Supplementary Materials: Structural Characterization of Humanized Nanobodies with Neutralizing Activity against the *Bordetella pertussis* CyaA-Hemolysin: Implications for a Potential Epitope of Toxin-Protective Antigen

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Figure S1. (a) Colony-PCR analysis of phage-transformed *E. coli* clones. 600-bp PCR products exclusively yielded by the vh/v_hh -positive clones are indicated. M, GeneRulerTM 1 kb DNA ladder (Thermo Scientific, Waltham, MA, USA). Each lane number corresponds to the clone number of phage-transformed *E. coli*; (b) Western blot analysis of lysate supernatants from the vh/v_hh -positive *E. coli* clones using anti-E tag antibodies. E-tagged VH/V_HH nanobodies expressed in the *E. coli* lysates were revealed as protein bands of ~17–22 kDa. M, pre-stained protein standards. Each lane number is referred to as the clone number of vh/v_hh -positive *E. coli*.



Figure S2. Expression of CyaA-Hly-specific nanobodies in pET vector system. (**a**) SDS-PAGE (Coomassie brilliant blue-stained 14% gel) analysis of lysates from *E. coli* expressing CyaA-Hly-specific His-tagged VHs/V_HHs under the control of T7/lac promoter; (**b**) Western blotting of **a** probed with anti-His tag antibodies. The expected ~17-kDa protein bands of VH/V_HH nanobodies are indicated. M, pre-stained protein standards. S and I, lysate supernatants and insoluble pellets after centrifugation, respectively.