

## APTAMER INFORMATION CMVgB\_377

## **Description:**

- *Identifiers*: CMVgB\_377
- Number of DNA nucleotides: 32 bases
- <u>Target for selection</u>: CMV gB

Aptamer was selected from a randomized 32-mer library against the CMV gB molecule. 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 CMV gB by MST (Microscale Thermophoresis) method:

- Buffer used for validation: 1X PBS, pH = 7.4, 1 mM MgCl<sub>2</sub>
- Average  $K_d = 15.1 \pm 2.1 \text{ nM}$

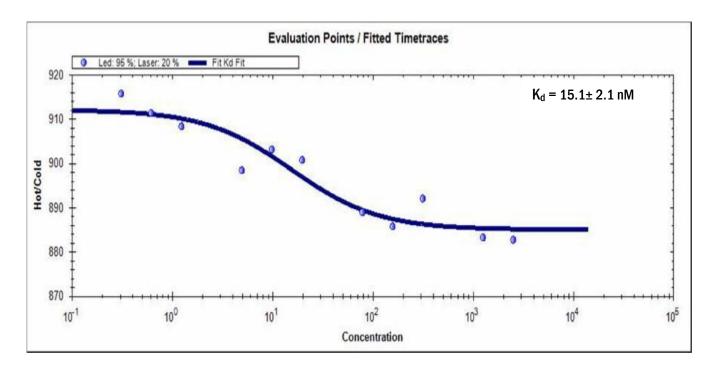


Figure 1. Aptamer-CMV gB binding. The thermophoresis is plotted versus the titrated CMV gB concentration.



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