

Supporting Information

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SI Text

Consider the capture efficiency of ligand molecules by the nanoswitches.

Here, we describe how almost all ligand molecules will be bound to the nanoswitch at low ligand concentrations when the dissociation constant of the more strongly binding antibody K_{D1} is lower than the concentration of nanoswitch $[N]$ and the dissociation constant of the more weakly binding antibody K_{D2} is much lower than the local concentration between the two binding sites on the nanoswitch $[C]$. When the ligand is at a much lower concentration than the nanoswitch, we can assume that the change in free nanoswitch concentration and the proportion of nanoswitches that bind to two ligands are both insignificant. With $[L]$ as the ligand concentration and $[N]$ as the nanoswitch concentration, the proportion of looped nanoswitch $[N_{loop}]$ is given by the following:

$$[N_{loop}] = \frac{[C][L][N]}{K_{D1}K_{D2} + K_{D2}[N] + K_{D1}[N] + [C][N]}$$

Here, K_{D1} is the dissociation constant of the more strongly binding sandwiching antibody. If we add the assumptions that the concentration of the nanoswitch $[N]$ is much greater than the dissociation constant of the more strongly binding antibody and the local concentration $[C]$ is much greater than the dissociation constant of the more weakly binding antibody, then this expression reduces to $[N_{loop}] \approx [L][1 - (K_{D2}/[C])(1 + K_{D1}/[N])] \approx [L][1 - K_{D2}/[C]]$. For our system, we have the parameters $K_{D1} = 10$ pM and $K_{D2} =$

100 pM. Additionally, the local concentration $[C]$ is ≈ 30 nM (11) and the nanoswitch concentration $[N]$ is 0.7 nM, resulting in almost all ligand captured by looped nanoswitch with $[N_{loop}] \approx 0.999[L]$.

Here, we demonstrate how a nanoswitch can capture more ligand than just using a capture antibody. When capturing ligand with an antibody instead of a nanoswitch, the proportion of antibody–ligand complex is given by the following:

$$[A \cdot L] = \frac{[A][L]}{K_D + [A]}$$

Therefore, if we assume that for the nanoswitch both antibodies have similar affinities, the ratio between capture from a nanoswitch to capture from single antibodies is given by the following:

$$\frac{\text{Cap}_{\text{nanoswitch}}}{\text{Cap}_{\text{single}}} = \frac{[C](K_D + [A])}{K_D^2 + 2K_D[A] + [C][A]}$$

This quantity is greater than 1 when $[C] > K_D + 2[A]$, which generally holds true since the local concentration $[C]$ is ≈ 30 nM (11). As an example, if we used antibodies that had $K_D = 1$ nM, then nanoswitches would capture 2.2 times than just using capture antibodies.

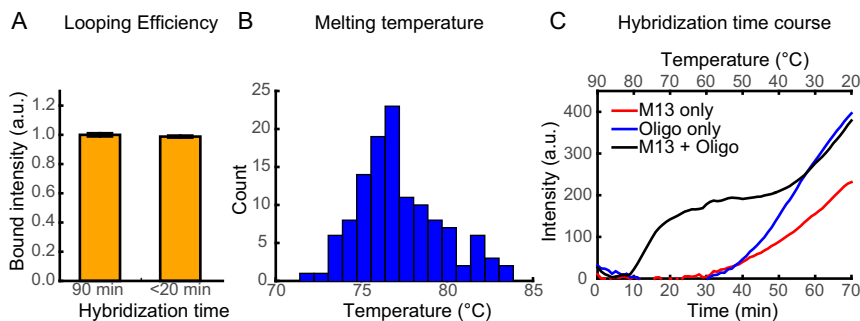


Fig. S1. (A) Bound intensity for long (90-min) and short (<20-min) nanoswitch construction times for 200 pM PSA. Hybridization conditions are described in *Methods*, and error bars are the SEM over four replicates. (B) Distribution of melting temperatures in the hybridization buffer (50 mM NaCl, 10 mM MgCl₂) for all 121 tiling oligonucleotides computed by the OligoAnalyzer Tool (IDT). (C) Measurement of annealing using SYBR Green I during a normal 90-min hybridization time course stepping down 1° per minute for only the M13 scaffold, only the tiling oligonucleotides, and both the M13 and tiling oligonucleotides.

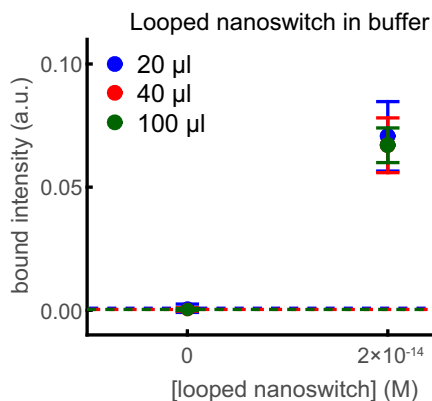


Fig. S2. Detection of looped DNA nanoswitch in buffer. Average bound intensity per lane as a function of concentration in buffer. Results are for 20 μL of sample run in one gel lane (blue), 40 μL of sample run across two gel lanes (red), or 100 μL of sample run across multiple gel lanes (green). Error bars are the SD over eight replicates, with the background given by the dashed line.

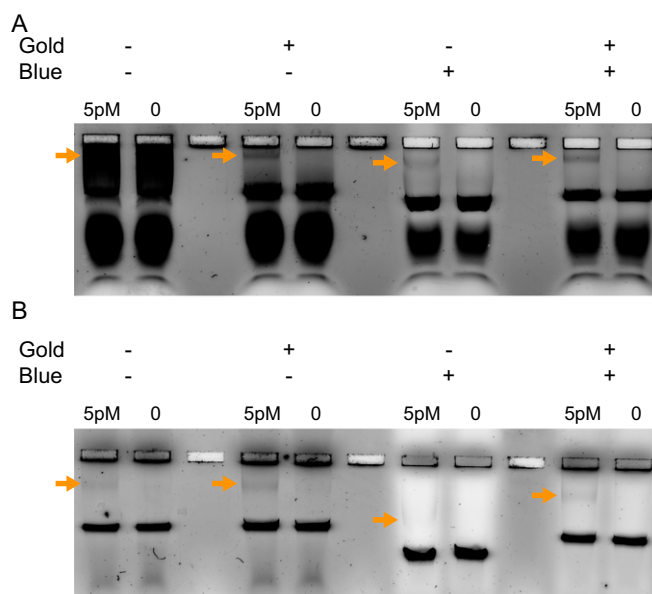


Fig. S3. Use of dyes to enable entry of nanoswitches into the agarose gel for separation. Gel electrophoresis results of 5 or 0 pM of PSA in (A) 20% FBS or (B) 20% bovine urine in the absence of any dyes, only SYBR Gold, only Coomassie Brilliant Blue G-250, or both SYBR Gold and Coomassie Brilliant Blue G-250. The location of the looped bands are specified by the orange arrows.

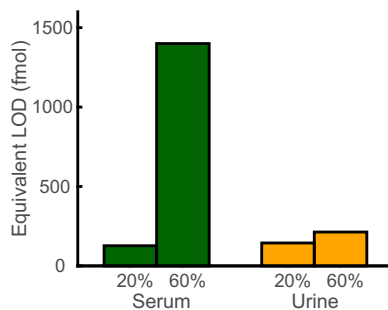


Fig. S4. Equivalent limit of detection (LOD) as a function of the percentage of biological material in the sample. The equivalent LOD is the LOD in 12 μL of undiluted sample. For a sample diluted to 20%, this means that the signal is averaged over three different lanes, while for 60% this is only one lane. The LOD was determined by extrapolating the concentration from the signal equal to background signal plus 3 SD of the background signal.

