



Figure S2: Ligand properties of β -selinene complexed with odorant binding protein 7 across a 100 ns MD simulation (a) Ligand RMSD (b) Radius of Gyration (rGyr) (c) Intramolecular Hydrogen Bonds (intraHB) (d) Molecular Surface Area (MolSA) (e) Solvent Accessible Surface Area (SASA) (f) Polar Surface Area (PSA)