

Table S2 Functional annotation of predicted target genes

Target gene		Interval	length	E-value	Target gene annotation
<i>VIT_206s0004g02800</i>	<i>REVOLUTA</i>	8460-8621	161	6.51E-13	homeobox leucine-zipper protein
<i>VIT_213s0019g04320</i>	<i>REVOLUTA-like</i>	2456-2617	161	6.04E-13	homeobox-leucine zipper protein revoluta
<i>VIT_204s0008g03250</i>	<i>HB-15</i>	8733-8897	164	1.61E-12	class iii hd-zip protein 8 homeobox-leucine zipper protein athb-15 hypothetical protein
<i>VIT_210s0003g04670</i>	<i>HOX32</i>	380-541	161	1.56E-13	homeobox-leucine zipper protein hox32-like

The Grapevine Genome CRIBI Biotech website (http://genomes.cribi.unipd.it/grape/search_form.php) was used to identify the target gene name, grape ID, coordinates and their functions.