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Supplementary Information

Cold atmospheric plasma driven self-assembly in serum proteins: Insights into the protein aggregation to biomaterials

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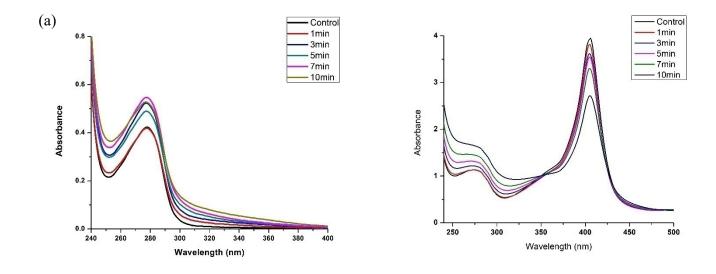


Figure S1 Ultra-Violet Visible Spectroscopy of control and plasma treated (a) BSA and (b) Hb

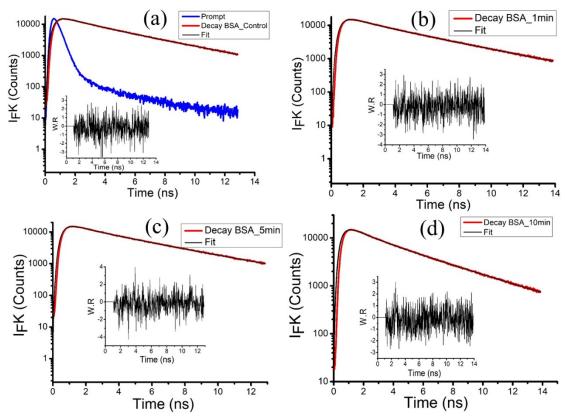


Fig S2: Time resolved spectra of BSA and CAP treated BSA

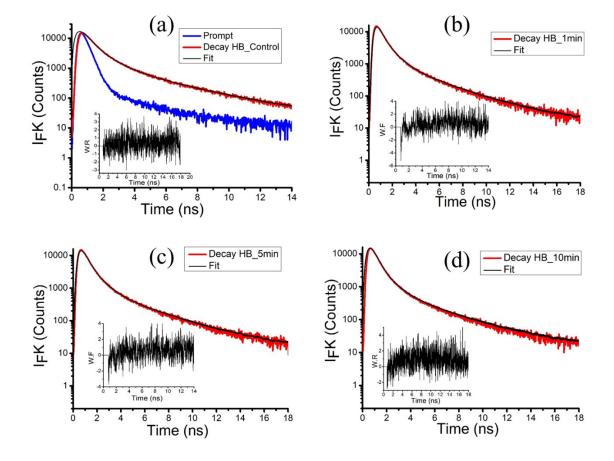


Fig S3: Time resolved spectra of Hb and CAP treated Hb

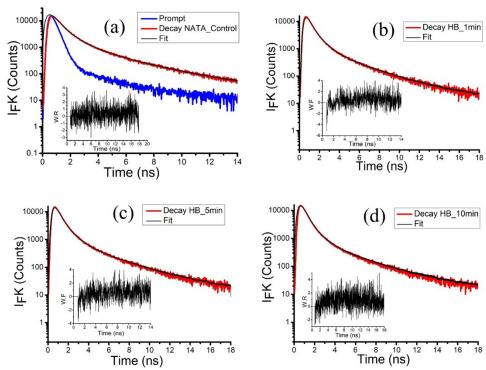


Fig S4: Time resolved spectra of NATA and CAP treated NATA

Table S1 - Lifetime of BSA at different time of plasma treatment

Treatment	$\tau_1(ns)$	% population	$\tau_2(ns)$	% population
time (Min)				
0	2.170 ± 0.009	27.30	6.217 ± 0.004	72.70
1	2.117 ± 0.008	29.74	6.280 ± 0.004	70.26
3	2.085 ± 0.008	29.39	6.327 ± 0.004	70.61
5	2.058 ± 0.008	29.59	6.258 ± 0.004	70.41
7	2.092 ± 0.008	31.34	6.250 ± 0.004	68.66
10	2.048 ± 0.007	32.54	6.064 ± 0.003	67.46

Table S2 - Lifetime of Hb at different time of plasma treatment

Treatment	τ_1 (ns)	%	τ_2 (ns)	%	τ_3 (ns)	%
Time		population		population		population
(Min)						
0	0.418 ± 0.001	73.97	1.263 ± 0.002	22.87	4.418 ± 0.009	3.17
1	0.212 ± 0.000	86.50	1.201 ± 0.002	11.17	4.598 ± 0.010	1.45
3	0.216 ± 0.001	84.50	1.184 ± 0.002	14.30	4.658 ± 0.011	1.60
5	0.260 ± 0.001	84.65	1.207 ± 0.002	13.84	4.653±0.012	1.53
7	0.268 ± 0.001	83.35	1.163 ± 0.002	15.80	4.586±0.012	1.55
10	0.280 ± 0.001	85.95	1.198 ± 0.002	12.26	4.672±0.012	1.35

Table S3 - Lifetime of NATA at different time of plasma treatment

Treatment Time (min)	$\tau_1(ns)$	% population	
0	3.153±0.002	100	
1	3.125±0.002	100	
5	3.070±0.002	100	
10	3.022±0.002	100	

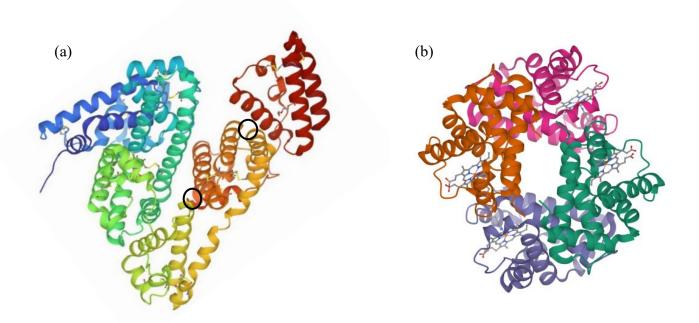


Figure S5 - Structure of (a) BSA with the Trp located in the position 213 and 134 marked in black circles. (Ref: PDB 4F5S) (b) Hemoglobin with the porphyrins and the six tryptophan residues (α 14, β 15 and β 37 in each dimer) Ref: 1HDA.