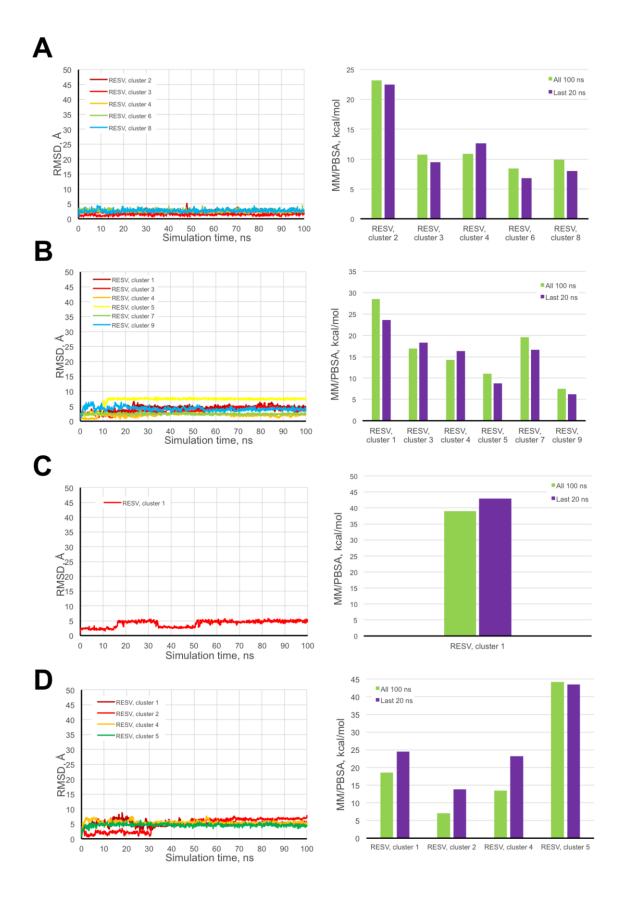


Supplementary Figure 1. Trajectories and MM/PBSA free energy analysis of RSV (left) and acarbose (right) forming complexes with the yeast form of GGA.



Supplementary Figure 2. Trajectories and MM/PBSA free energy analysis of RSV clusters forming complexes with human GAA (**A**), yeast GGA (**B**), human α -Man I (**C**), and human α -Man II homology model (**D**).