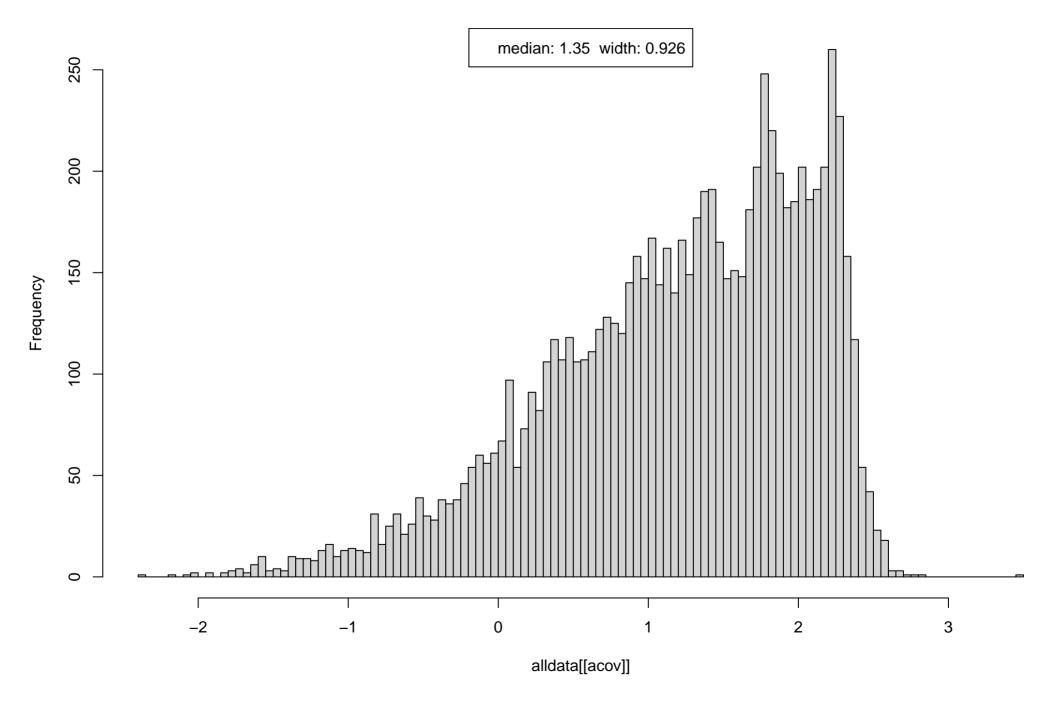
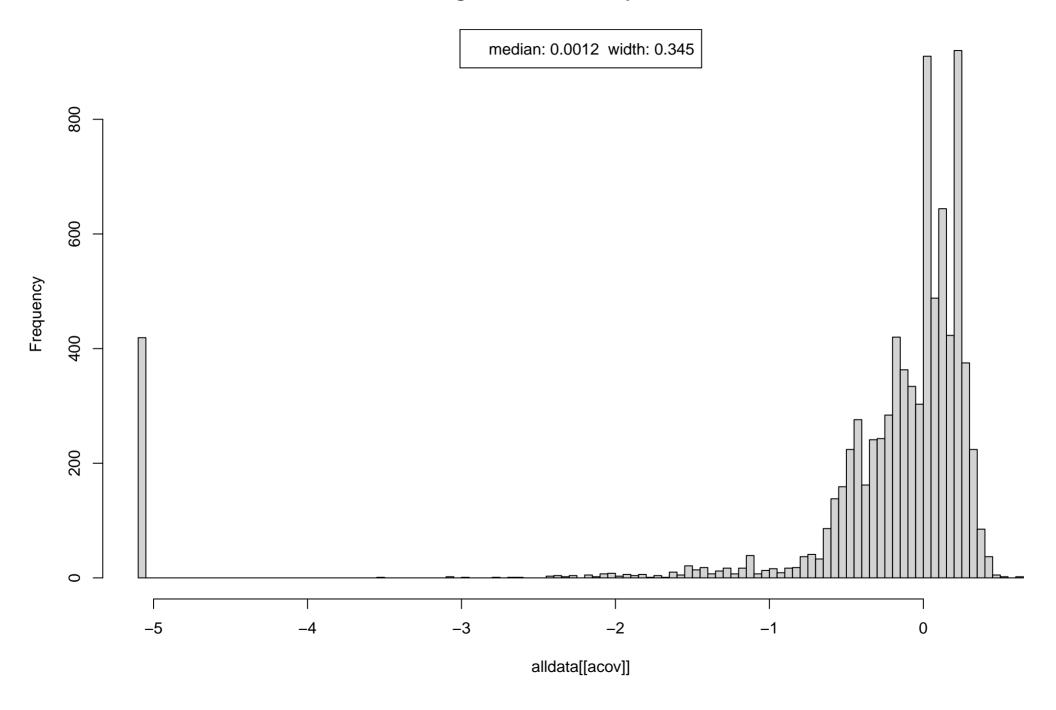
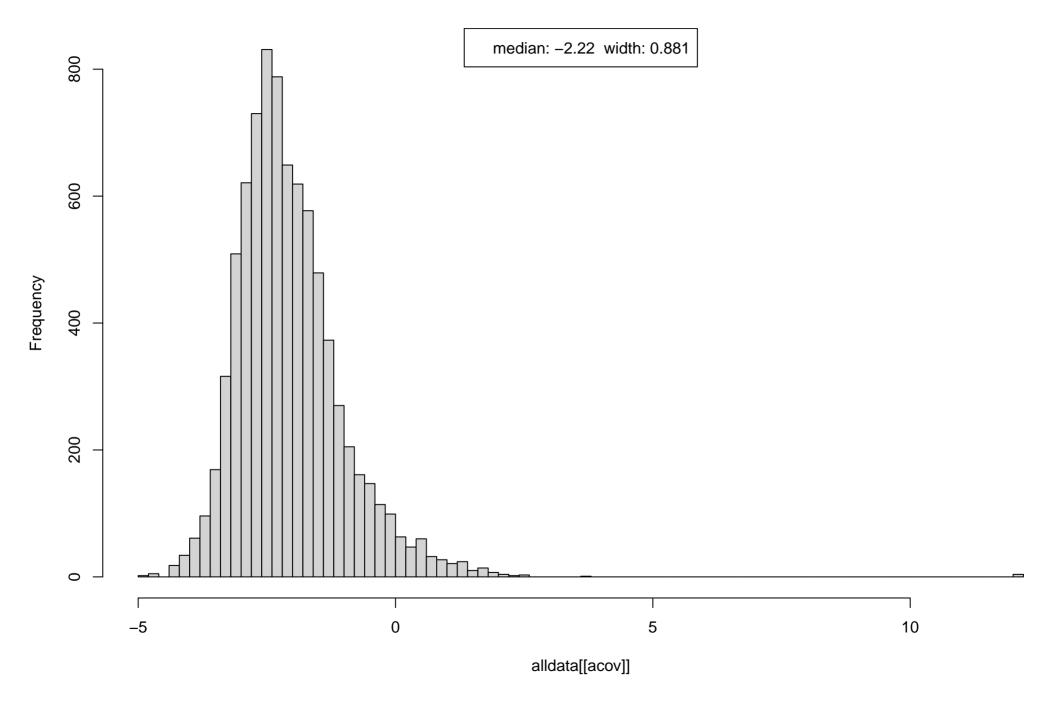
log_RMSD



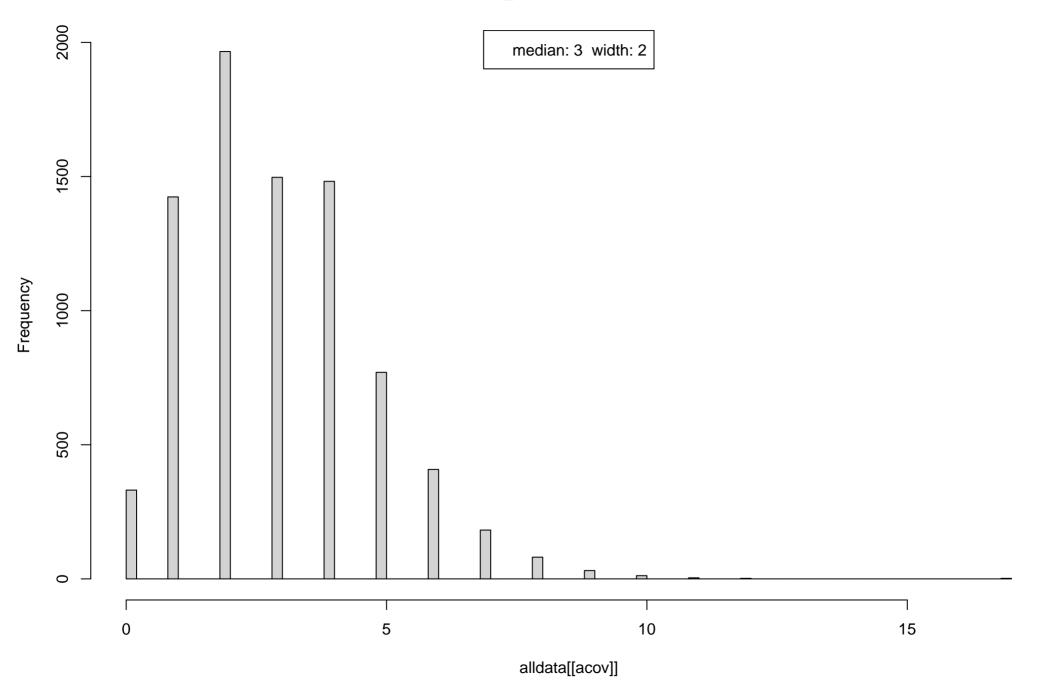
log_mcs_unbonded_polar_sasa



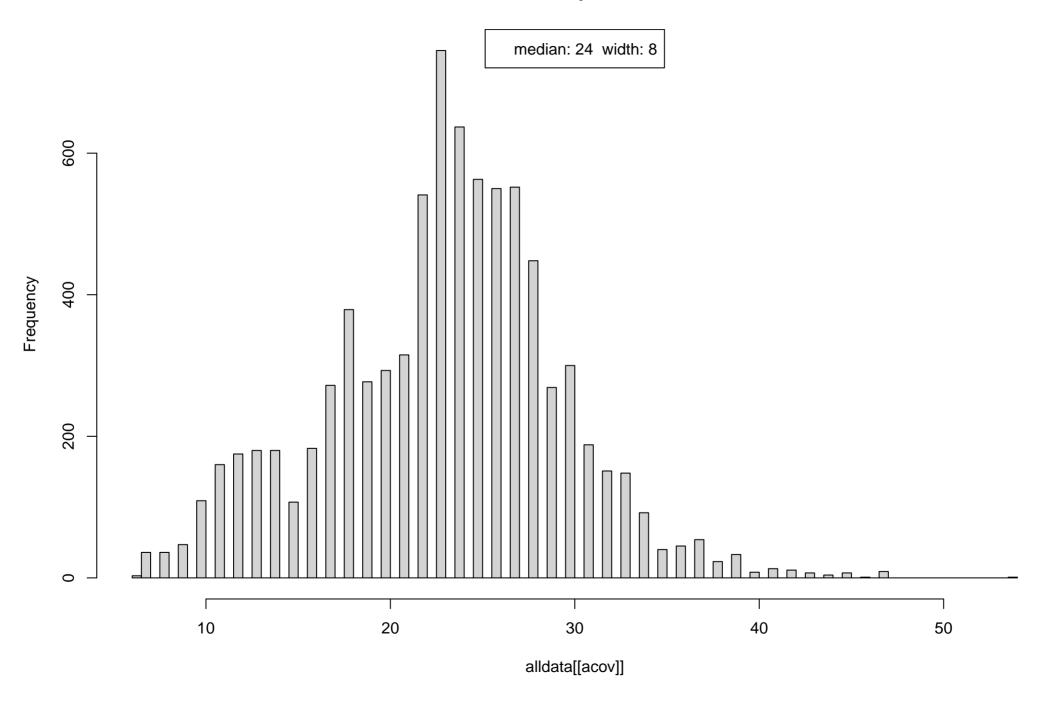
logit_ec_tanimoto_similarity



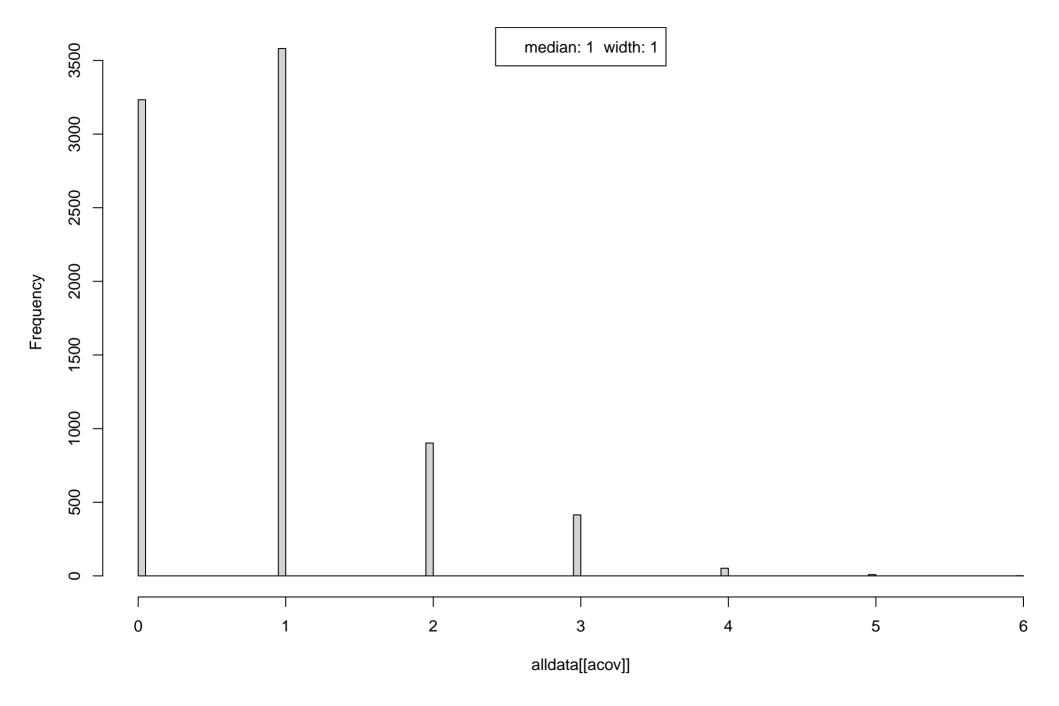
$mcs_NumHeteroAtoms$



${\bf docked_HeavyAtomCount}$







docked_NumRotatableBonds

