

Fig. S5 Comparison of the CAPS marker results obtained for the same set of rapeseed plants using the initial standard method (with the purification step) and the simplified method (without the purification step). 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 (for better comparison, only two steps are shown for the initial standard method)

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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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