

Supplementary material

Low-fouling SPR detection of lysozyme and aggregates

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Table S1. Comparison of the analytical characteristics of lysozyme sensors

| Detection mode ^a | Detection limit (nM) | Reference |
|---|------------------------|-----------|
| SPR | 20 | This work |
| SPR-previous work | 70 | [29] |
| SPR, graphene coated interface | 0.5 | [21] |
| SPR, MIP | 21 | [54] |
| SPR, competitive test, polyclonal antibody | 0.01 | [55] |
| Fluorescence/graphene-based platform, amplification | 5.6 | [56] |
| SWV/Au NP amplification | 2.0 x 10 ⁻⁵ | [57] |
| SWV/Three-way junction DNA structure, Fc-tagged cDNA, | 0.2 | [58] |
| EIS, target-induced aptamer displacement | 0.07 | [19] |
| EIS/MWCNT-SPE | 862 | [20] |
| EIS/Chitosan-graphene oxide | 28.5 | [59] |
| EIS, aptamer, on recognition-induced charge switching | 14 | [60] |

^a Abbreviations: SWV: Square Wave Voltammetry. EIS: Electrochemical Impedance Spectroscopy. MWCNT: Multi Wall Carbon Nanotubes. SPE: Screen-printed Electrode. NP: nanoparticle. MIP: Molecularly Imprinted Polymer

Table S2 Properties of proteins investigated in the selectivity study

| Analyte | Size (kDa) | Isoelectric point |
|----------------------|------------|-------------------|
| Lysozyme | 14.3 | 11 |
| Myoglobin | 17 | 7.2 |
| Cytochrome C | 12 | 10.2 |
| Bovine Serum Albumin | 66 | 4.7 |
| Salmon Calcitonin | 3.4 | 8.9 |

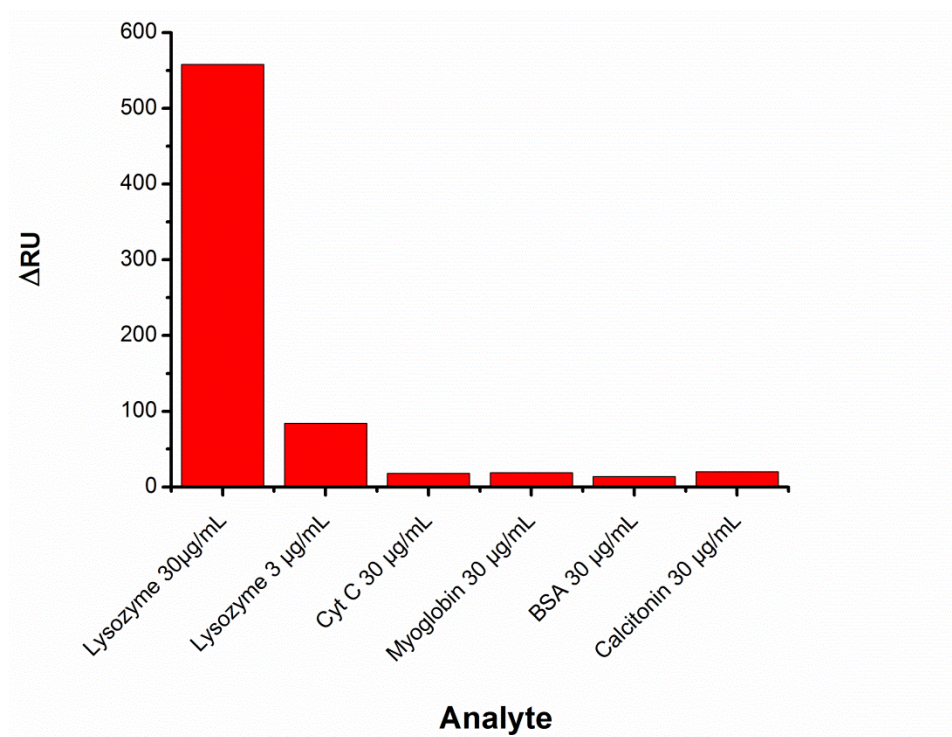


Figure S1. Net signal (calculated as $\Delta RU = \Delta RU_{\text{aptamer channel}} - \Delta RU_{\text{reference channel}}$) recorded with the aptasensor for lysozyme (at 3 $\mu\text{g/mL}$ and 30 $\mu\text{g/mL}$) and for several proteins at 30 $\mu\text{g/mL}$. Running buffer: 20 mM Tris-HCl pH 7.4 with 100 mM NaCl and 5 mM MgCl_2 . The protein solutions were injected for 5 minutes at 100 $\mu\text{L/min}$, followed by rinsing with buffer for 5 minutes at the same flow rate.