



APTAMER INFORMATION Laminin -110 aptamer #203

1a. Description:

- *Identifiers*: 110 (Oligo# 203)
- Number of DNA nucleotides. 40 bases (without 3'-8AT), 58 bases (with 3'-8AT)
- Molecular weight (includes 3'-8AT and biotin): 18,287.1 g/mol
- <u>Target for selection:</u> Laminin, Sigma (Cat #L2020)

Aptamer was selected from a randomized 40-mer library against Laminin. 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 hCG protein using surface plasmon resonance (SPR):

- Immobilized Ligand: Laminin aptamer #203 with 3'-8AT and biotin
- Analyte: Laminin
- By Kinetic Analysis (Qdat), avg K_d = 5.0 ± 0.9 nM

(non-linear regression, pseudo-first order 1:1 interaction, Levenberg-Marquardt algorithm):







Overlaid fit:

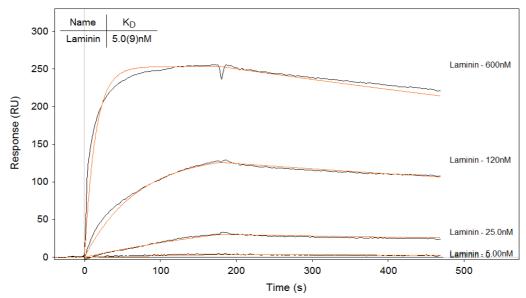


Figure 1. Overlaid fit of association and dissociation results of Laminin aptamer binding with Laminin protein concentrations of 600, 120, 25 nM;

	k _a (M ⁻¹ s ⁻¹)	k _d (s ⁻¹)	Rmax (RU)	K _D
Laminin protein	1.16 ±0.01e5	6 ±1e-4	255.3 ±0.5	5.0 ±0.9 nM

Table 1. Kinetics table showing rate and affinity constants obtained for Laminin aptamer binding to Laminin protein using the SensíQ system.

