

Fig. S3 Comparison of the CAPS marker results obtained for the same set of rapeseed plants using automatic FastPrep / CTAB and FastPrep / FastDNA SPIN Kit DNA extraction procedures. For each analyzed plant, two samples (designated with letters P and D, which are explained in Fig. 2) representing two steps of the CAPS protocol were applied on the gel

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Cleaved amplified polymorphic sequences (CAPS) marker for identification of two mutant alleles of the rapeseed *BnaA.FAD2* gene Marcin Matuszczak, Stanisław Spasibionek, Katarzyna Gacek, Iwona Bartkowiak-Broda

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