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>Chardonnay_62_1223

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>Kabernet_Cortis_47_1223

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>Sibirkovii_9_1223

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>Sypun_Chernii_170_1223

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CGGGTAGG

Fig. 2. Sequencing of 850 bp amplicons of *VvMybA1* alleles in colored (Cabernet Cortis and Sypun black) **and uncolored** (Chardonnay and Sibirkovy) **varieties of European wine grape (*Vitis vinifera* L.)** (Anapa zonal ampelographic collection SKANCSVV) (for the figure, see http://www.agrobiology.ru).

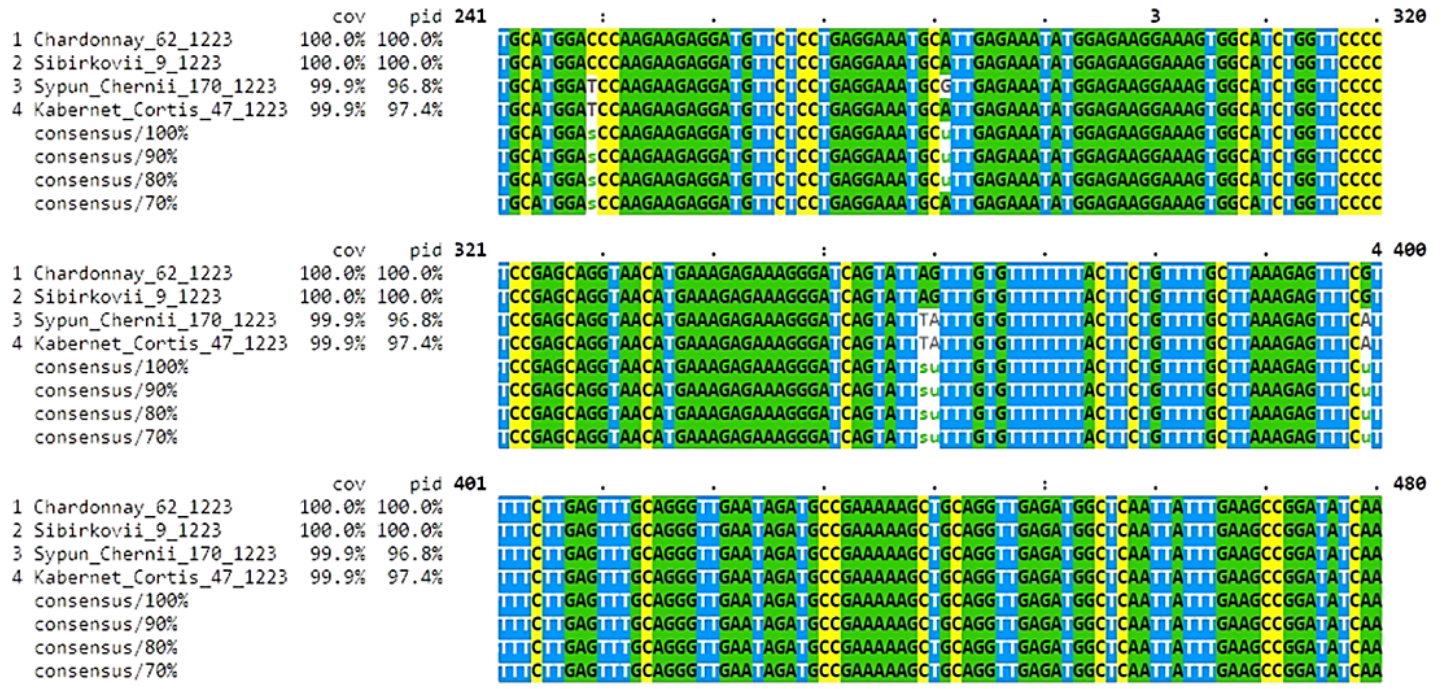

Fig. 3. Analysis of *VvMybA1* gene allele regions in four native and introduced varieties of European wine grape (*Vitis vinifera* L.) (colored Cabernet Cortis and Sypun black and uncolored Chardonnay and Sibirkovy) (ClustalO program; Anapa zonal ampelographic collection SKANCSVV). Nucleotide polymorphisms identified by alignment of the analyzed sequences in four genotypes are highlighted as in white. A unique nucleotide substitution at nucleotide position 281 in Sipun black variety is marked by white background. For the figure, see http://www.agrobiology.ru.

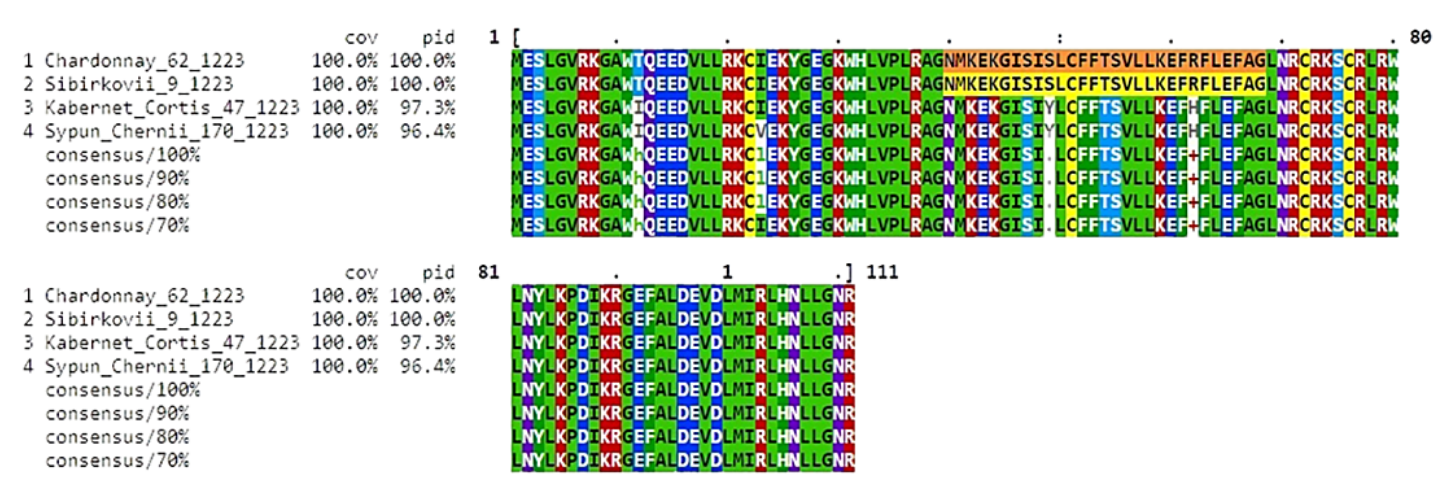

Fig. 5. Alignment of amino acid sequences (with an intron insertion marked) **translated from *VvMybA1* gene alleles in four native and introduced varieties of European wine grape (*Vitis vinifera* L.)** (colored Cabernet Cortis and Sypun black and uncolored Chardonnay and Sibirkovy) (ClustalO program; Anapa zonal ampelographic collection SKANCSVV). In Sypun black variety, in the position 23 (highlighted in white background) there is an amino acid substitution (isoleucine characteristic of the other three varieties is replaced to valine). For the figure, see http://www.agrobiology.ru.