

**Figure S1.** Comparison of the DNA methylation map of the full-length T-DNA sequence among the transgenic gentian plant lines #3, #15, and #19.

The percentage of methylated cytosine is represented by bar charts (red, CpG; green, CpWpG; black, CpHpH and CpCpG), and the position of each cytosine is represented below. Each promoter, coding, terminator, and T-DNA border region is indicated by a different color. Crossbars at the bottom within the 35S promoter region indicate the region of the probes for EMSA.

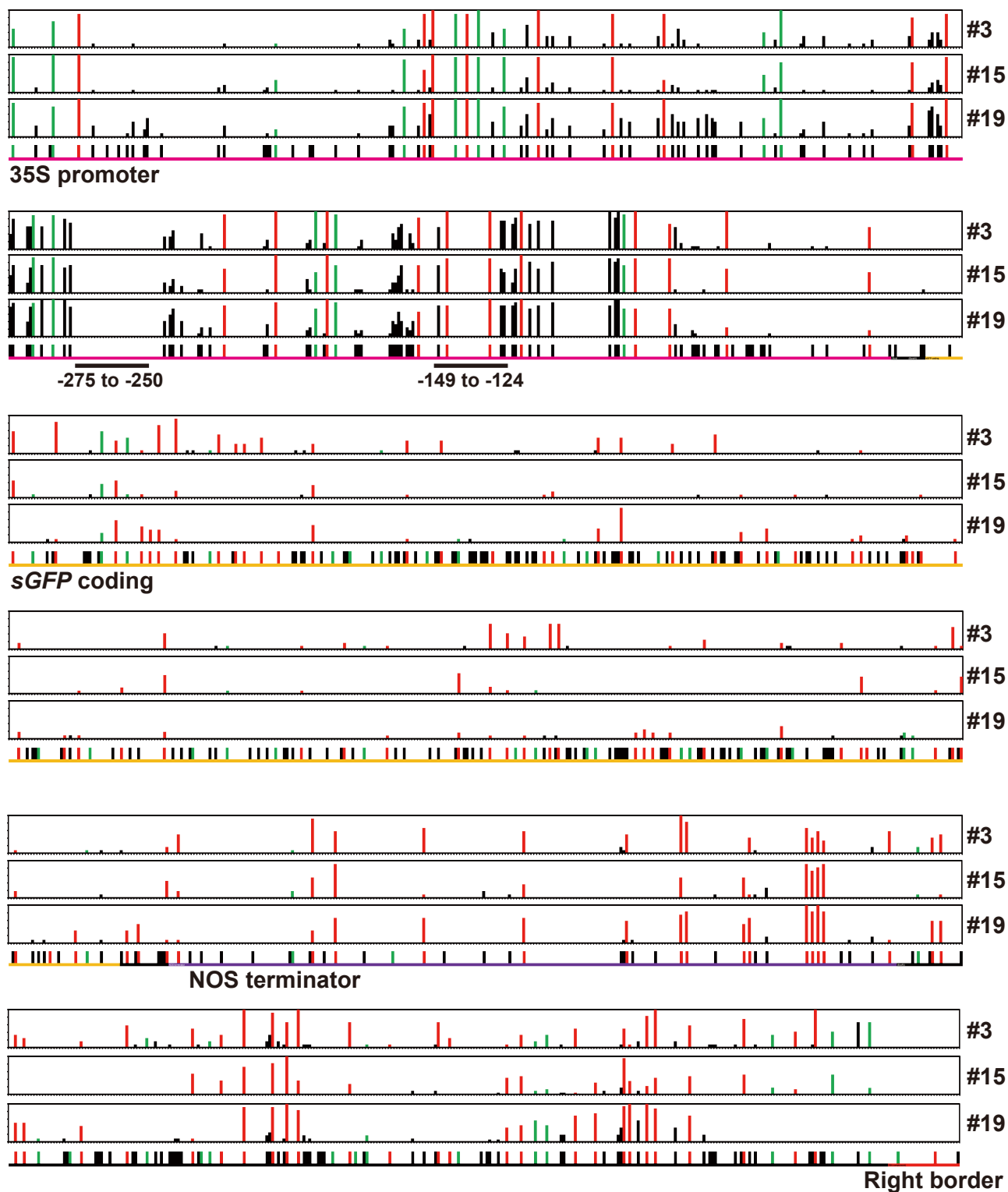


Figure S1. (Continued.)