Supplementary Fig. S3 Relative expression of *LhMYB6* in the basal and upper tepal parts at flower bud developmental stage (St) 1, 2, 3, and 4. Levels of gene expression were analyzed by qRT-PCR, and *LhACTIN* was used to normalize the expression of the genes. Vertical bars represent the standard error (SE) of the means of three flower buds. The same letters above the columns indicate that the values are not statistically significant (p<0.05) by Tukey HSD. Note that raised spots were not included in the tepal parts.