

## Supplementary Material

Structure-Activity Correlations of Variant Forms of the B Pentamer of *Escherichia coli* Type II Heat-Labile Enterotoxin LT-IIIb With Toll-like Receptor 2 Binding

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## **Methods: Production and Purification of Toll-like Receptors TLR1 and TLR2**

In brief, the genes for TLR1 and TLR2 were supplied in the truncated transfer vector pVL1393 (Becton Dickinson, and Co, San Diego) as described (Jin et al., 2007). These plasmids were co-transfected separately with 0.5  $\mu$ g of BaculoGold Baculovirus DNA following the manufacturer's protocol. A positive control (pVL1392-XylE) was co-transfected as well. The negative control contained only sf9 cells. Cultures of  $2.0 \times 10^6$  cells were plated in sterile Petri dishes for each co-transfection and were incubated overnight at 27°C. Cells reached >75% confluence. Cultures with the DNA insert were incubated overnight at 27°C for four days. On day 6, viral supernatant was harvested and labeled P1 virus and stored at -80°C. P2 and P3 viral supernatant was generated by amplification of the P1 stock. P2 amplification was made with  $0.5 \times 10^7$  sf9 cells that were infected with 500  $\mu$ L of viral supernatant (P1) in 15 mL of BakPak Complete medium (Clonetechn) in a vented T-75 flask. The amplification was allowed to incubate at 27°C for three days. Harvested supernatant (P2 viral stock) was determined to be  $2.21 \times 10^7$  pfu/mL via plaque assay using the manufacturer's recommendations. A second round of amplification (P3) was performed in a 75 mL suspension culture at an MOI of 0.1 for three days. Titer assay showed  $2.45 \times 10^8$  pfu/mL. P3 virus was used for all large scale infections/expression (6-8L of sf9 cells) studies that were carried out in oxygenated spinner flasks that were stirred at 80 rpm with oxygenation of 30%. Cells were infected with virus optimized at a MOI of 7.5 for TLR1 and at MOI of 11.0 for TLR2. Cells were incubated for 72 hours or until cell viability was between 68-75% via trypan blue exclusion. Protein expression levels were reduced to less than 1 mg/L if cells were harvested with viability higher than 75%. Typical yields were 2.9 mg/L for TLR1 and 1.8 mg/L for TLR2.

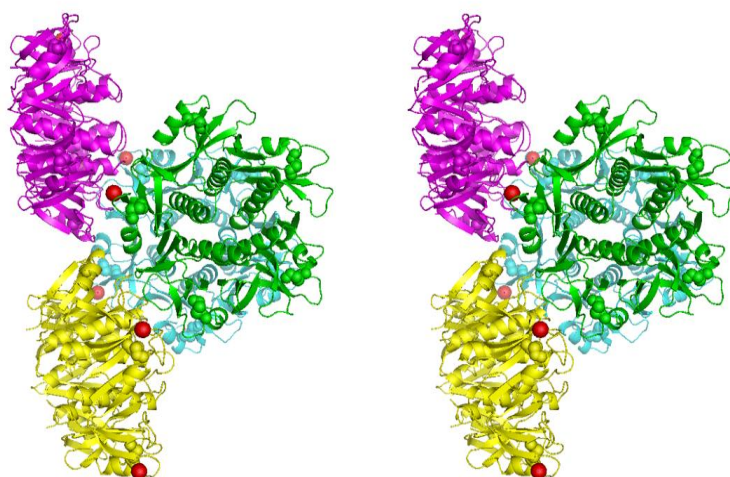
### **Purification of TLR1 and TLR2**

Cells were resuspended in Buffer A (50mM Tris HCl, pH-7.4, 150 mM NaCl, and 0.2 mM TCEP) and then lysed using a Branson Sonifier set at 70% intensity with 3 cycles of 30s on followed by 30s off while on ice. Lysed cells were centrifuged in a Ti-45 rotor set at 45,000 rpm for 45 minutes at 4°C to pellet debris. The resulting supernatant was filtered through a 0.22  $\mu$ M syringe

filter and applied to a Protein A column. TLR1-IgG1 (Fc) or TLR2-IgG (Fc) was eluted using Buffer B (100mM Na citrate, pH 4.5). The resulting peak containing the TLR of interest were quickly exchanged into Buffer A using a Centricon YM-10 filter due to instability at low pH conditions. The protein was added to a GF200 column pre-equilibrated with Buffer A. Peak fractions were verified via SDS-PAGE and flash frozen in liquid nitrogen and stored at  $-80^{\circ}\text{C}$ .

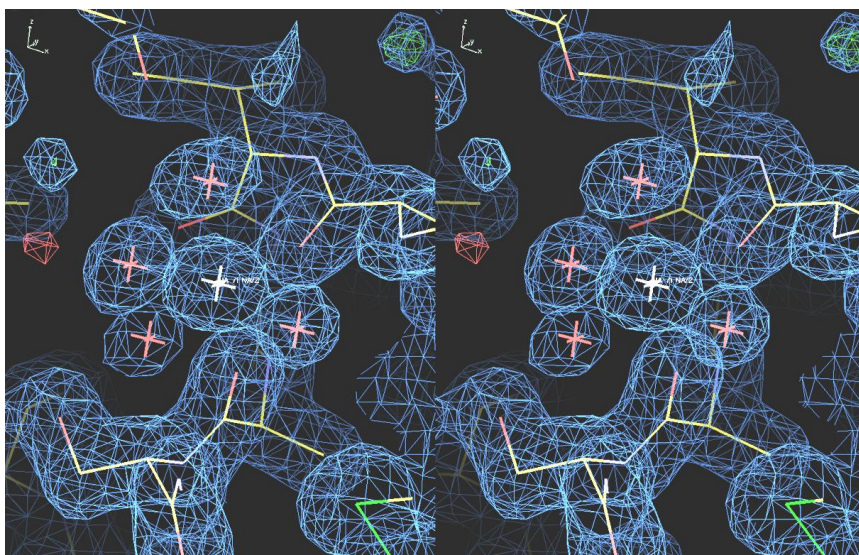
### **Packing arrangement in the LT-IIb-B<sub>5</sub>(T13I) variant**

Structural results for the LT-IIb-B<sub>5</sub>(T13I) variant revealed the presence of eight Na<sup>+</sup> ions present in the crystallization media. The Na<sup>+</sup> ions form octahedral binding interactions utilizing the backbone carbonyls of residues Arg12 and Ala15 and with water with contact distances between 2.3 – 2.5Å (Fig. S1). The Na<sup>+</sup> ions are coordinated with three types of interactions: Arg12, Ala15 and four waters (Fig. S2a); Arg12, Ala15, Ser16 and three waters; or Arg12, Ala15, Asn33 from a symmetry-related molecule (Fig. S2b). These interactions help stabilize the four unique pentamers in the asymmetric unit of the crystal lattice.

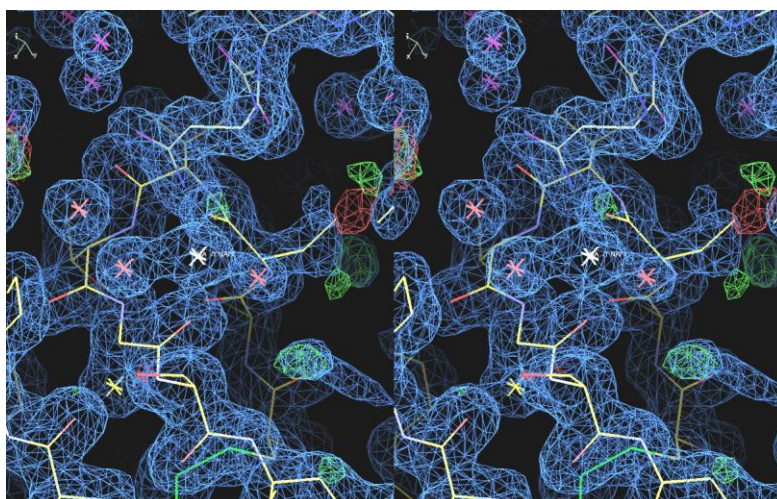


**Supplementary Fig. S1.** Packing arrangement of the four pentamers (green,cyan,violet and yellow) in the structure of the LT-IIb-B<sub>5</sub>(T13I) variant (residue 13 as space filling representation) highlighting the interactions of the Na<sup>+</sup> ions (red spheres) present in the crystallization buffer.





(a)



(b)

Supplementary Fig. S2. (a) Example of the environment around one of the  $\text{Na}^+$  ions (white X) in the structure of the LT-IIb-B<sub>5</sub>(T13I) variant. The octahedral coordination uses the carbonyl oxygens from Arg12 and Ala15 and water (red X) molecules (2.3-2.5Å). (b) Example of the environment around one of the  $\text{Na}^+$  ions (white X) in the structure of the LT-IIb-B<sub>5</sub>(T13I) variant. The coordination uses the carbonyl oxygens from Arg12 and Ala15, the amine from Asn33 from a symmetry-related molecule, and water (red X) molecules (2.3-2.5Å).