PS-10-8 Poster Session

## Determining the role of TIR domain of Interleukin-1 receptor 8 (SIGIRR) in regulating TLR4 signalling

Surekha Nimma<sup>1</sup>, Jeffrey Nanson<sup>1</sup>, Thomas Ve<sup>1,2</sup> and Bostjan Kobe<sup>1,3</sup>

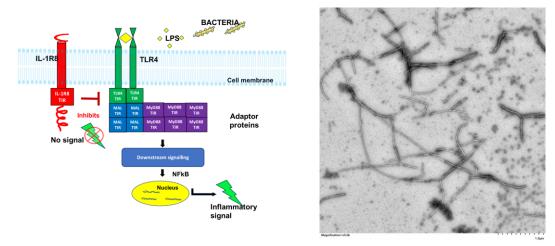
<sup>1</sup>School of Chemistry and Molecular Biosciences and Australian Infectious Diseases Research Centre, University of Queensland, Brisbane, QLD 4072, Australia,

<sup>2</sup>Institute for Glycomics, Griffith University, Southport, QLD 4222, Australia, <sup>3</sup>Institute of Molecular Bioscience, University of Queensland, Brisbane, QLD 4072, Australia

s.nimma@uq.edu.au

The TIR superfamily includes membrane receptors, Interleukin-1 receptors (IL-1Rs) and Toll-like receptors (TLRs) and also TIR-containing cytoplasmic adaptor proteins such as MAL and MyD88. These proteins play a major role in immune signalling and are vital to innate host defense, inflammation, injury and stress [1]. IL-1R8, also known as single immunoglobulin interleukin-1 receptor-related protein (SIGIRR) is an inhibitory receptor from IL-1R family which regulates signalling of both IL-1Rs and TLRs. The mechanism of inhibition is not yet known, but the only available genetic evidence suggests that the conserved intracellular TIR domain of IL-1R8 alone is necessary to inhibit LPS-induced TLR4 signalling [2]. The recent cryo-EM structure of the MAL protofilament has revealed the molecular mechanism of TIR-TIR interactions in the MAL and MyD88 dependent TLR4 signalling [3]. Based on this, we hypothesize that a similar TIR:TIR interaction between the TIR domain of IL-1R8 and the TIR domains of either TLR4/MAL/MyD88 would be involved in the inhibition mechanism.

The TIR domain of human IL-1R8 was cloned, expressed and purified using E. *coli* host system. Turbidity assays, negative-stain electron microscopy (EM) and single-molecule fluorescence spectroscopy (SMFS) analysis indicated a potential interaction between IL-1R8<sup>TIR</sup> and MAL<sup>TIR</sup>. MAL<sup>TIR</sup> forms filamentous assemblies when incubated with IL-1R8<sup>TIR</sup> (Fig. 1). We are currently focusing on solving the 3D structure of MAL<sup>TIR</sup>/IL-1R8<sup>TIR</sup> filaments using negative-stain EM and cryo-EM to obtain molecular insights into the interaction interfaces and binding sites of IL-1R8<sup>TIR</sup> and MAL<sup>TIR</sup>. This study will eventually lead to an understanding of how TLR4 signalling is regulated by IL-1R8 and can potentially pave way in development of new therapeutic agents in future.



**Figure 1.** Left: Model representing the inhibition of TLR4 signalling by IL-1R8. Right: Negative-stain EM image of MALTIR/IL-1R8TIR filaments taken using Hitachi HT 7700 TEM.

[1] Boraschi, D. et al. (2018). Immunol Rev. 281, 97-232

[2]. Qin, J. et al. (2005). J Biol Chem. 280, 25233-25241

[3]. Ve, T. et al. (2017). Nat. Struc. Mol. Biol. 24, 743-751

Keywords: Interleukin-1 receptor 8; TLR4 signalling; MAL/TIRAP; TIR domain; cryo-EM

Acta Cryst. (2021), A77, C785