

Fig. S8 Results of testing the ability of the studied CAPS marker to detect the alternative source of high oleic acid content in rapeseed. The analyzed plants originated from the new series of mutagenesis experiments which could generate the new alleles of the *BnaA.FAD2* gene. All the tested DNA samples are described in the figure. 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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