

Supplementary Information

Interaction Rule and Mechanism of Perfluoroalkyl Sulfonates contained different carbon chain with Human Serum Albumin

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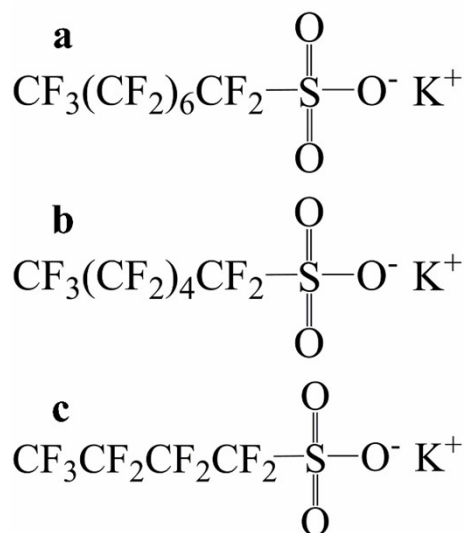
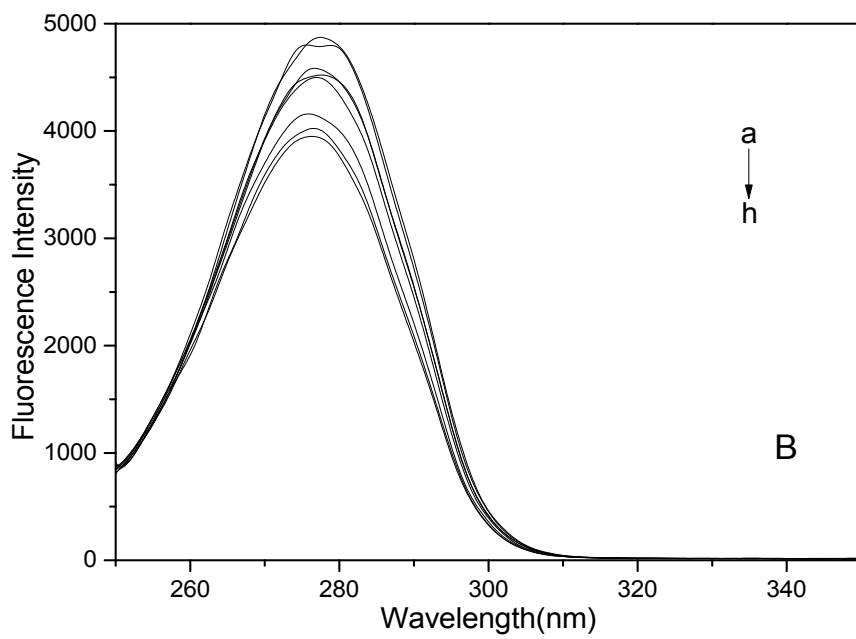
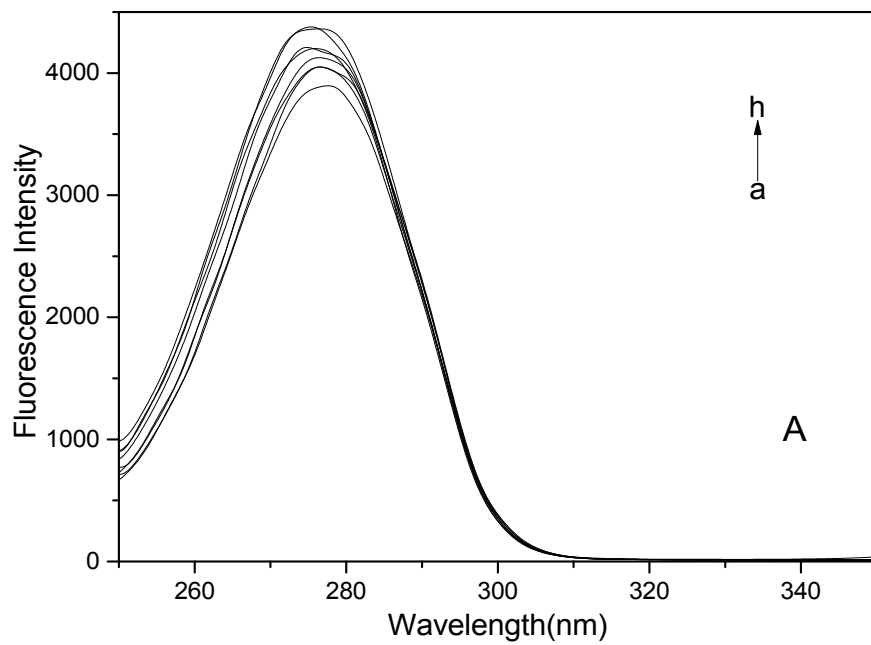


Fig. S1 Chemical structure of perfluorooctane sulfonate, perfluorohexane sulfonate and perfluorobutane sulfonate: a (potassium salt of PFOS), b (potassium salt of PFHS), c (potassium salt of PFBS).



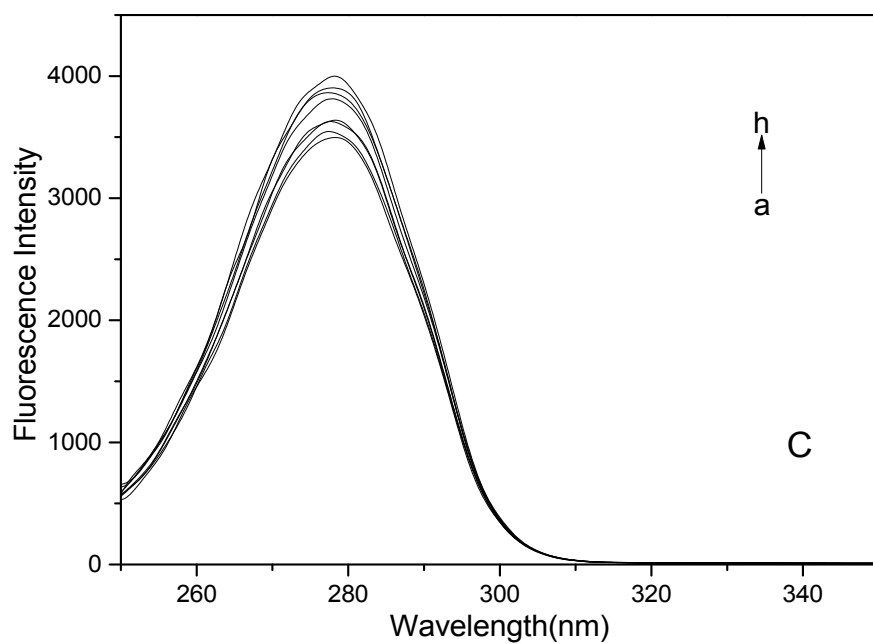
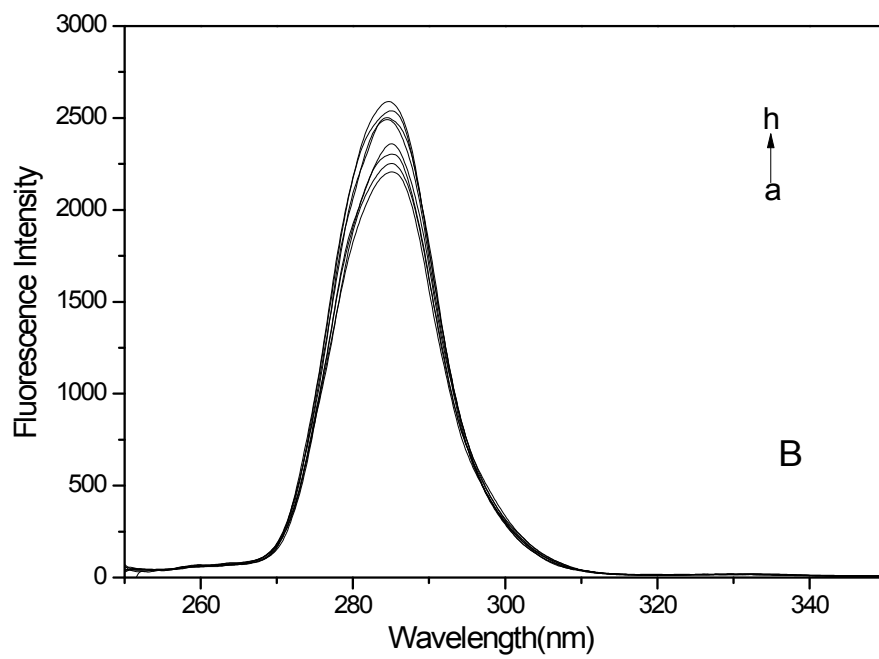
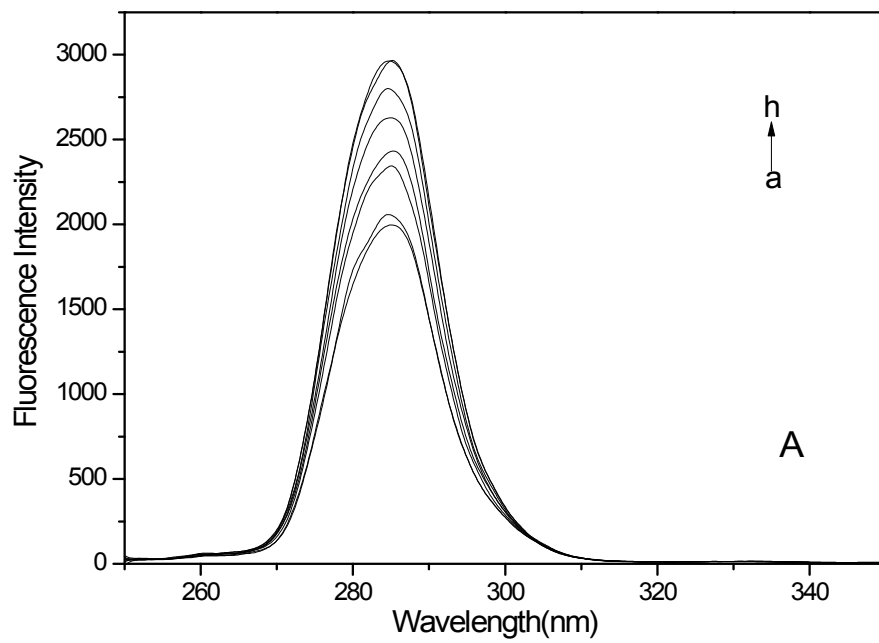


Fig. S2 Synchronous fluorescence spectra of HSA with different concentrations of PFASs. Conditions: HSA: 1×10^{-6} mol/L; A (PFOS), B (PFHS) and C (PFBS) (a-h): $0, 2 \times 10^{-6}, 6 \times 10^{-6}, 1 \times 10^{-5}, 3 \times 10^{-5}, 5 \times 10^{-5}, 7 \times 10^{-5}, 1 \times 10^{-4}$ mol/L; pH=7.4; T=298 K; $\Delta\lambda=60$ nm.



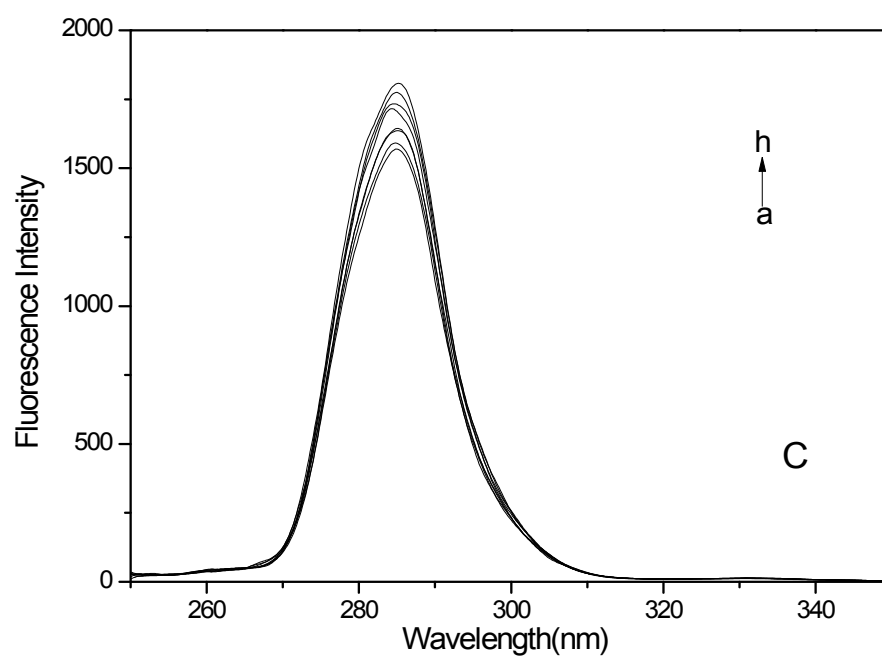
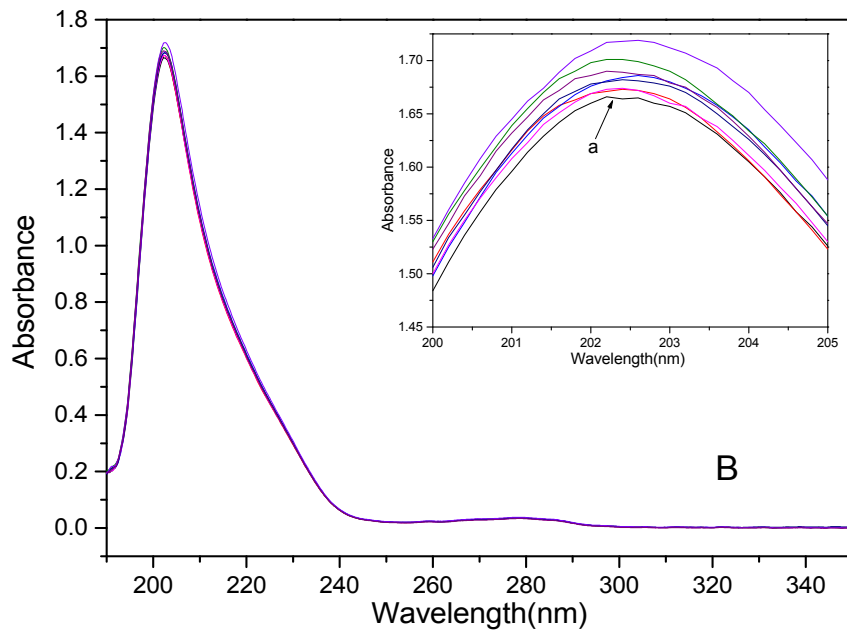
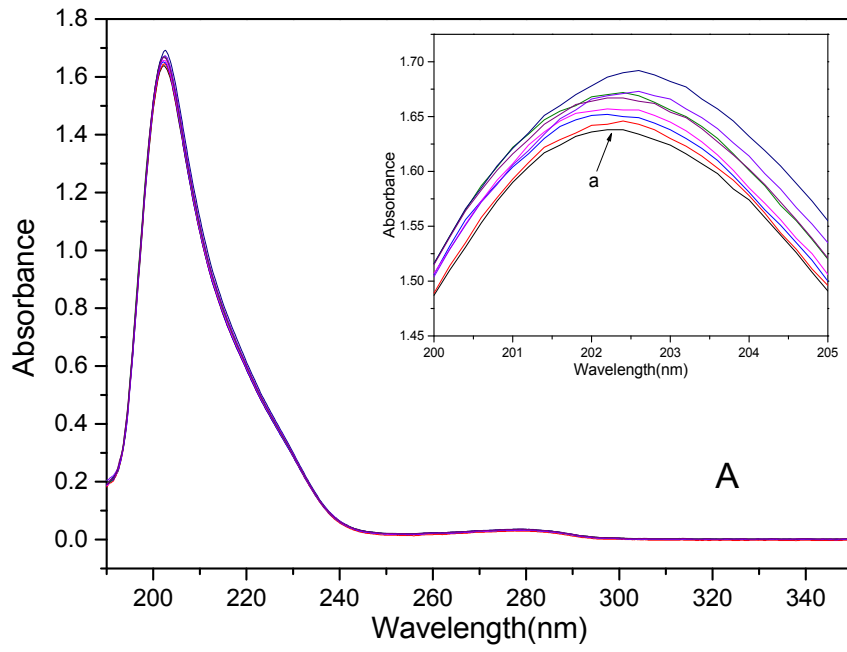


Fig. S3 Synchronous fluorescence spectra of HSA with different concentrations of PFASs. Conditions: HSA: 1×10^{-6} mol/L; A (PFOS), B (PFHS) and C (PFBS) (a-h): 0, 2×10^{-6} , 6×10^{-6} , 1×10^{-5} , 3×10^{-5} , 5×10^{-5} , 7×10^{-5} , 1×10^{-4} mol/L; pH=7.4; T=298 K; $\Delta\lambda=15$ nm.



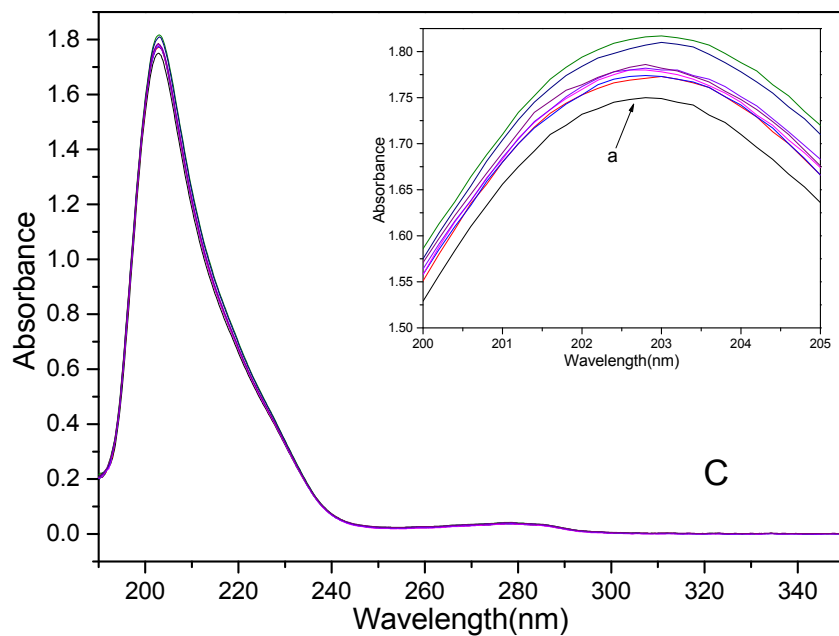
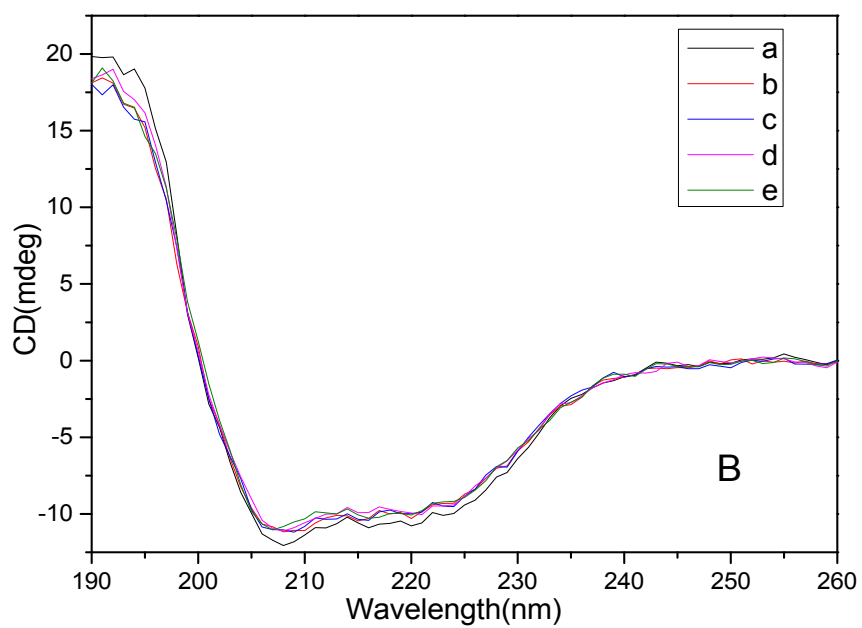
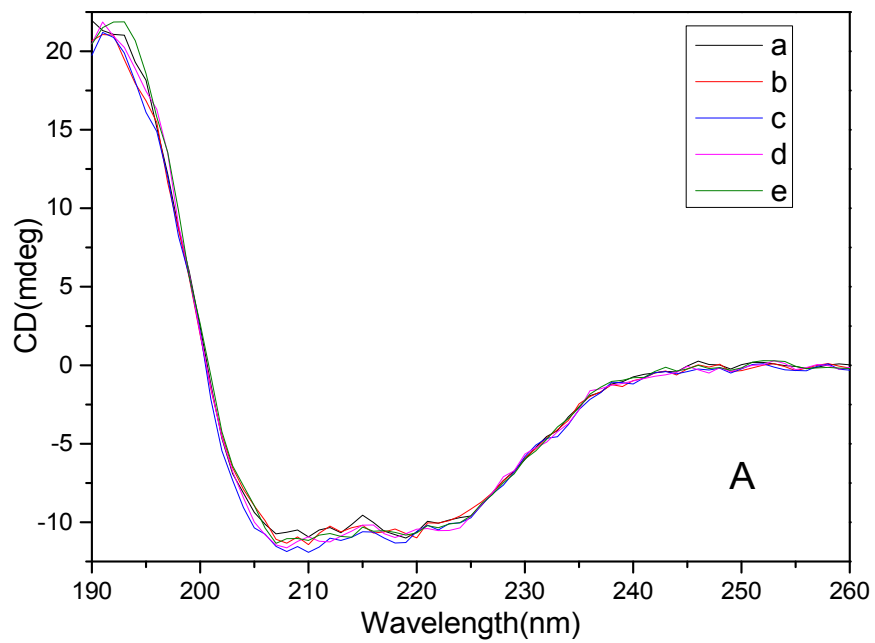


Fig. S4 UV-visible absorption spectra of HSA with different concentrations of PFASs. Conditions: HSA: 1×10^{-6} mol/L; A (PFOS), B (PFHS) and C (PFBS): 0, 2×10^{-6} , 6×10^{-6} , 1×10^{-5} , 3×10^{-5} , 5×10^{-5} , 7×10^{-5} , 1×10^{-4} mol/L; pH=7.4; T=298 K;



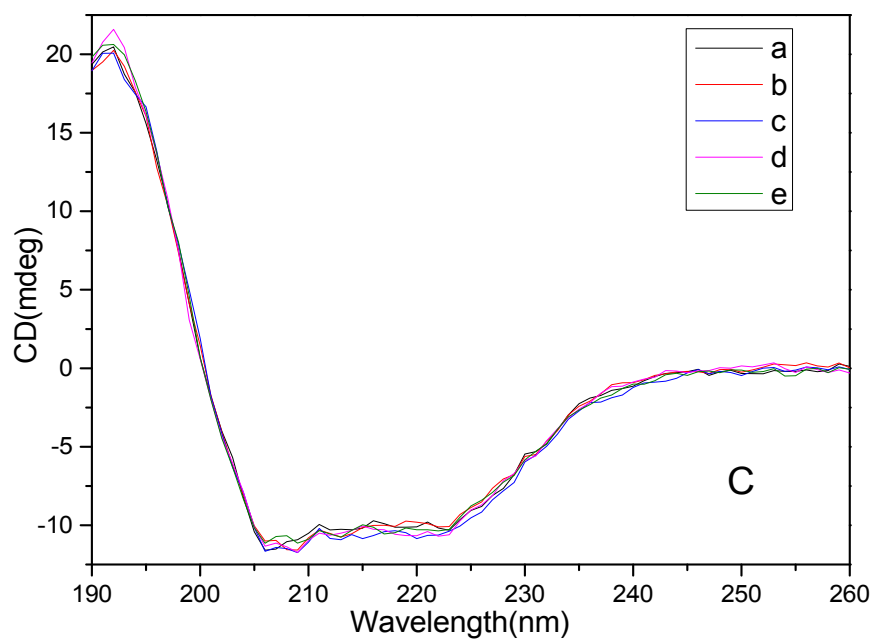


Fig. S5 CD Spectra of HSA with different concentrations of PFASs. Conditions: HSA: 1×10^{-6} mol/L; A (PFOS), B (PFHS) and C (PFBS) (a-e) : 0, 5×10^{-6} , 1×10^{-5} , 5×10^{-5} , 1×10^{-4} mol/L; pH=7.4; T=298K.

Table S1 The interaction data between PFOS and HSA

Ligand	Receptor	Residue	Type	Distance
H 8881	OD 5973	Asn 391	H-don	1.39
O 8880	NH 6284	Arg 410	H-acc	2.65
O 8878	OH 6306	Tyr 411	H-acc	2.48

Table S2 The interaction data between PFHS and HSA

Ligand	Receptor	Residue	Type	Distance
H 8880	OD 5973	Asn 391	H-don	1.28
O 8879	NH 6284	Arg 410	H-acc	2.50
O 8877	OH 6306	Tyr 411	H-acc	2.53

Table S3 The interaction data between PFBS and HSA

Ligand	Receptor	Residue	Type	Distance
H 8887	OD 5973	Asn 391	H-don	1.32
O 8886	NH 6284	Arg 410	H-acc	2.60
O 8877	OH 6306	Tyr 411	H-acc	2.49