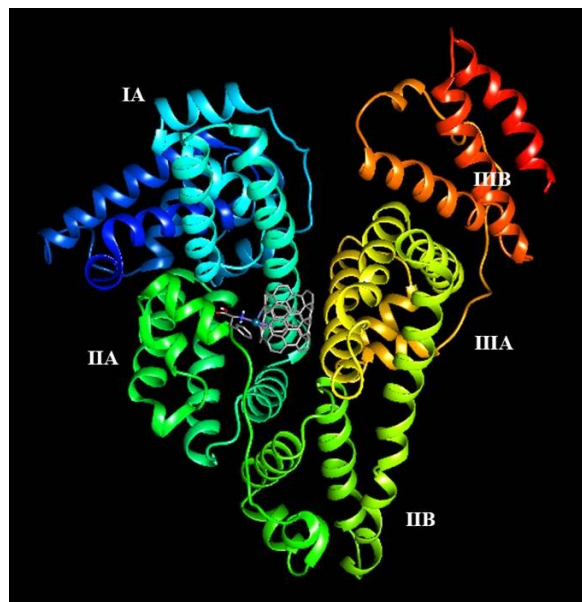
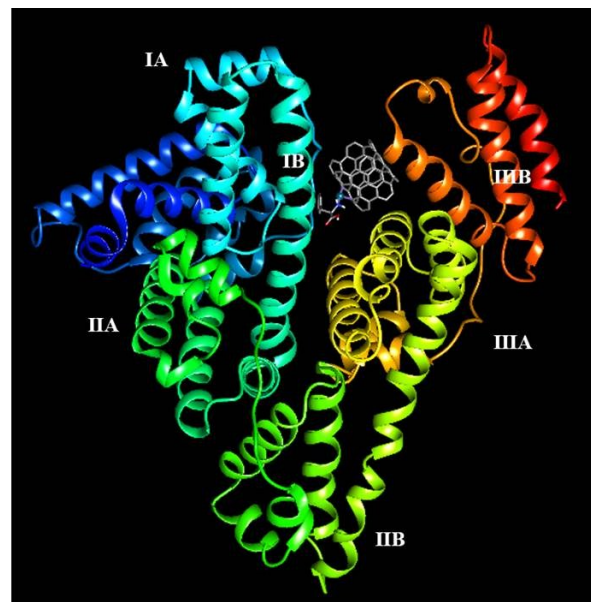


Table 1. The binding free energies of the purposed binding to HAS. The binding free energies (kcal.mol⁻¹) and Inhibition Constant of the biosensor-HAS complex were estimated by the scoring function of AUTODOCK.

Number	Binding site	$\Delta G_{\text{binding}}$	ΔG_{nes}	ΔG_{es}	ΔG_{tor}	K_i
1	IIA	-10.05	-11.70	-0.27	1.92	42.68
2	IB	-10.05	-11.71	-0.27	1.92	42.81



1



2

Table 2. Hydrogen bonding parameters

Hydrogen bonding interaction	Bond length (Å)	Bond angles (°)
Phe-Pd/SWCNT ... Glu292	2.76	127
Phe-Pd/SWCNT ... Lys190	2.45	145
Phe-Pd/SWCNT ... Glu425	2.47	136