

APTAMER INFORMATION

Hemoglobin aptamer 5E01 #391

1a. Description:

- *Identifiers:* 5E01 (Oligo #391)
- *Number of DNA nucleotides:* 40 bases (without 3'-6T), 46 bases (with 3'-6T)
- *Molecular weight (includes 3'-6T and biotin):* 14,729.7 g/mol
- *Target for selection:* Hemoglobin protein, Sigma (Cat #H7379)

Aptamer was selected from a randomized 40-mer library against Hemoglobin protein. Proprietary methods were then used to select the aptamer.

Aptamer folding instruction before use:

Once the aptamer is in its working concentration, it needs to be heated to 85-90 °C for 2 minutes, and then cooled to room temperature before use.

1b. Validation data with Hemoglobin protein by BLI (Bio-Layer Interference) method:

- *Immobilized Ligand:* Hemoglobin aptamer 5E01 with 3'-6T and biotin
- *Analyte:* Hemoglobin protein
- *Buffer used for validation:* 20 mM Tris, 100 mM NaCl, 0.005% Tween20 in nuclease free water, pH 7.4
- Equilibrium Binding Analysis, Avg $K_d = 4.3\text{nM}$

1c. Kinetics Screening Assay using Streptavidin Biosensors :

We validate the binding data by single reference method.

- *Single reference data:* All curves are referenced to a sensor dipped in buffer alone (no protein) (see Figures 1 & 2).

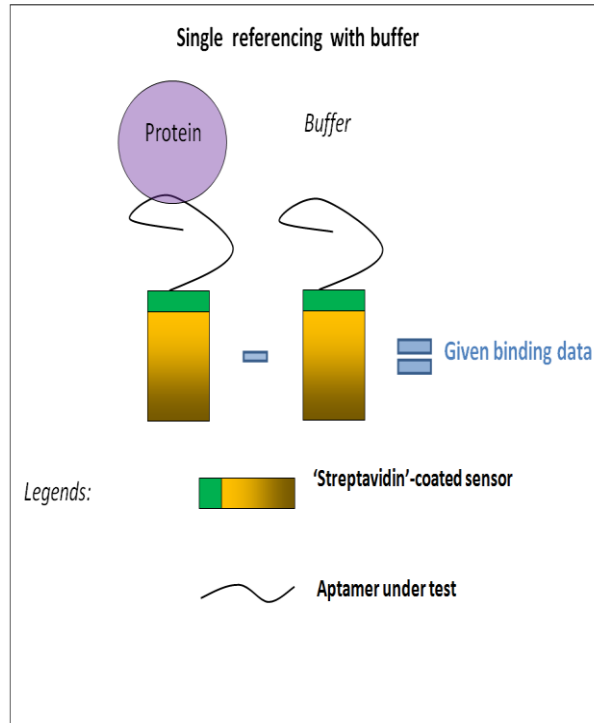


Figure 1. Diagram showing aptamer: protein binding validation scheme.



1d. Single reference data:

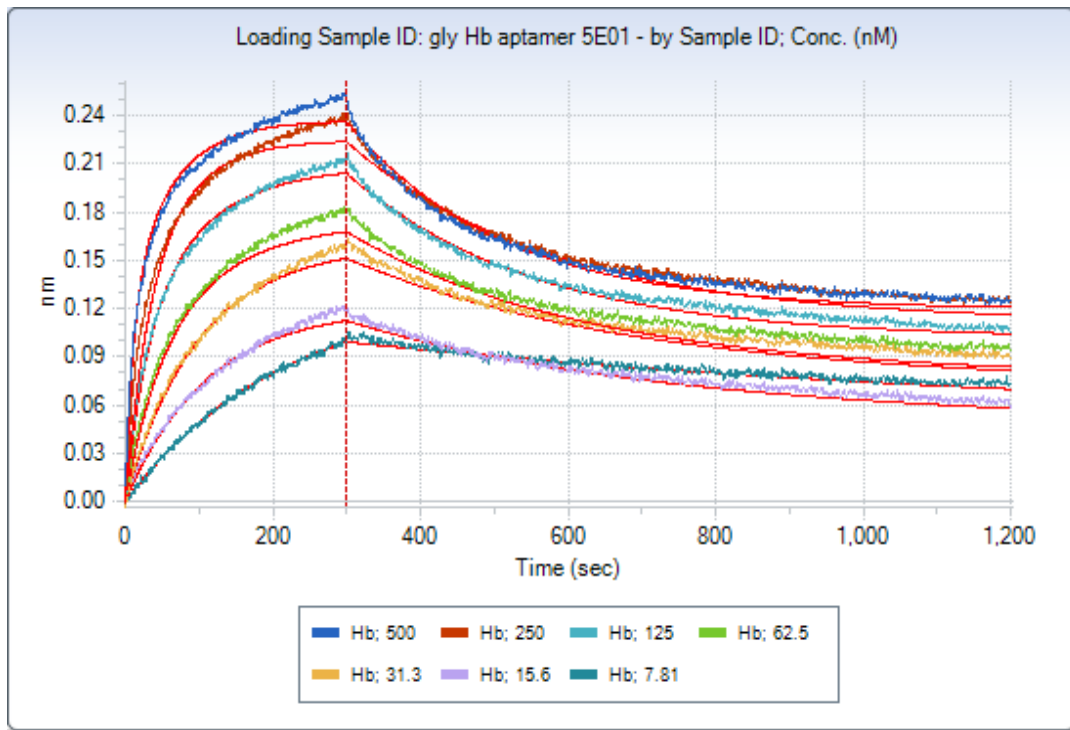


Figure 2. Association and dissociation graph of 2:1 (heterogenous ligand) fitting model of Hemoglobin aptamer 5E01(biotinylated) to Hemoglobin protein concentrations 500, 250, 125, 62.5, 31.3, 15.6, 7.81 nM by single reference method.

