Study of the lines of common wheat of breeding of National Center of Grain named after P.P. Lukyanenko on allele variants of *Waxy*-genes

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The aim of this work was the molecular identification of promising genotypes common wheat by allele variants of Waxy genes (Wx) to create varieties with improved technological qualities of grain. 352 lines of common wheat were studied by allelic variants of genes WxA1, WxB1, WxD1. All lines were obtained in the department of breeding and seed production of wheat and triticale in the National Center of Grain named after P.P. Lukyanenko, by crossing with a mutant line carrying null alleles WxA1b, WxB1b and WxD1b. There were identified 205 lines carrying the wild type allele WxA1, 229 lines with the wild-type allele of the gene WxB1 and 249 with the wild-type allele of the gene WxD1. 64 lines carried functional allele WxB1e, different from that wild type. The lines carrying zero alleles of genes WxA1, WxB1 were selected (147 and 58 respectively). 13 lines, showing the heterozygous condition for the gene WxD1, and one line for the WxB1 gene, were identified. Full technological evaluation was given to 57 promising lines and several standard varieties. The obtained results show that the lines carrying zero-allele WxA1b or WxB1b, as well as a combination of alleles WxA1b+WxB1b, are generally characterized by high protein content (15.3, 15.4, 15.5 % respectively), gluten (28.3, 28.6, 30.3 %). All three groups of lines are close to the "strong" wheat by the strength of flour (285 alveograph units, 270 a.u., 279 a. u.). The lines with the identified null alleles of the Wx genes are distinguished by a high water absorption capacity compared to the Tanya variety, which does not have these alleles in its genotype. The optimal combination of liquefaction indicators and high valorimetric evaluation probably contributes to the formation of high-volume bread with a high overall baking rating.