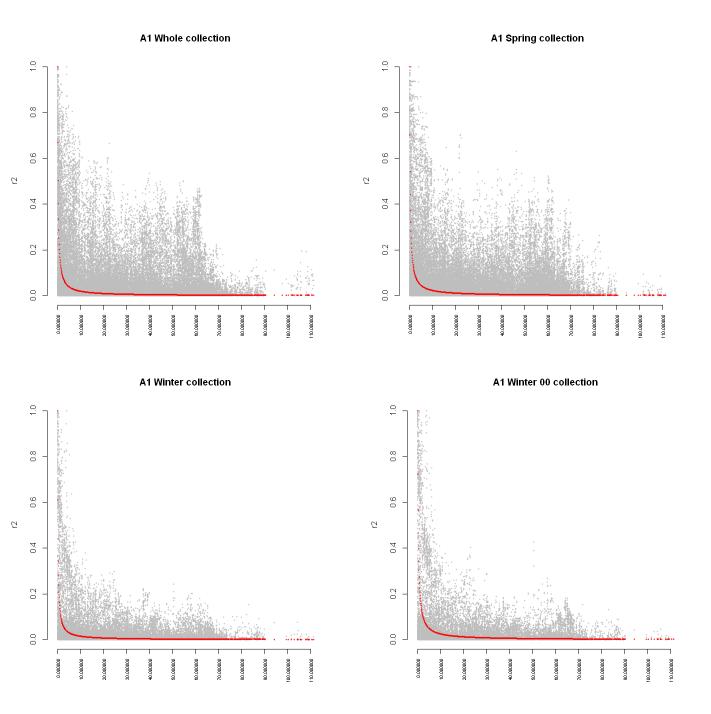
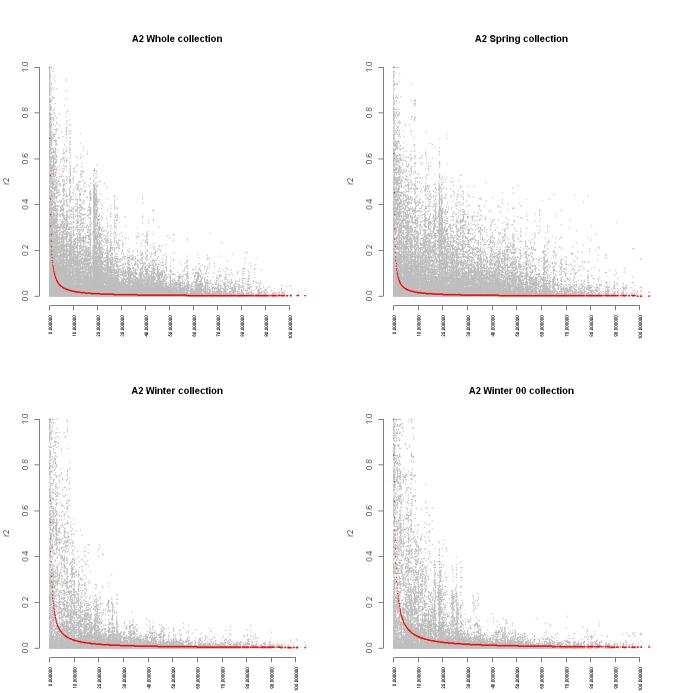
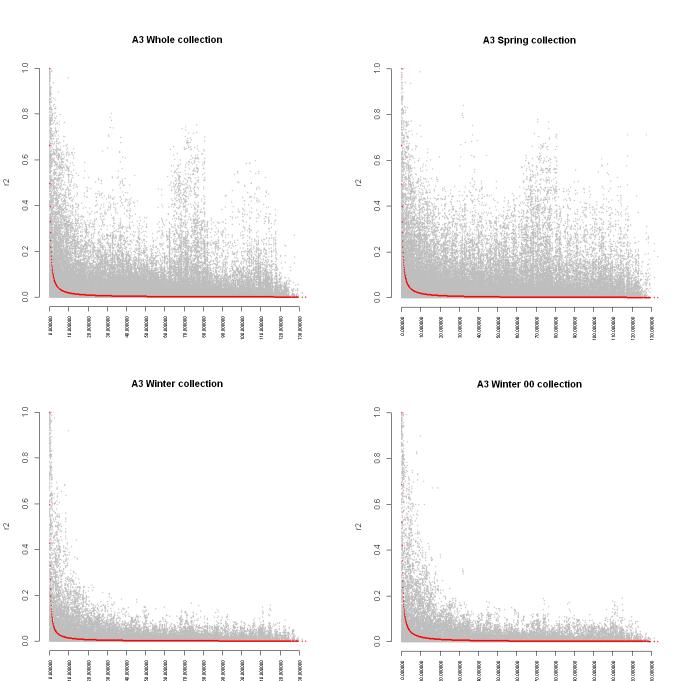
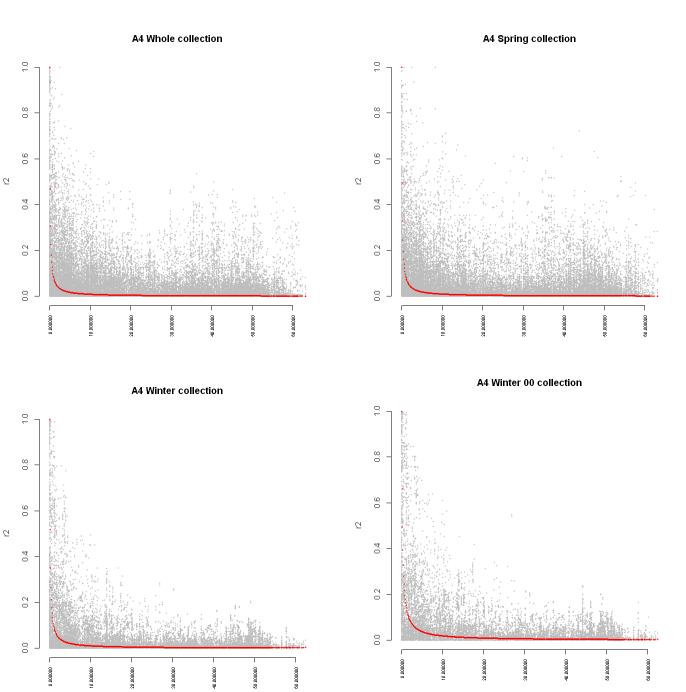
**Supplementary figure 4:** Plots of  $r^2$  as a function of genetic distance (in cM) between pairs of SNPs on each linkage group in the whole, spring, winter and « 00 » winter oilseed rape collections. Red curves show the non-inear regressions trend line of  $r^2$  versus genetic distance



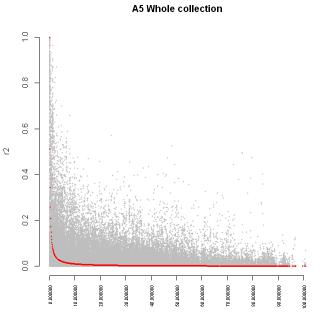


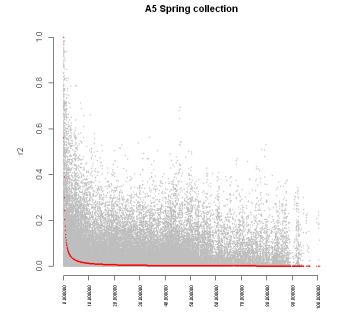


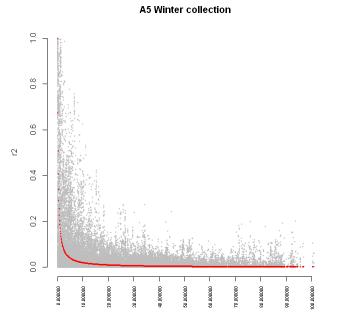


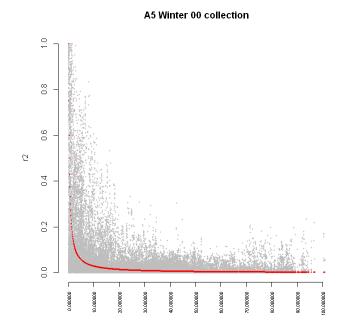
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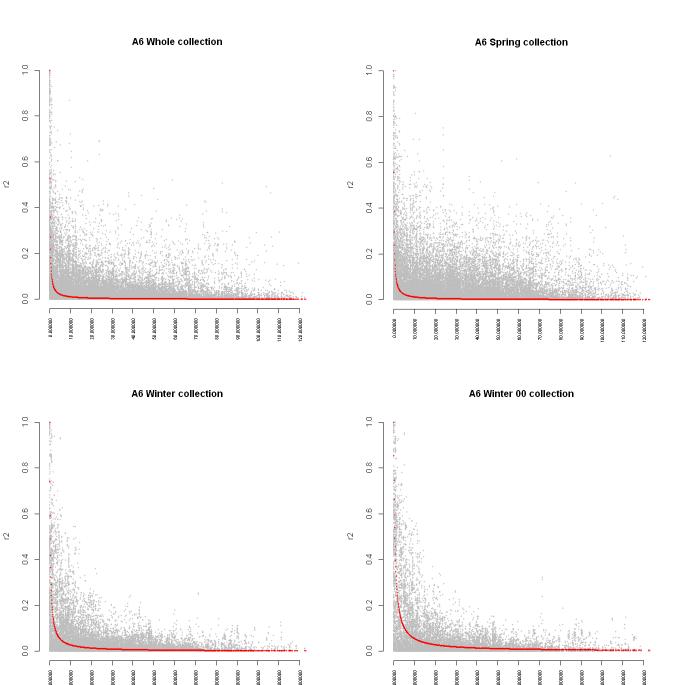
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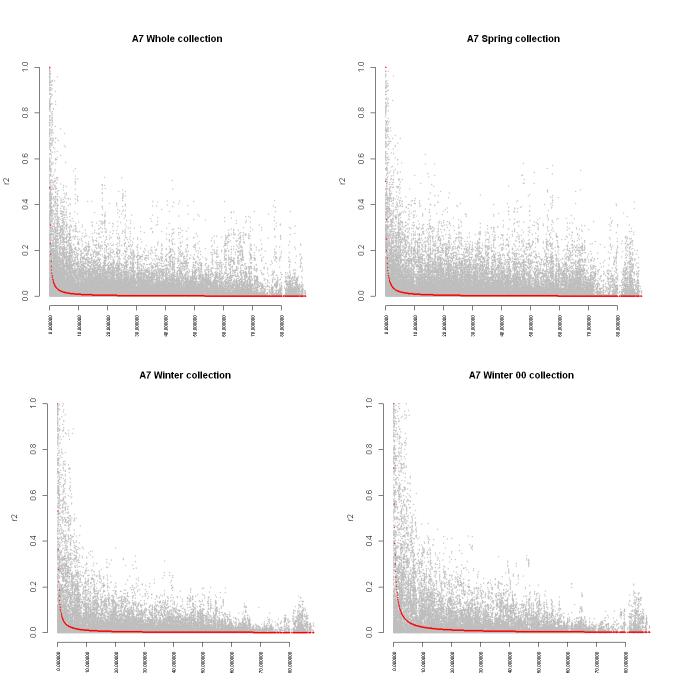


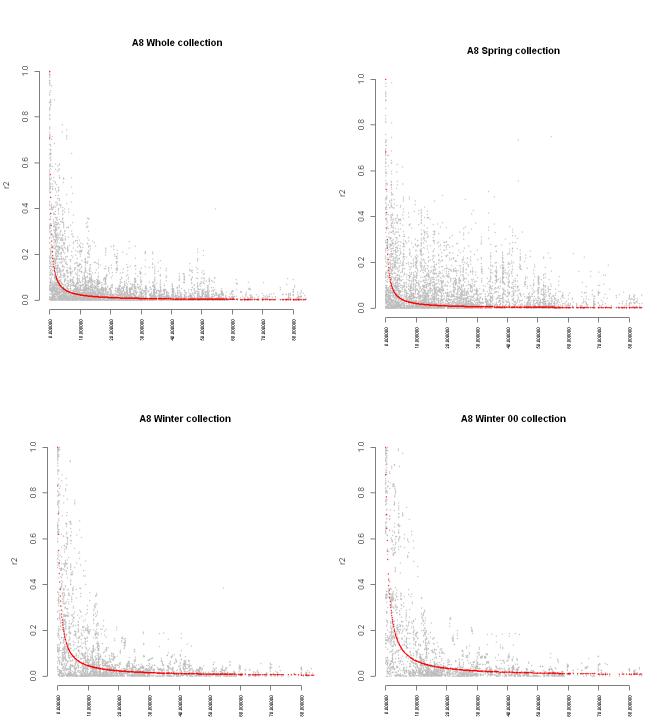


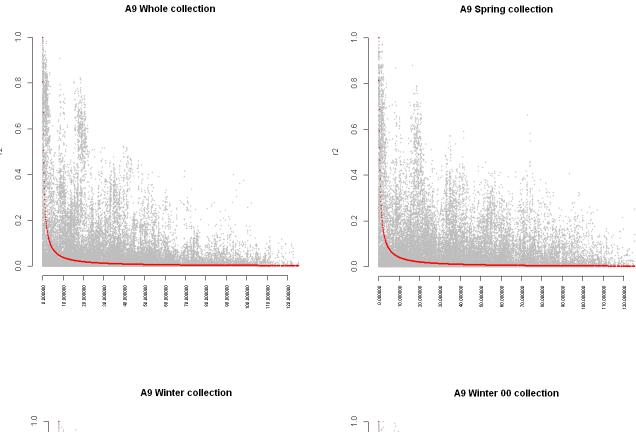


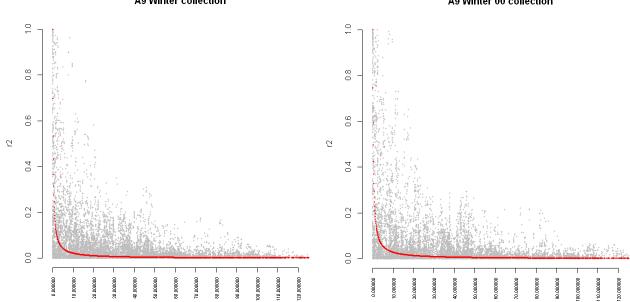


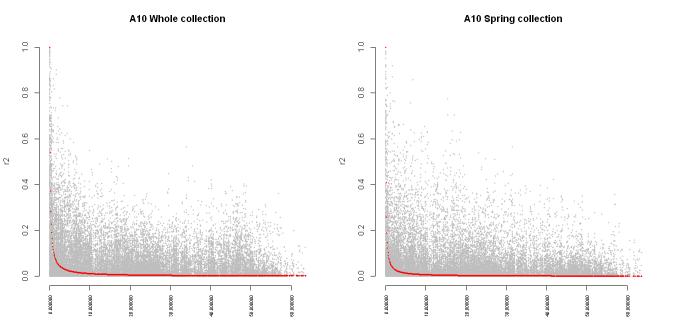


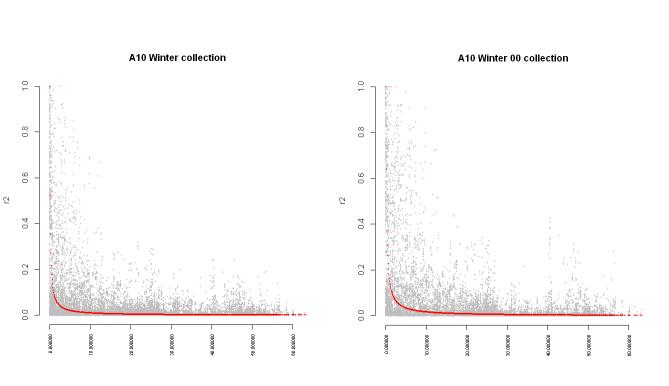


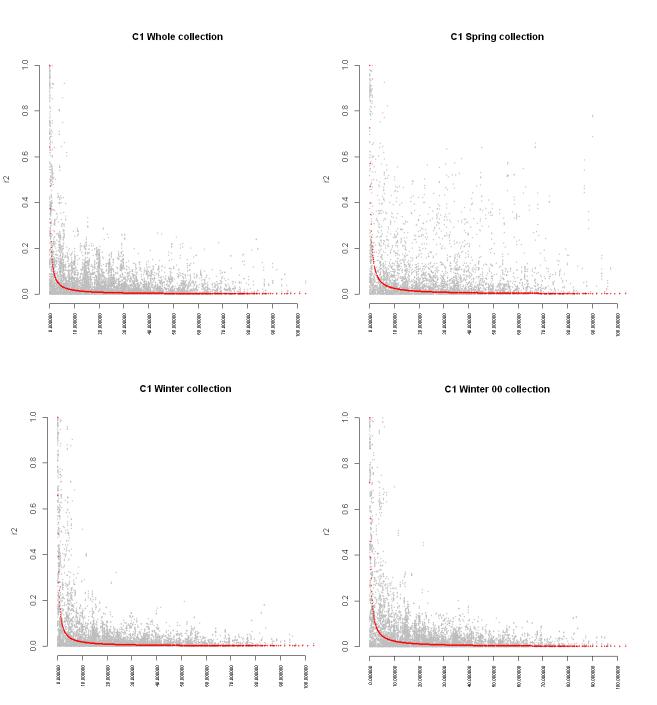


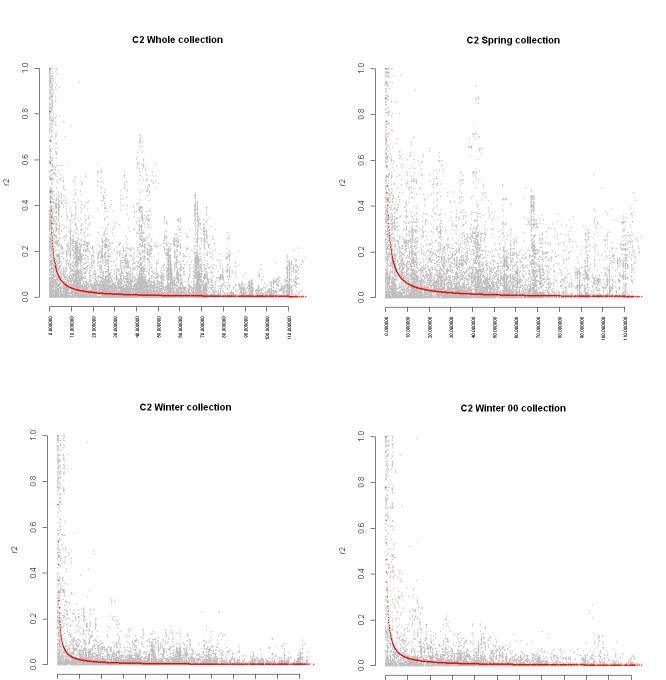


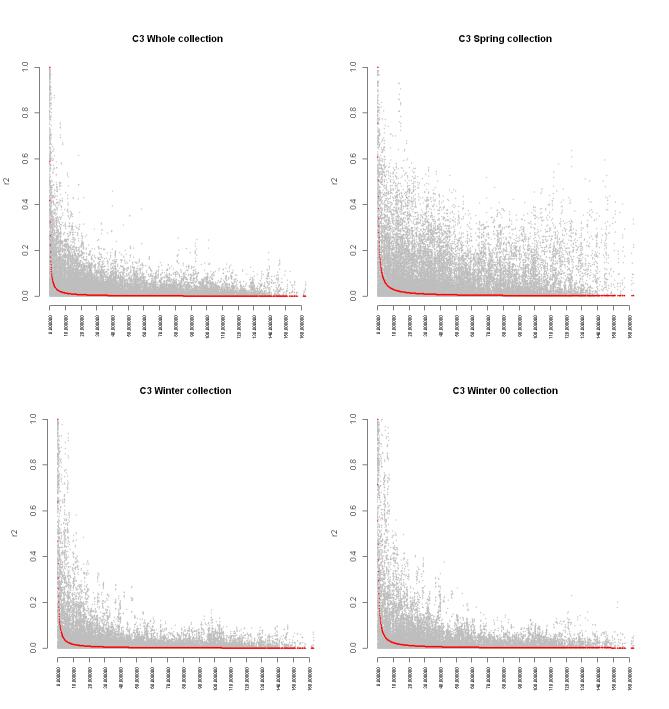


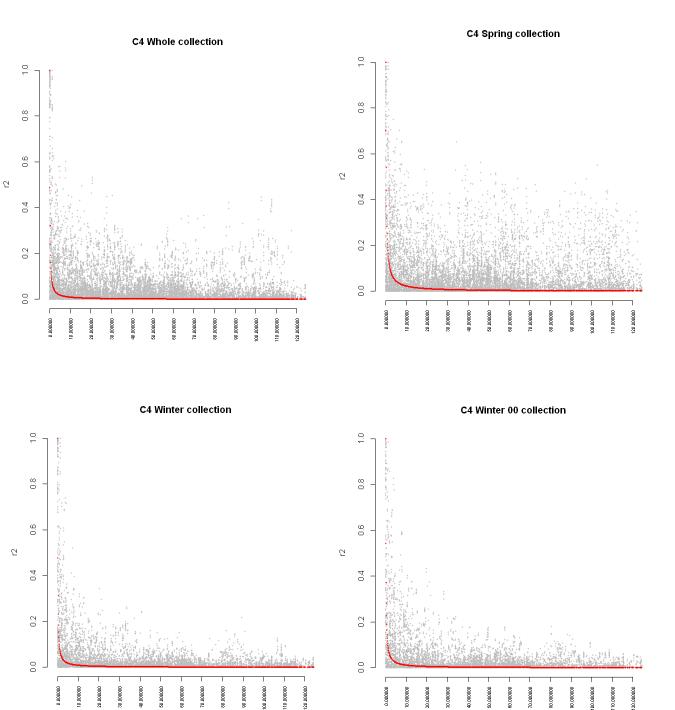


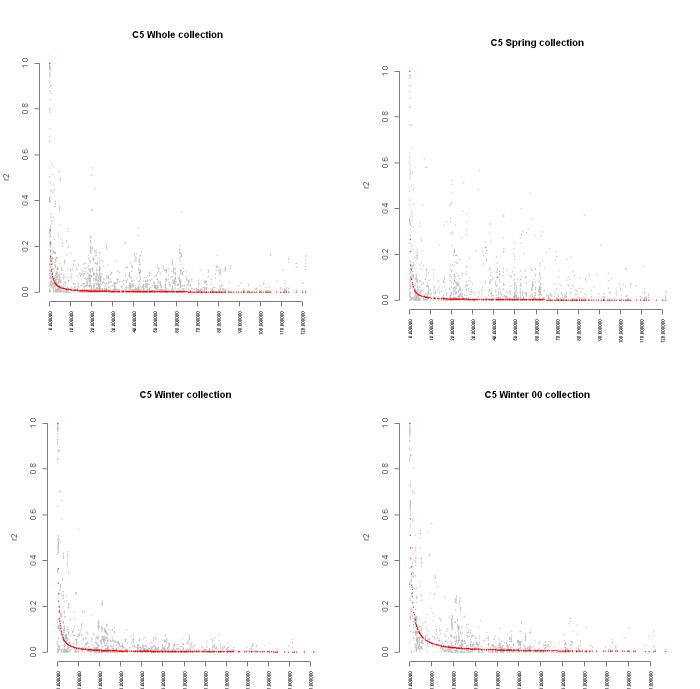


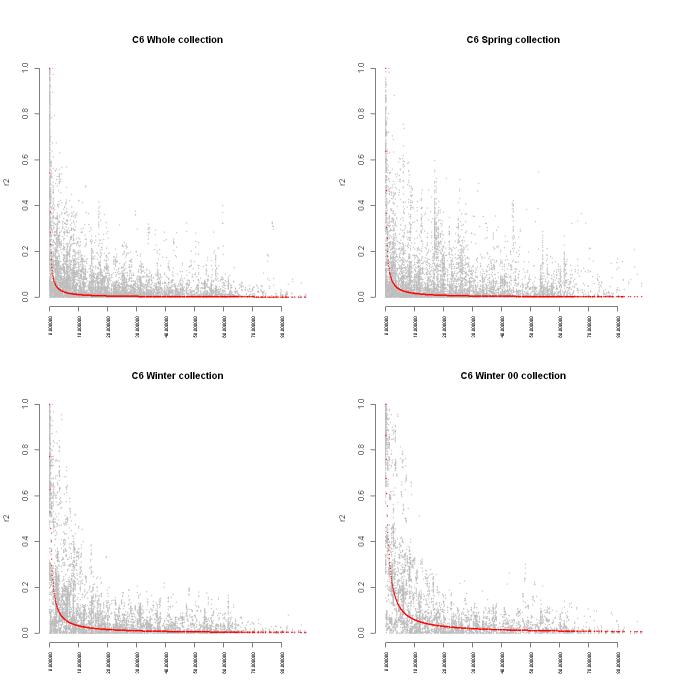


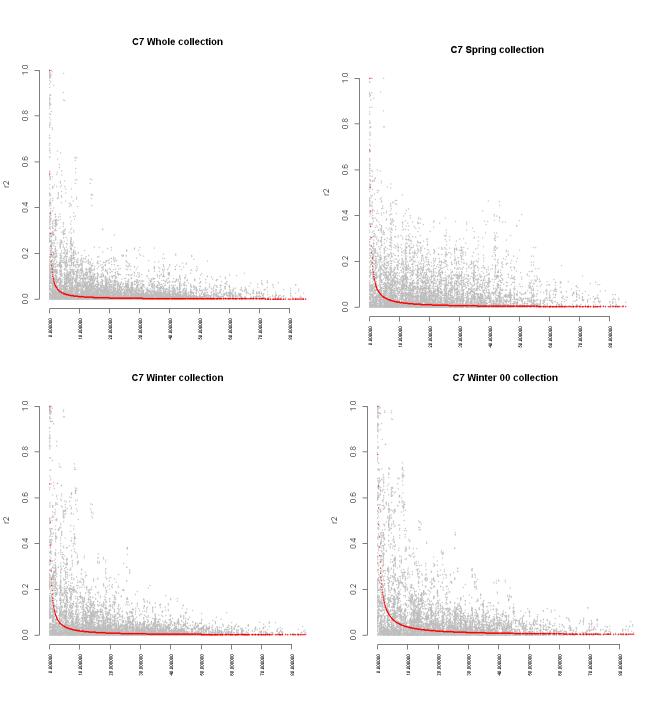


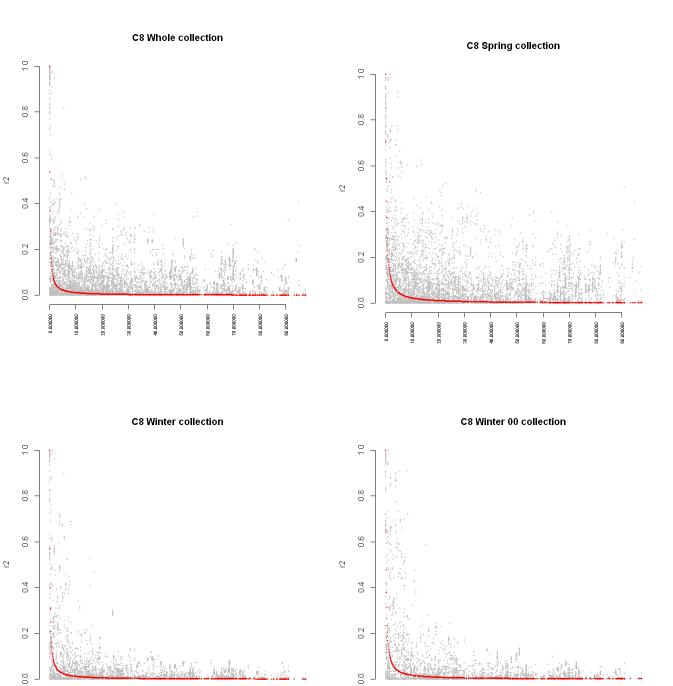












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