



Figure S3. Time evolution of dihedral angles in L-BH (ligand code 5FH) and L-IMH from MD simulations. (A-C) Simulations 5FH(g^-) MD_001–MD_003 (L-BH bound to Mhp1, started from the extended conformation as seen for the IMH-bound structure). (D) L-BH in solution. (E-H) Simulations 5FH(g^+) MD_001–MD_004 (L-BH bound to Mhp1, started from the U-shaped folded over conformation as modeled in 2JLO). (I-K) Simulations IMH(g^-) MD_001–MD_003 (L-IMH bound to Mhp1 as seen in the crystal structure). (L) L-IMH in solution. (J) Extended to 500 ns. (L) L-NMH in solution.