Supplementary Data



Figure S1. Scanning electron microscopy image of the nanofilm after the nanoprecipitation of 20 mg/mL BSA solution.



Figure S2. Stability of proteins in NP.

- A. UV-visible spectroscopy data of BSA solution (1mg/mL) in water (green line), BSA NP diluted in 30 times (black line), the corresponding supernatant after BSA NP centrifugation after nanoprecipitation (red line) and the supernatant after the storage of the same BSA NP at +4 ° C for 3 months (blue line).
- B. Results of SDS-PAGE of original BSA (lane 1), BSA NP immediately after nanoprecipitation (lane 2) and BSA NP after their storage at +4 ° C for 10 months (lane 3).







Figure S4. Fluorescent microscopy image of BSA-RhoB nanoparticles. Scale bar is 10 μ m.



Figure S5. Competitive inhibition of binding of BSA-RhoB nanoparticles with immobilized polyclonal antibodies against BSA in the presence of unlabelled BSA proteins at different concentrations.



Figure S6. Atomic force microscopy image (part A) and normalized distribution of heights (part B, mean size = 16 nm, median size = 13 nm, standard deviation = 9 nm) of the lysozyme NP.



Figure S7. Enzymatic activity of lysozyme (black line) and lysozyme NP (red line) according to ²¹.

| | BSA NP | Fibrinogen NP | Polylysine NP | lg NP | Lysozyme NP |
|------------------------|--------|------------------|------------------|-------|----------------|
| Mean size, nm | 129 | 116 | 116 | 175 | 33 |
| Median size, nm | 114 | 94 | 108 | 158 | 31 |
| Standard deviation, nm | 38 | 40 | 29 | 36 | 6 |

 $Table \ S1. \ {\rm Dynamic \ light \ scattering \ of \ nanoparticles \ fabricated \ from \ different \ proteins.}$

| | BSA in water | BSA NP | Fibrinogen in water | Fibrinogen NP | Lysozyme in water | Lysozyme NP | |
|----------------|-----------------|--------|------------------------|------------------|----------------------|----------------|--|
| Helix,% | 57,6 | 63,8 | 25,5 | 24,1 | 32,9 | 27,7 | |
| Antiparallel,% | 3,9 | 3,2 | 11,2 | 12,1 | 8,3 | 9,3 | |
| Parallel,% | 4,7 | 4,0 | 11,2 | 11,6 | 8,8 | 10,9 | |
| Beta-Turn,% | 13,1 | 12,3 | 18,3 | 18,9 | 16,9 | 17,7 | |
| Random Coil,% | 20,1 | 17,4 | 38,6 | 38,7 | 33,4 | 37,6 | |

Table S2. Deconvolution analysis of circular dichroism spectra of proteins in water solutions and in nanoparticles

Table S3. Enzyme-linked immunosorbent assay data of three independent experiments ofdetection of antibodies against HBsAg and the corresponding nanoparticles from Mab HV301against HBsAg

| Antibodies | ELISA reverse titers | | | Average geometric | 80 | | | stical cance |
|--|----------------------|-----------------|-----------------|----------------------|----|-------------|--------|-------------------|
| | Experiment 1 | Experiment 2 | Experiment 3 | titer (GEOMEAN) | 50 | IG(GEOMEAN) | ig(SD) | Statis signifi |
| Mab HV301 against HBsAg | 160 | 160 | 320 | 200 | 90 | 2.3 | 0.2 | P>0.05 |
| NP from Mab HV301 HBsAg against | 80 | 80 | 160 | 100 | 50 | 2.0 | 0.2 | P>0.05 |