## Supporting Information

## Study of the Interaction of Folic Acid-modified Gold Nanorods and Fibrinogen through Microfluidics: Implications on Protein Adsorption, Incorporation and Viability of Cancer Cells

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Figure S-1: FTIR spectra of functionalization of GNR-CTAB with SH-PEG-COOH and SH-PEG-OMe.



Figure S-2. Microchips in their different conformations. Y-shape, B) S-shape.



Figure S-3: Hydrodynamic diameters of GNR-CTAB.





Figure S-4: Hydrodynamic diameters of GNR-PEG.



Figure S-5: Hydrodynamic diameters of GNR-PEG-FA.



Figure S-6: Hydrodynamic diameters of GNR-FB-10'.







Figure S-8: Hydrodynamic diameters of GNR-FB-60'.

			Size (d.nm):	% Intensity:	St Dev (d.nm):
Z-Average (d.nm):	8,459	Peak 1:	3,406	51,0	1,090
Pdl:	0,430	Peak 2:	56,97	49,0	25,36
Intercept:	0,727	Peak 3:	0,000	0,0	0,000
Result quality :	Refer to quality report				



Figure S-9: Hydrodynamic diameters of GNR-FB  $Q_1Y$ 







Figure S-11: Hydrodynamic diameters of GNR-FB  $Q_{a}Y$ 



Figure S-12: Hydrodynamic diameters of GNR-FB  $Q_1S$ 



Figure S-13: Statistical analysis of FB molecules per GNR-FA obtained by TEM images analysis. \*\*\*\*p < 0.0001, \*\*\*p < 0.0003 related to the comparison of bulk 60',  $Q_3$ Y and  $Q_3$ S.



Figure S-14: Scheme of the movements (elasticity) of the fibrinogen based on the fluctuations size of the slowest vibration-modes of the protein. (The colored arrows indicate the fibrinogen movements directions).



Figure S-15. Matrix cross-correlation maps of flexibility of chain N of FB showing the correlations between residue fluctuations which are plotted like a function of i vs j residue together with the corresponding flexibility perturbation in the 3D crystallographic chain (red positive correlation, blue negative correlation).



Figure S-16: Circular dicroism spectra of A) Fibrinogen, inset SELCON3 and CONTIN analysis. B) bulk experiments at 10, 30 and 60 min, C) Y-shape experiments, D) S-shape experiments.