

APTAMER INFORMATION CMVgB_40

Description:

- *Identifiers*: CmvgB_40
- 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 Peptide #3 by MST (Microscale Thermophoresis) method:

- Buffer used for validation: 1X PBS, pH = 7.4, 1 mM MgCl₂
- Average K_d = 17.2 ± 2.1 nM

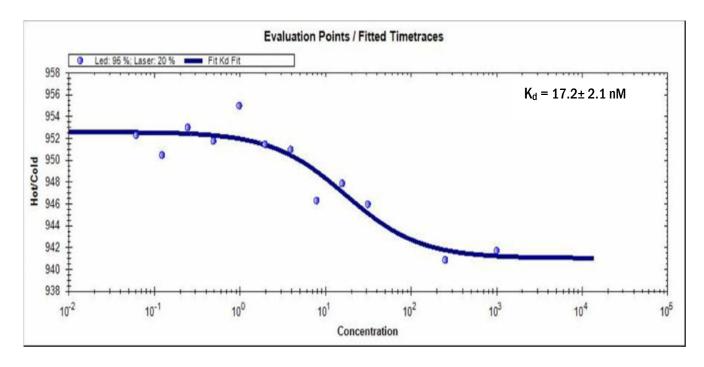


Figure 1. Aptamer-CMV gB binding. The thermophoresis is plotted versus the titrated CMV gB concentration.