

Deciphering the interaction of plumbagin with human serum albumin: A combined biophysical and molecular docking study

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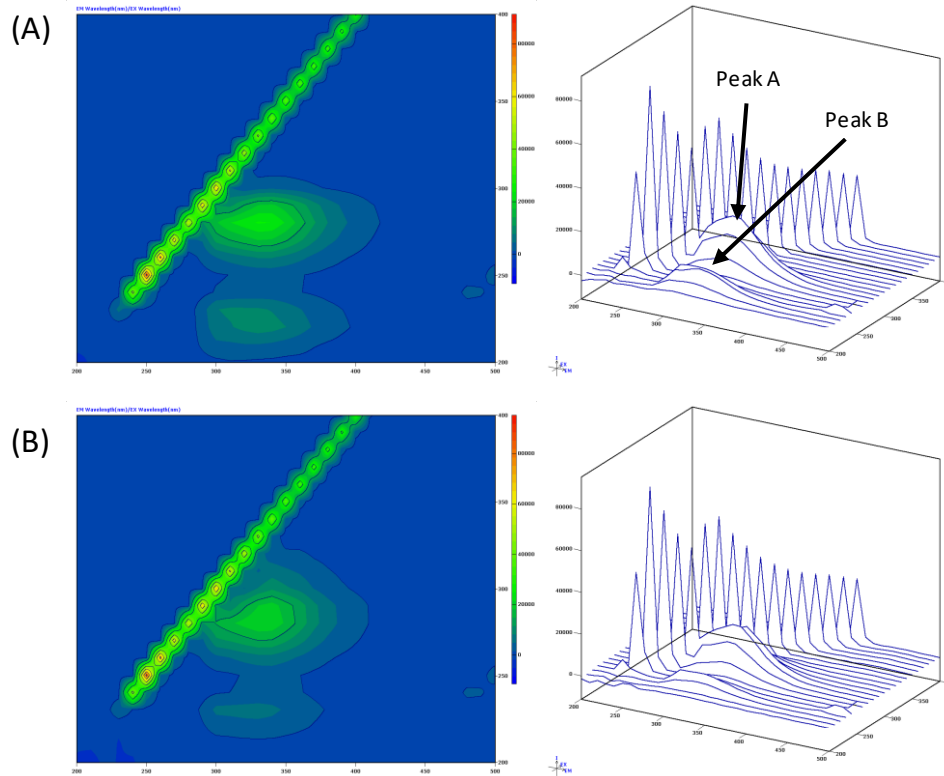
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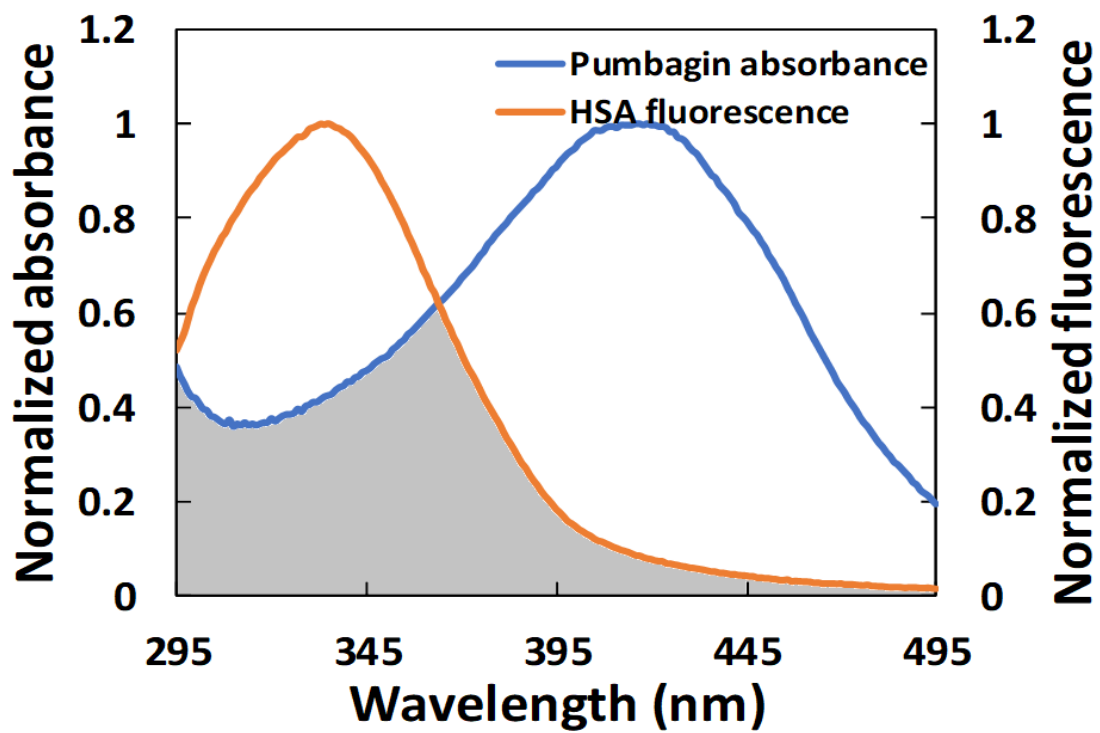
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Supplementary Figures



Supplementary Fig S1. Three-dimensional fluorescence spectra of (A) HSA (B) HSA-plumbagin complex.



Supplementary Fig S2. Spectral overlap of the fluorescence spectrum of HSA and absorption spectrum of plumbagin.

Supplementary Tables

Supplementary Table S1. α -helical content of HSA in the absence and presence of plumbagin.

Molar ratio ([HSA]/[plumbagin])	MRE _{208nm}	% α -helix
1:0	-20818.19	57.99
1:1	-18608.64	50.37
1:2	-17716.28	47.29

Supplementary Table S2. Förster resonance energy transfer data obtained from the spectral overlap of HSA and plumbagin.

Parameters	Values
Energy transfer efficiency (E)	29.04%
r	2.27 nm
Critical distance (R ₀)	1.96 nm
Spectral overlap (J)	2.64×10 ⁻¹⁵ cm ³ L/mol

Supplementary Table S3. Changes in accessible surface area (Δ ASA) in Å² for the HSA residues interacting with plumbagin.

S. No.	Residues	ASA (Å ²) in HSA	ASA (Å ²) in HSA–plumbagin complex	Δ ASA (Å ²)	Domain
1.	Lys199	28.861	10.555	18.306	IIA
2.	Arg222	31.178	19.248	11.93	IIA
3.	Leu238	30.529	0.454	30.075	IIA
4.	Arg257	15.368	5.231	10.137	IIA
5.	Leu260	15.548	4.653	10.895	IIA
6.	Ile290	12.978	0	12.978	IIA
7.	Ala291	40.991	7.57	33.421	IIA