

Camelid single-domain antibody ciA-B5(ABCD_AW306) recognizes light chain of botulinum neurotoxin type A by ELISA

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Abstract

Camelid single-domain antibody (VHH) ciA-B5 (ABCD_AW306) detects by ELISA the light chain (LC) of the botulinum neurotoxin type A (BoNT/A).

Introduction

Botulinum neurotoxin type A (BoNT/A, UniProt #P0DPI1) poses a great threat to humans due to its most potent toxicity with the longest duration of paralysis. BoNT/A consists of a heavy chain (H_C) encompassing the receptor binding domain (H_C) and the translocation domain (H_N), and a light chain (LC) which is a zinc-dependent endopeptidase, capable of specific cleavage on neuronal SNARE proteins. Previous research has showed that VHH ciA-B5 can bind to the H_N domain of the heavy chain of BoNT/A and has demonstrated its ability to neutralize BoNT/A in a mouse model (Mukherjee *et al.*, 2012; Lam *et al.*, 2020). In the present study, we describe the reactivity in ELISA of this VHH against LC of BoNT/A.

Materials & Methods

Antibodies: The gene coding for ciA-B5 (ABCD_AW306, ABCD nomenclature, <http://www.expasy.org/abcd>) was codon-optimized for expression in *E. coli*, synthesized by Genscript, and cloned into pET22b (Novagen #69744-3) expression vector, which was modified to carry a FLAG tag (DYKDDDDK) at the C-terminal end for detection. His-tagged recombinant VHH was expressed in *E. coli* BL21(DE3) and purified by Ni-NTA spin columns (Qiagen #31014) following the manufacturer's instructions.

Antigen: The bont/LCA1 fragment (residues 1 - 420 of BoNT/A1) was cloned into pET45b vector with an N-terminal His-tag (Novagen #71327-3). LC/A1 was expressed in *E. coli* RosettaTM 2(DE3) and purified using Ni-NTA spin columns (Qiagen #31014), following the manufacturer's instructions. Hc/A1 fragment or receptor-binding domain of BoNT/A1 (residue 871-1296 of BoNT/A1) was produced the same way and used as a negative control.

Protocol: High Bind StripwellTM Microplates (Corning #07-200-24) were coated with 2 µg/mL recombinant antigens (LC/A1 and Hc/A1) in Phosphate Buffer Saline (PBS 1X) at 4°C overnight. Each well was rinsed twice with 300 µl of washing buffer (PBS 1X supplemented with 0.05% (w/v) Tween-20), then blocked for 1 hour with 300 µl of PBS 1X supplemented with 1% (w/v) BSA. After washing, wells were incubated for 2 hours at 37°C with 100 µl of the recombinant VHH diluted in PBS 1X with 0.5% (w/v) BSA. The wells were washed three times with washing buffer, then incubated with anti-DYKDDDDK (FLAG) Antibody Rabbit - HRP Conjugate (Immunology Consultants Lab #RFLG-45P) (dilution 1:10000) for 1 hour at 37°C. After six washes, Tetramethylbenzidine (TMB) substrate (Sigma #T5569) was added (100 µl per well). Reactions were stopped by adding 100 µl of 1 M HCl to each well. The experiment was run in triplicate, the optical density of each sample was analyzed at 450 nm with a reference reading at 630 nm. EC₅₀ values (the half-maximal effective concentration) were calculated via non-linear regression analysis using GraphPad Prism.

Results & Discussion

VHH ciA-B5 bound in a concentration-dependent manner to LC/A1 but did not bind to the Hc/A1 negative control (Fig.1). From the ELISA dose-response curve, EC₅₀ value was calculated to be approximately 67.3 nM. Of note, ciA-B5 displayed moderate affinity toward LC/A1 comparing to a panel of VHHs (produced in our laboratory) consisting of JPU-A5, JPU-C10 and JPU-D12 from Lam *et al.* (2020) and A1, A16 from Conway *et al.* (2010) (data not shown) (Conway *et al.*, 2010; Lam *et al.*, 2022).

In a previous study, ciA-B5 was reported to bind to the H_N region of BoNT/A1 (Lam *et al.*, 2020). This conclusion was based on the crystallization of a hetero-tetrameric complex (LCH_N/A-ciA-B5-ciAD-12-ciA-H7) including both LC and H_N of BoNT/A1, along with three VHHs (ciA-H7, ciA-D12, ciA-B5), in which ciA-H7 interacts with residues in the long helix α5 (Loop120, 170, and 250), and α10 of LC/A1; ciA-D12 interacts with residues in the C-terminal loop, the N-terminal loop, and α10 of

LC/A1; and ciA-B5 interacts to residues 600–616 in H_N domain of BoNT/A1. However, this report did not conclusively exclude the possibility that ciA-B5 can bind to the LC/A1. As both ciA-H7 and ciA-D12 interact with numerous regions of LC/A1, it is conceivable that interaction between ciA-B5 and LC/A1 could be obscured in the crystallographic analysis. In the present study, our results clearly demonstrated that ciA-B5 binds to LC/A1 in a concentration-dependent manner in ELISA assay. Therefore, we hypothesized that ciA-B5 can interact with both the LC and H_N domains of BoNT/A1.

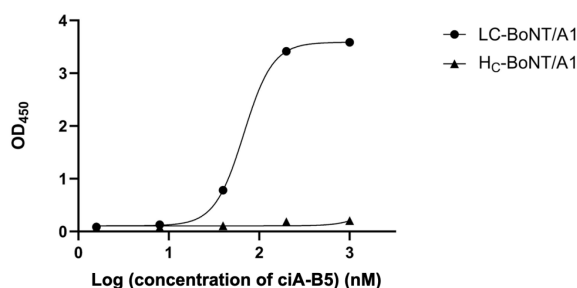


Fig. 1. ELISA dose-response curve of VHH ciA-B5 to LC-BoNT/A1 and H_c-BoNT/A1 as negative control.

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Conflict of interest

The authors declare no conflict of interest.

Data Availability Statement

The data that support the findings of this study are available from the corresponding author upon reasonable request.