

1 **Supporting Information for "Ensembler: Enabling high-throughput molecular simulations at the**  
2 **superfamily scale"**

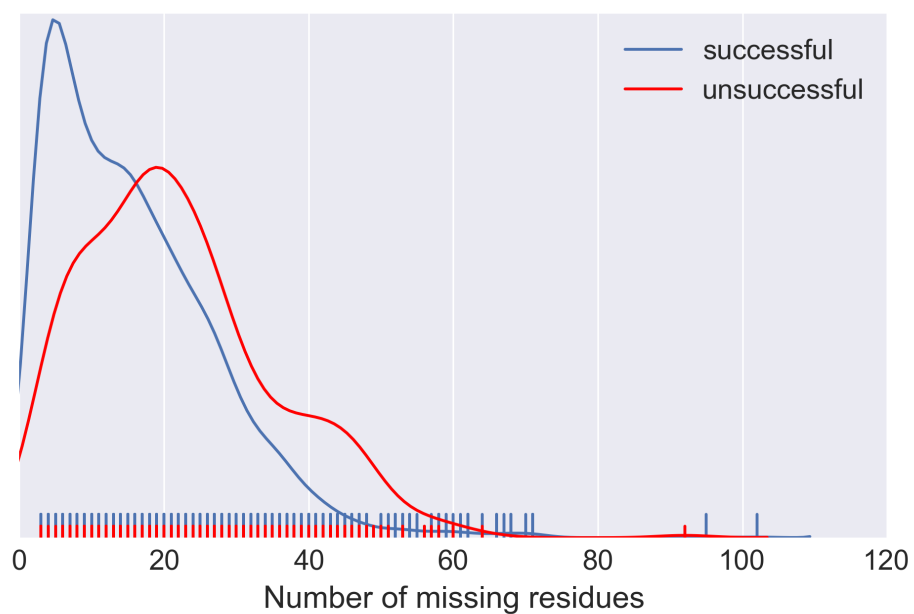
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**FIG. 1. Distributions of the number of missing residues for templates for which remodeling (with the `loopmodel` command) was either successful or unsuccessful.** The plotted distributions are smoothed using kernel density estimation, and the raw data points are shown as a rug plot.