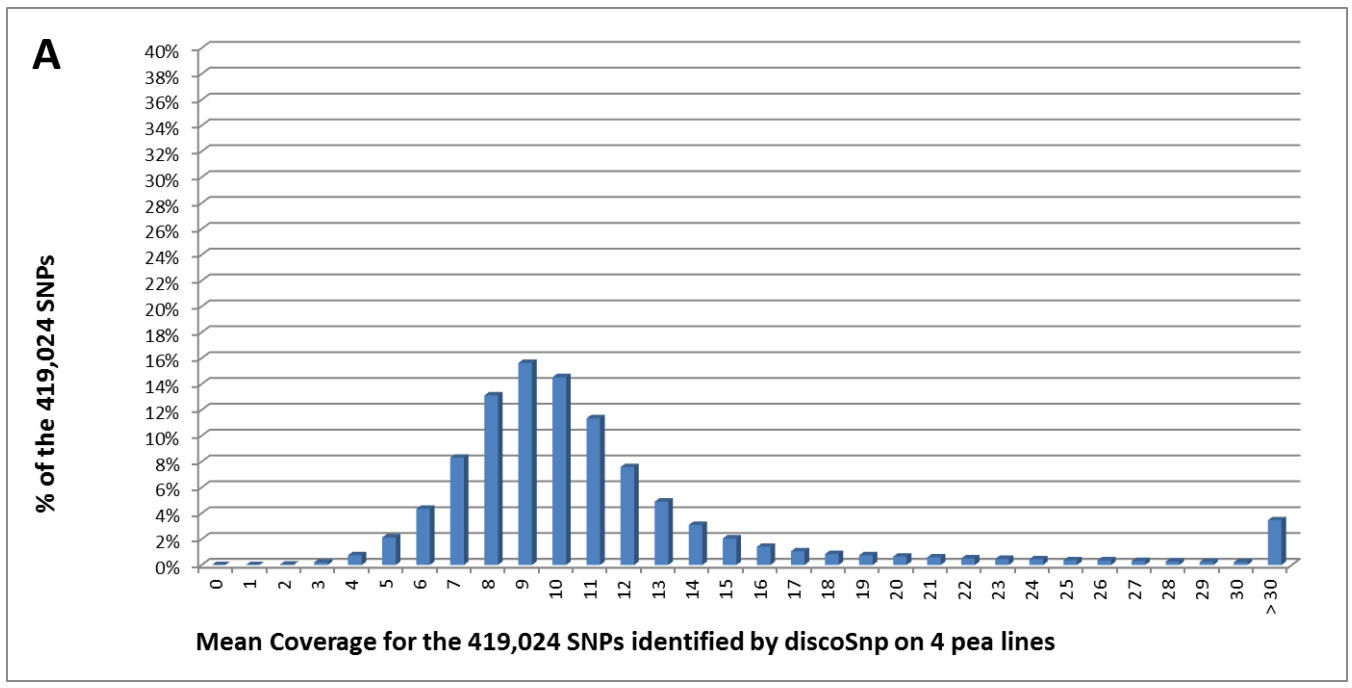
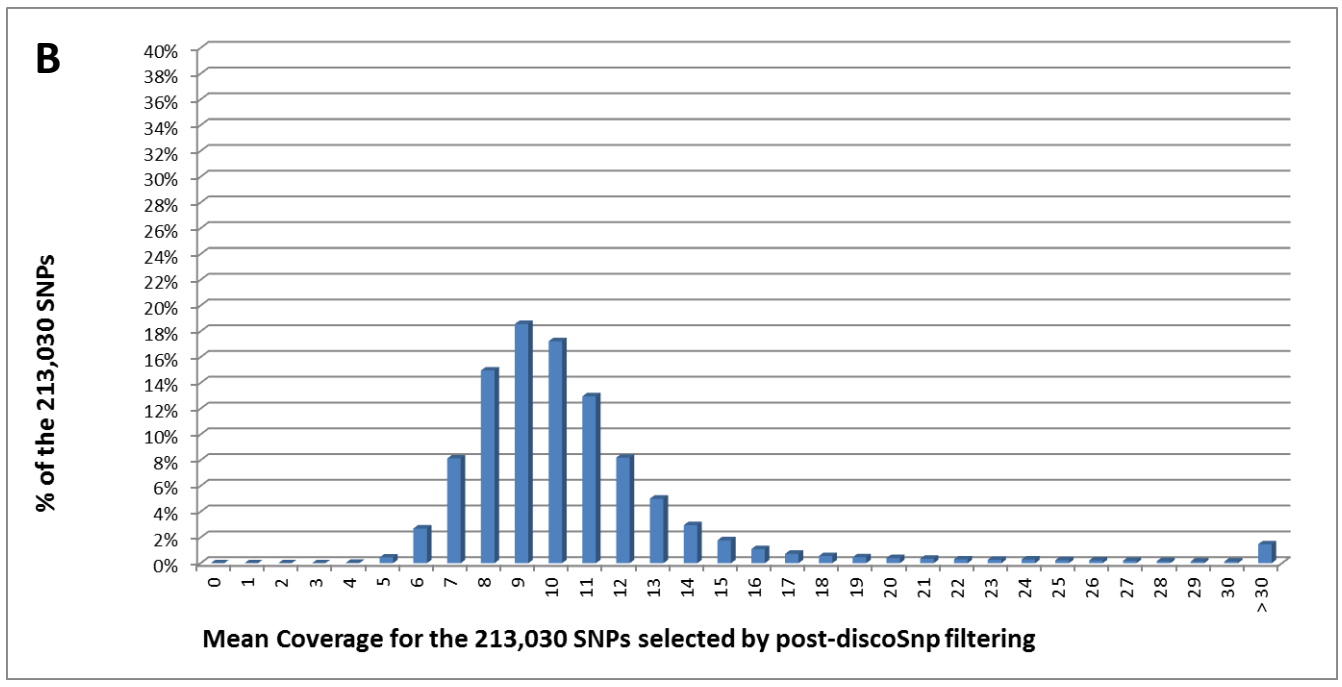
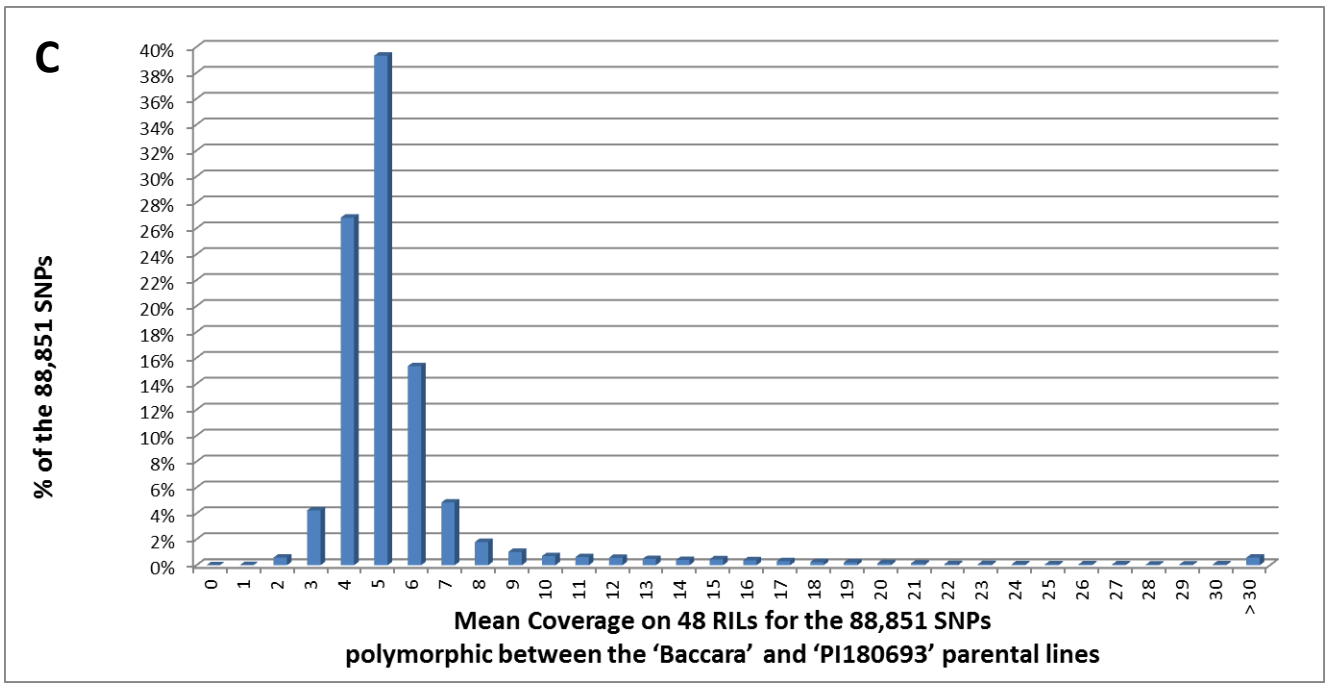
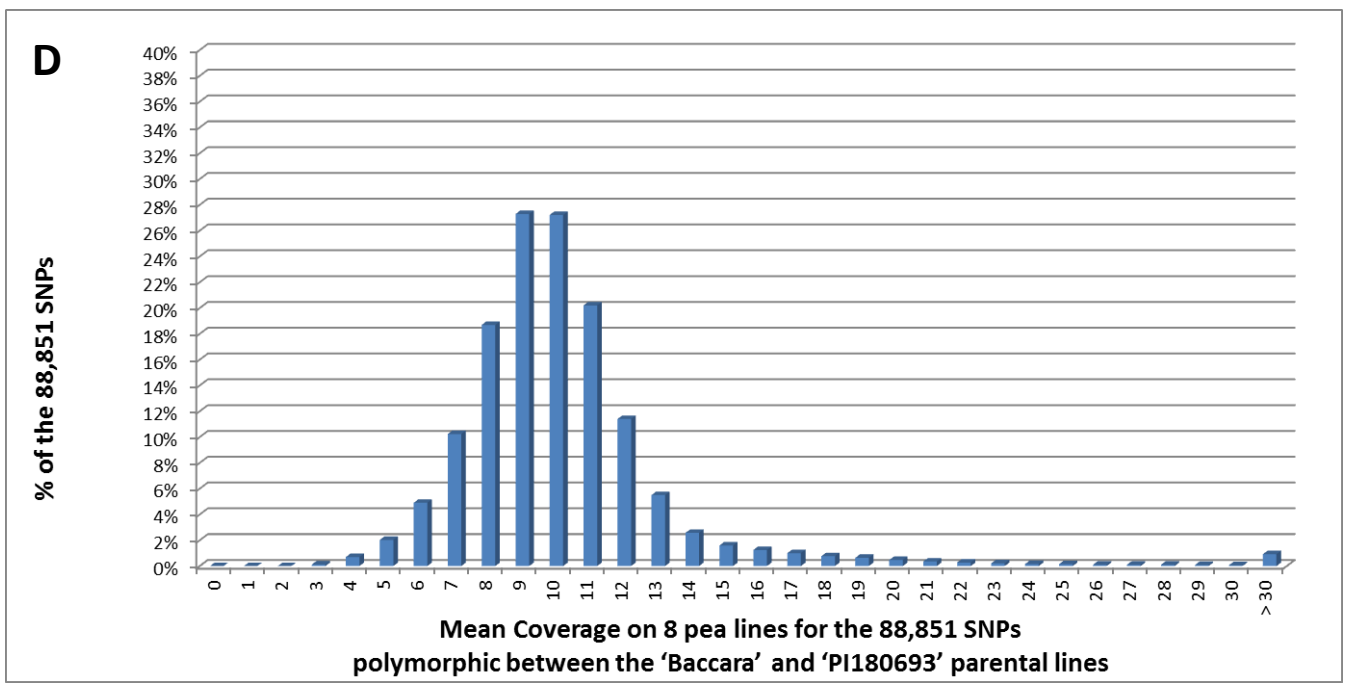

A
% of the 419,024 SNPs
Mean Coverage for the 419,024 SNPs identified by discoSnp on 4 pea lines


B
% of the 213,030 SNPs
Mean Coverage for the 213,030 SNPs selected by post-discoSnp filtering


C
% of the 88,851 SNPs
Mean Coverage on 48 RILs for the 88,851 SNPs
polymorphic between the 'Baccara' and 'PI180693' parental lines


D
% of the 88,851 SNPs
Mean Coverage on 8 pea lines for the 88,851 SNPs
polymorphic between the 'Baccara' and 'PI180693' parental lines