

Unravelling SAA-mediated immune responses to bacterial-derived amyloids in the central nervous system

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Recent studies indicate that gut inflammation can exacerbate the risk of developing Alzheimer's Disease (AD). Our lab found that bacterial amyloids act as potent immune inducers in the gastro-intestinal tract and its enteric nervous system (1). In this context, serum amyloid A3 (SAA3) emerged as a key regulator driving a pro-inflammatory feed-forward response, characterized by cytokine secretion and T-cell infiltration. Since SAAs are acute-phase proteins with pro-inflammatory and amyloidogenic properties, which are elevated in the brains of AD patients, we asked whether they play a role in pathogenic gut-brain communication in AD. To investigate this, we administered intraperitoneal injections of the bacterial amyloid curli in C57Bl6 mice. This caused a general increase in circulating SAA protein levels and a specific, restricted elevation of Saa3 expression at brain borders. As a more direct means to interrogating the putative function of SAAs in AD pathology progression, we stereotactically injected adeno-associated viral vectors that cause constitutive Saa3 expression into the hippocampus of one-month-old APP/MAPT mice. Through RNA-Scope and quantitative immunofluorescence staining, we will evaluate the impact on microglial activation and A β load, respectively. Lastly, we are investigating whether this SAA loop is conserved in humans by exposing iPSC-derived microglia progenitor cells to curli and A β . Preliminary results indicate that these cells produce SAA1/2 in response to curli, making them a good human model to further investigate. By uncovering the role of SAAs in early immune-driven mechanisms of AD pathogenesis, we aim to provide novel insights that contribute to the development of disease-modifying therapies for this devastating disorder.

(1) Verstraelen et al (2024). CMGH, 18(1), 89–104.