Additional file 2: Bubble plot showing the pair-wise percentage of overlapping markers called by the different pipelines. This plot complements the visualization in Figure 3, showing all pair-wise proportions of shared SNPs among the five evaluated pipelines. This plot indicates, for example, that GBS-SNP-CROP-MR01 calls 33.7% (1,844 SNPs) of the 5,471 SNPs called by GBS-SNP-CROP-RG; but those same 1,844 SNPs comprise only 8.6% of the 21,318 SNPs called by GBS-SNP-CROP-MR01.