

Supporting Information

Mukherjee et al. 10.1073/pnas.0811357106

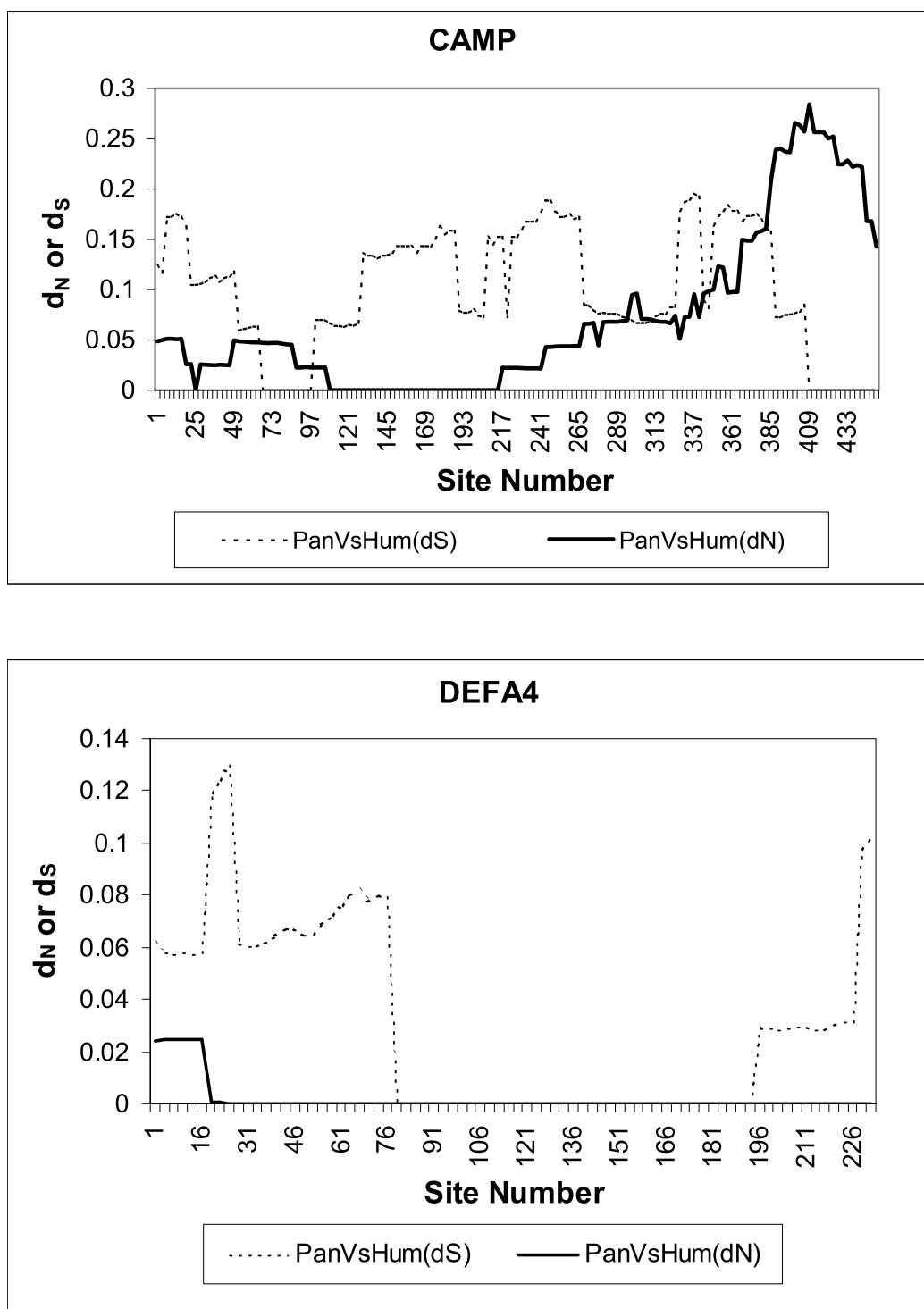


Fig. S1. Plots of the numbers of synonymous (d_S) and non-synonymous (d_N) substitutions per site for 11 innate immunity genes, calculated by comparing data on humans (generated in this study) with chimpanzee (*Pan*) data (downloaded from public database), using a sliding window approach, with window width = 60 bases and slide length = 3 bases. These plots could not be prepared for *TLR5* because of unavailability of chimpanzee data.

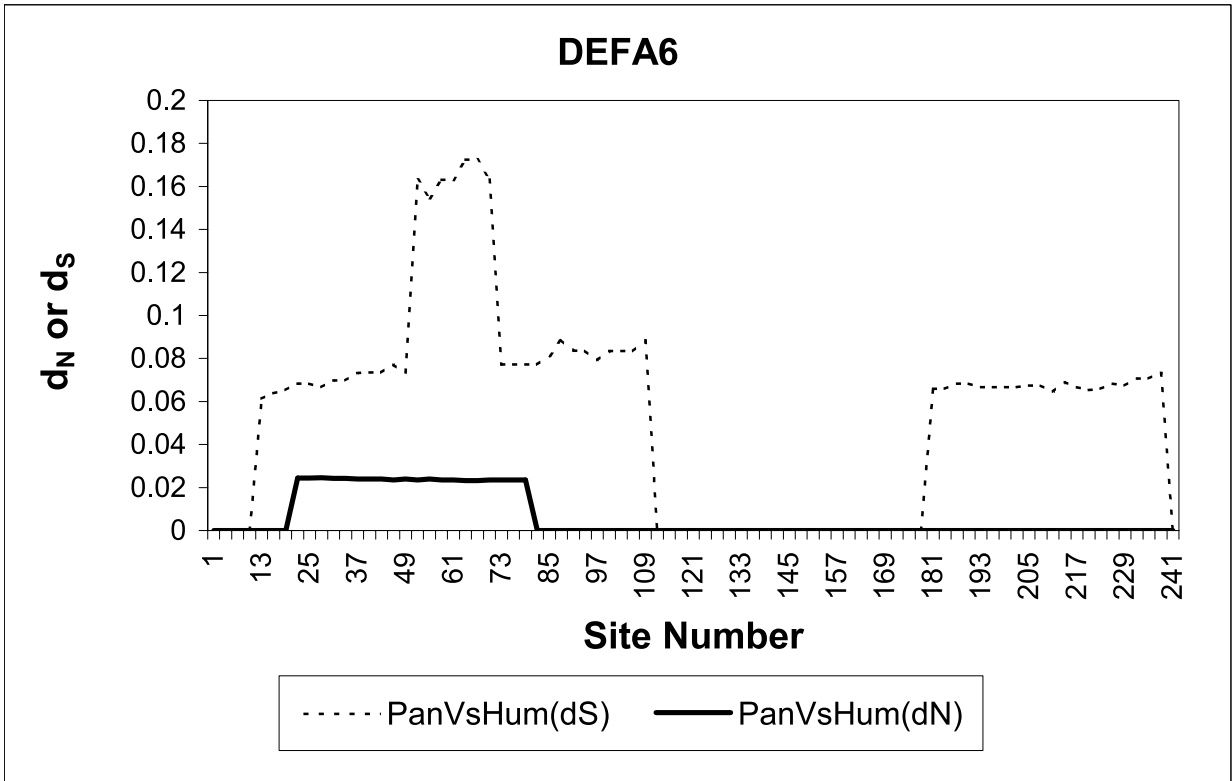
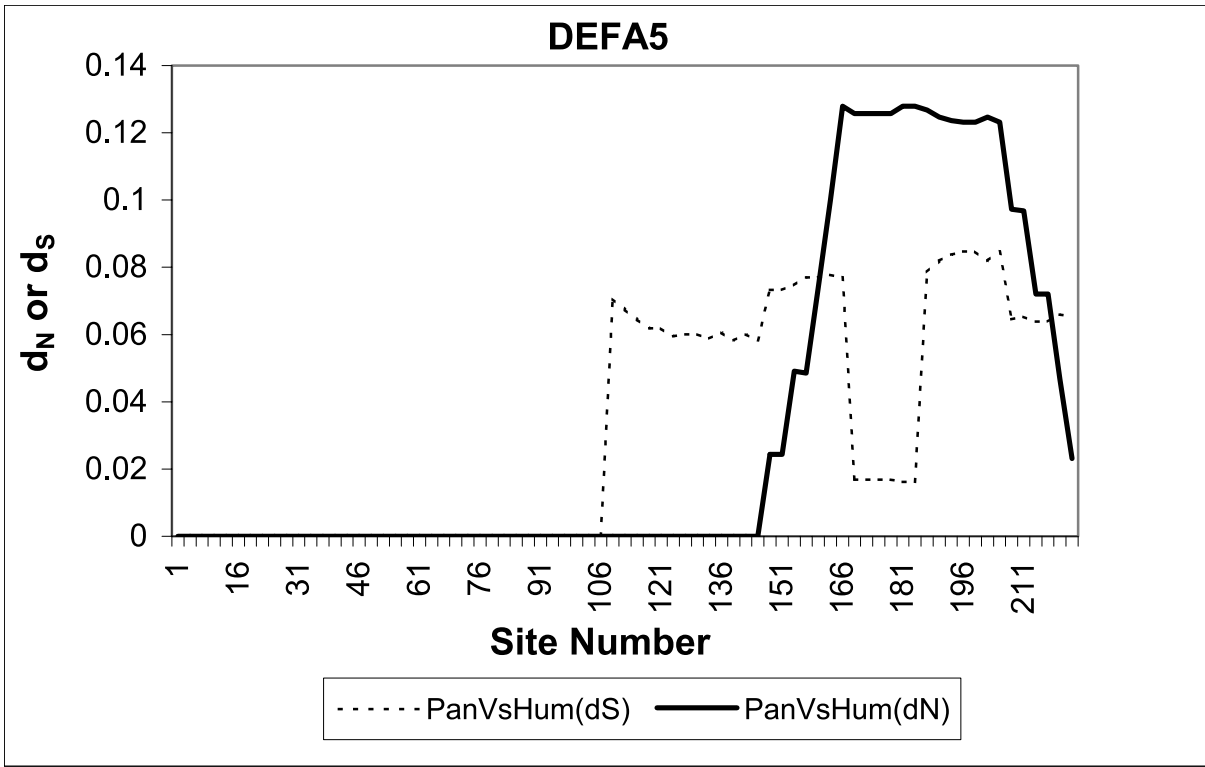


Fig. S1 (continued).

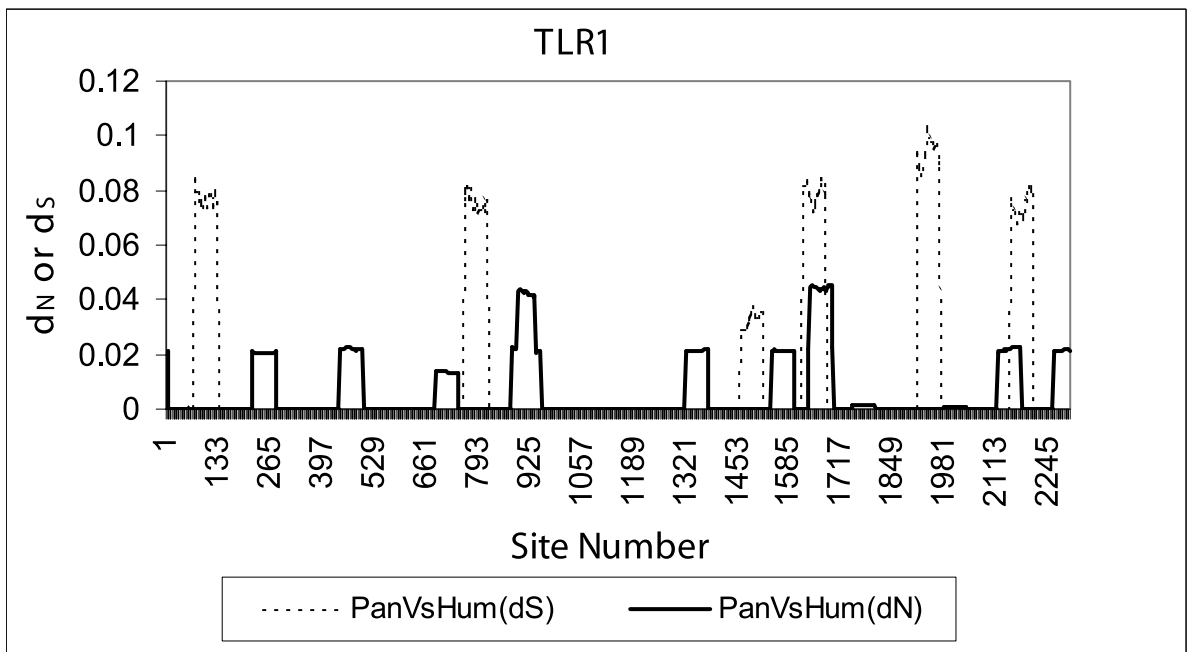
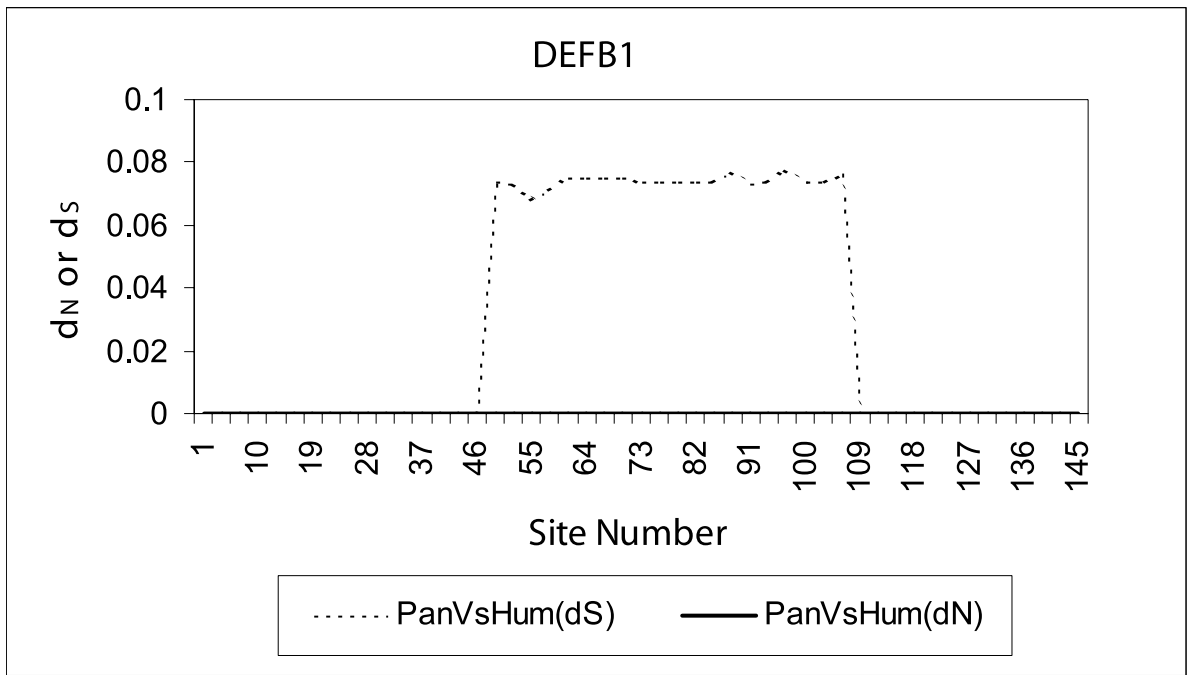


Fig. S1 (continued).

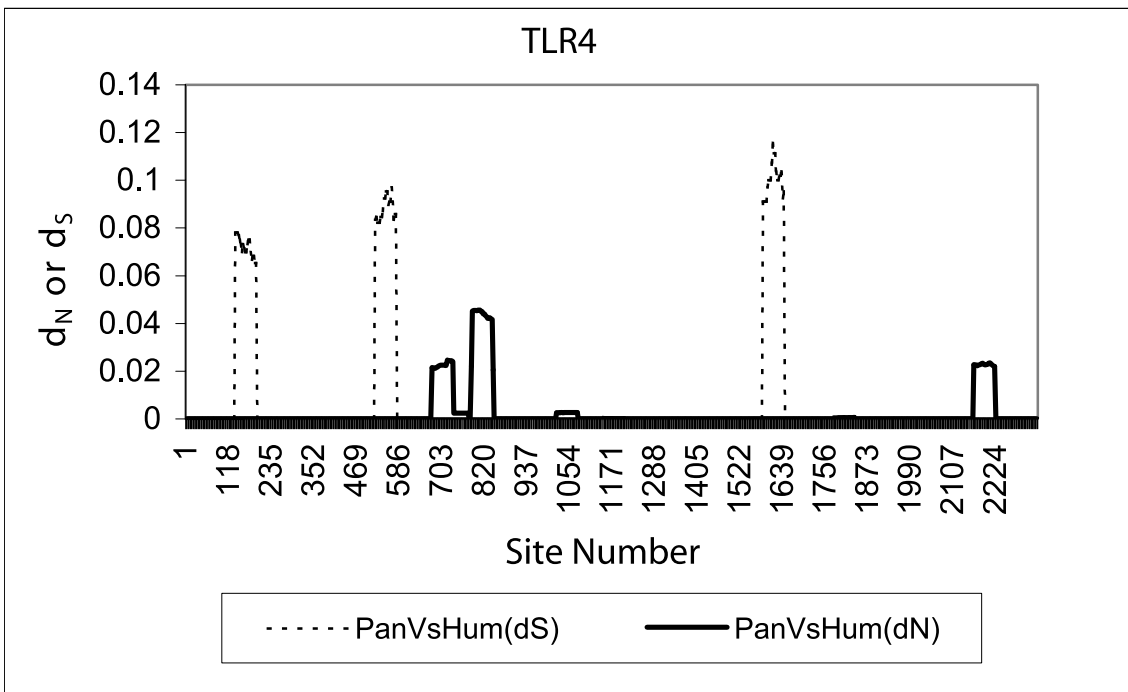
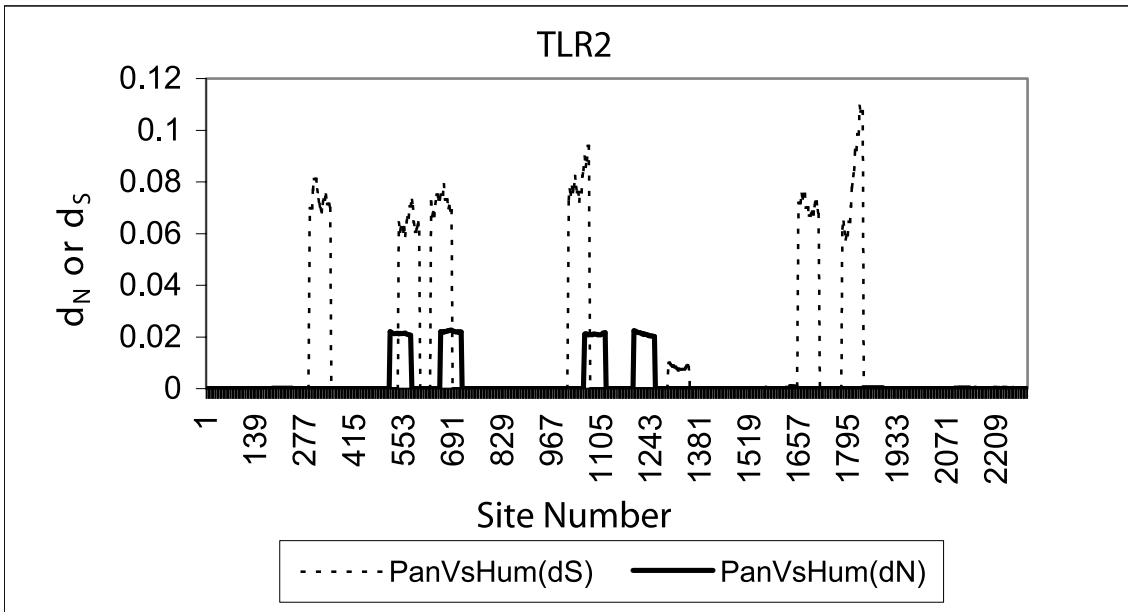


Fig. S1 (continued).

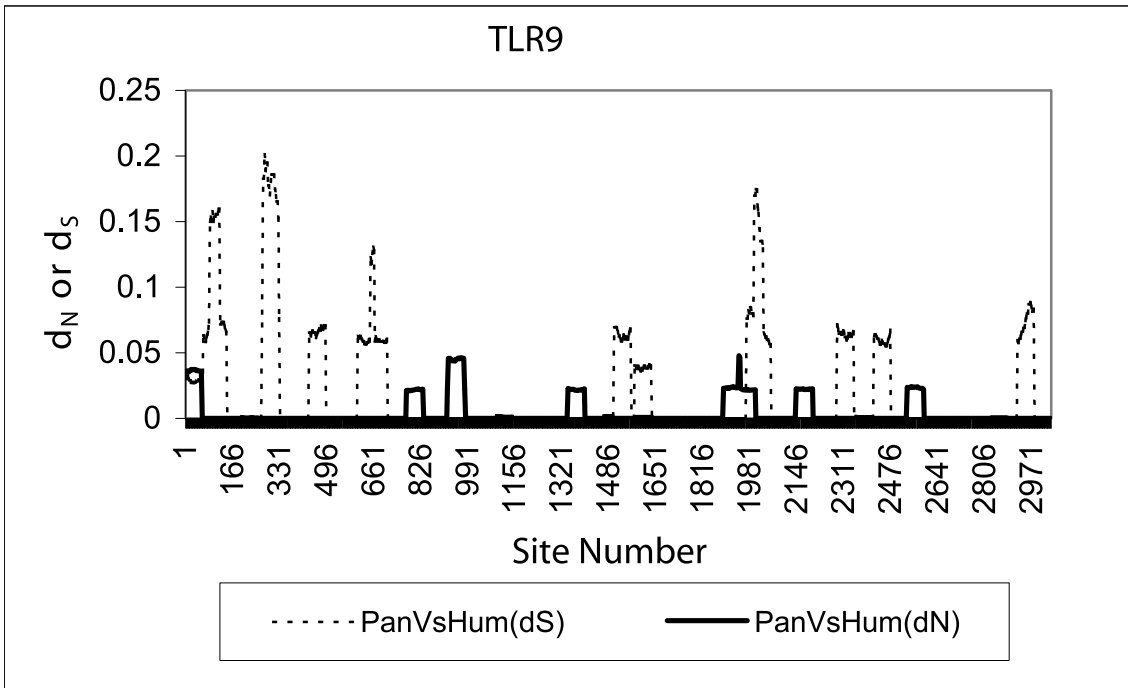
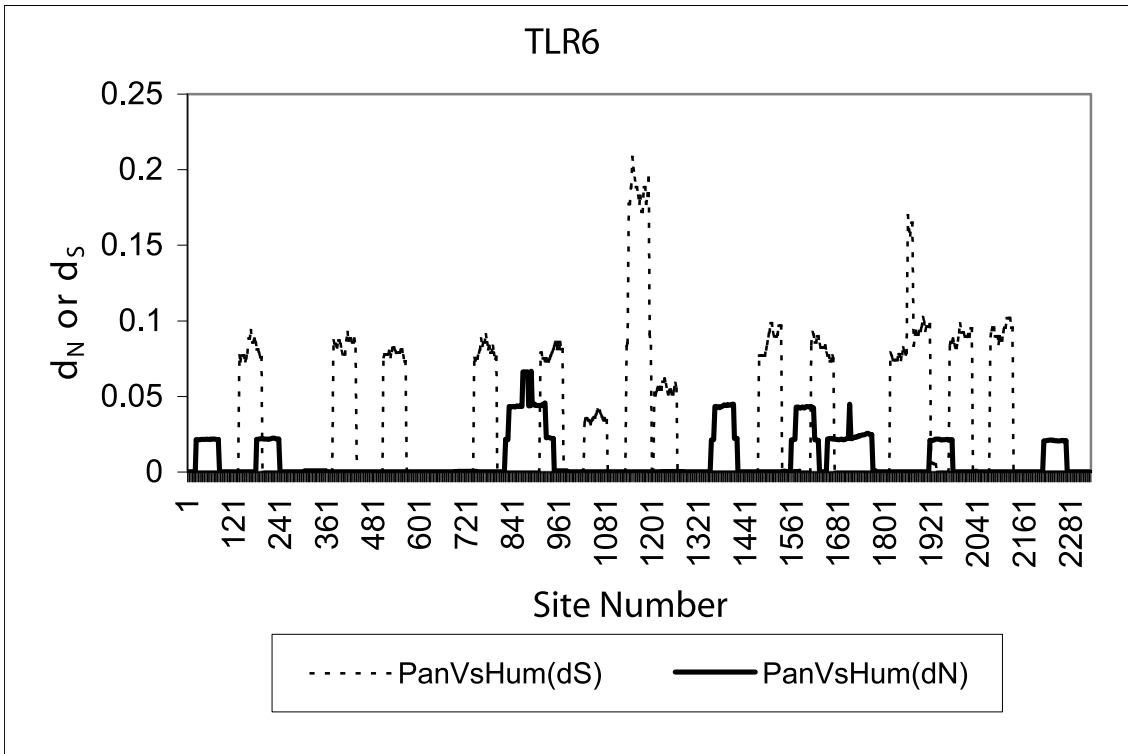


Fig. S1 (continued).

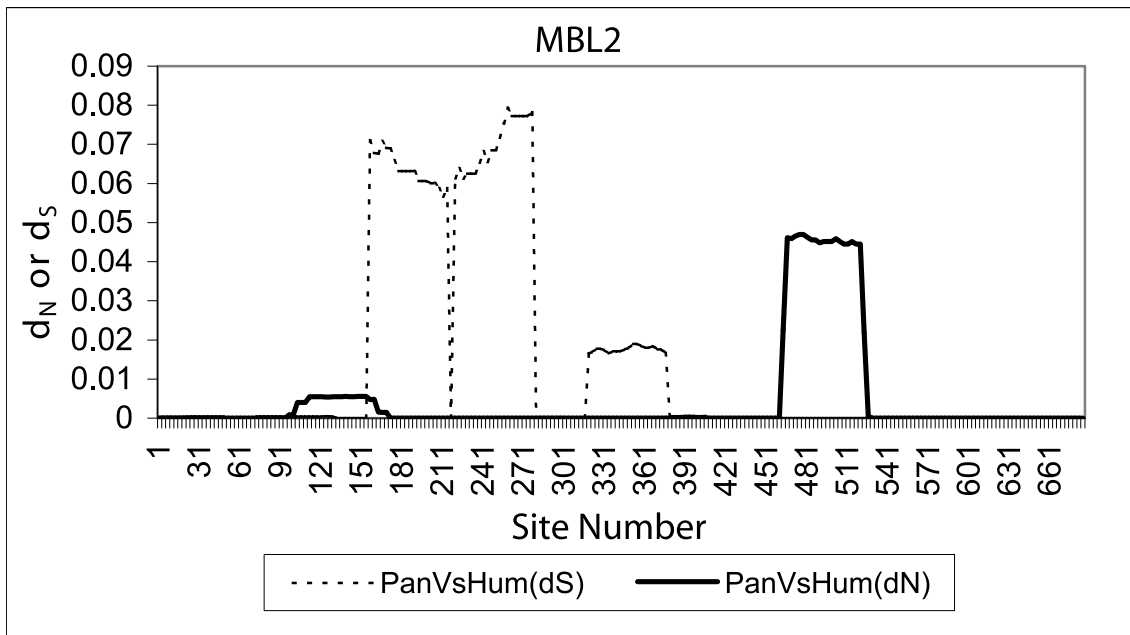


Fig. S1 (continued).

Other Supporting Information Files

[Table S1](#)