

APTAMER INFORMATION Tetanus aptamers Tet15 & Thj899

1. Tetanus aptamer Tet15

1a. Description:

- Identifiers: Tet15
- Number of DNA nucleotides: 70 bases
- Target for selection: Tetanus toxoid, List Biological Laboratories, Inc. (Cat. #191A)

Aptamer was selected from a randomized 32-mer library against the Tetanus toxoid 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 Tet15by BSI (Back-Scattering Interferometry) method:

- <u>Buffer used for validation:</u> 1X PBS, pH = 7.4, 1 mM MgCl₂
- Average K_d = 15.9 ±9.0 nM



Figure 1. Aptamer-Tetanus toxoid binding. The phase shift of bound aptamer is plotted versus the titrated tetanus concentration. The values are the mean values from three independent measurements.

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2. Tetanus aptamer Thj899

2a. Description:

- Identifiers: Thj899
- Number of DNA nucleotides: 32 bases
- Target for selection: Tetanus toxoid, List Biological Laboratories, Inc. (Cat. # 191A)

Aptamer was selected to form a sandwich pair with the Tet15- Tetanus toxoid protein complex. See Figure 2.



Figure 2. Sandwich-array assay. The labeled Tet15-Tetanus toxoid protein complex was pre-incubated together then introduced to an aptamer microarray in an effort to discover a sandwich pair for this target. After binding the Tet15- Tetanus complex to the microarray, the array was washed and scanned for fluorescence. Bound aptamers subjected to MST analysis.

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 Thj899by MST (Microscale Thermophoresis) method:

- <u>Buffer used for validation</u>: 1X PBS, pH = 7.4, 1 mM MgCl₂
- Average K_d = 1.04 ± 0.17nM









