

Figure S4a. Binding mode of IAA with TIR1. Hydrogen bonds are drawn as green lines.

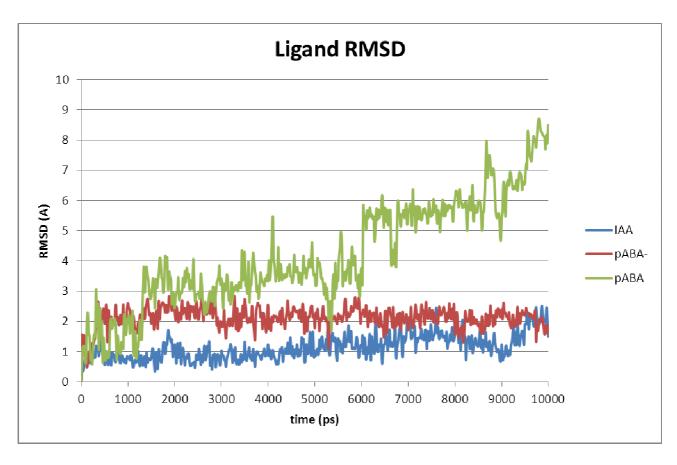


Figure S4b. Root Mean Square Deviation of the ligand with respect to the MD starting.