

表 1 砂梨木质素合成相关转录因子基因 *MYB* 启动子区特征序列位置预测

Table 1 Predicted position of the characteristic sequence in the promoter regions of sand pear putative transcription factor gene *MYB* related to lignin biosynthesis

预测项 Predicted item	启动子(对应基因) Promoter (Corresponding gene)			
	<i>PpMYBx1</i>	<i>PpMYBx2</i>	<i>PpMYBx3</i>	<i>PpMYBx4</i>
TATA box 位置 TATA box position	578 bp	664 bp	753 bp, 288 bp	709 bp
启动子位置 Promoter position	612 bp	677 bp	787 bp, 322 bp	741 bp
增强子位置 Enhancer position	484 bp	541 bp	527 bp	594 bp
克隆长度 Clone length	821 bp	874 bp	972 bp	914 bp

