Identification of the stem rust resistance genes in the introgression lines of spring bread wheat using molecular markers

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A total of the 58 introgression lines and 11 cultivars of spring bread wheat developed by Agricultural Research Institute of South-East Region and cultivated in the Volga Region were analyzed. The lines were obtained with the participation of CIMMYT synthetics, durum wheat cultivars, direct crossing with alien species such as Agropyron elongatum, Ag. intermedium, Aegilops tauschii, different species of the genus Triticum L., Secale cereale and triticale Satu. Cultivars and lines were evaluated for resistance to Lysogorsk and Omsk stem rust pathogen populations (Puccinia graminis f. sp. tritici) and to the Ug99 race group in Kenya (KARI) as well as analyzed for the presence of the known Sr resistance genes (Sr22, Sr25, Sr26, Sr31, Sr35, Sr36, Sr38, Sr39) using molecular markers. The gene Sr31 remained effective to the local pathogen population. The 26 wheat lines out of 58 were resistant to all local pathogen populations taken into analysis and the 15 introgression lines were resistant to stem rust in Kenya. The genes Sr31/Lr26, Sr25/Lr19, Sr22, Sr35 and Sr38/Lr37 were identified in the introgression lines. The gene Sr31/Lr26 was identified in 13 lines. All lines carrying 1RS.1BL translocation (Sr31/Lr26) were resistant to all local pathogen populations taken into analysis. The gene Sr25/Lr19 was identified in 40 lines. The genes combination Sr31/Lr26+Sr25/Lr19 was identified in 10 lines. The gene Sr22 was identified in 2 lines, this fact will be checked in further work, gene Sr35 – in one line and gene Sr38 – in two lines. The genes combinations Sr38/Lr37 + Sr25/Lr19, Sr35 + Sr25/Lr19 were identified. The lines with genes combinations Sr38/Lr37 + Sr25/Lr19 were resistant to local pathogen populations. The line with gene Sr22 was resistant to local pathogen populations and to the Ug99 race group in Kenya. The genes Sr26, Sr36 and Sr39 were not detected in the analyzed wheat lines.

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