

# Solid State NMR Explorations of Protein-Lipid Interactions in Cardiovascular Disease

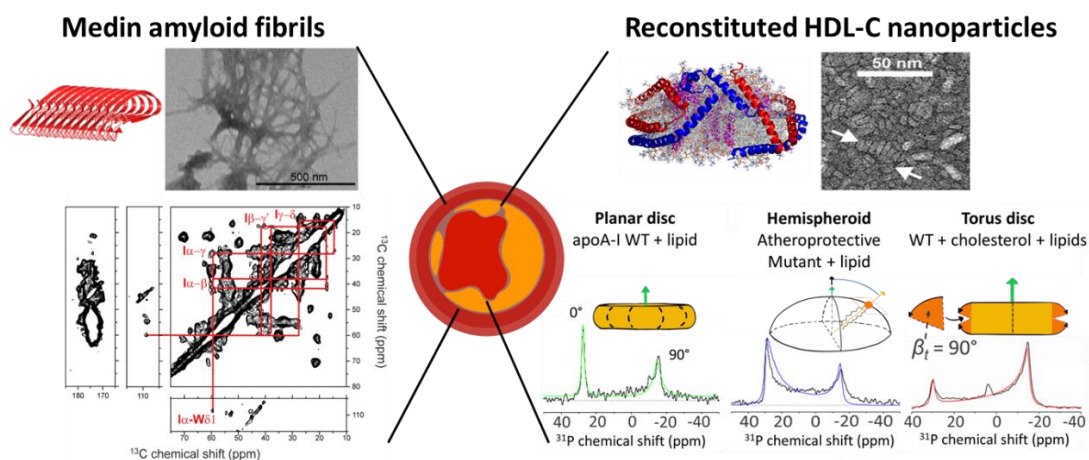
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## Abstract

We have utilised and developed solid-state NMR (ssNMR) methods to investigate functionally important phospholipid-protein interactions associated with cardiovascular disease. This could be a common factor in the formation of arterial plaques and cholesterol transport. Medin is a highly prevalent polypeptide found in blood vessel amyloid plaques of individuals over the age of 50, and has possible associations with Alzheimer's disease via its co-aggregation with the A $\beta$  peptide.<sup>1, 2</sup> Using the high-field ssNMR facility, we obtained the first structural details of medin fibrils.<sup>3</sup> Recently, we have used MAS and oriented ssNMR to characterise medin on the surface of liposomes mimicking extracellular vesicles isolated from smooth muscle cells.<sup>4</sup> The results provide insight into the aortic amyloid formation pathway and the role of phosphatidylserine lipids.

Secondly, we have developed ssNMR methods to probe protein-lipid interactions in high-density lipoprotein (HDL-C, "good cholesterol"), as HDL-C carries cholesterol to the liver for excretion. Our oriented <sup>31</sup>P ssNMR method can detect the rHDL-C morphology<sup>5</sup>, and using the dynamically averaged <sup>13</sup>C-<sup>13</sup>C and <sup>13</sup>C-<sup>1</sup>H dipolar couplings we have determined the orientational distribution of [2,3,4-<sup>13</sup>C<sub>3</sub>]cholesterol in discoidal rHDL-C.<sup>6, 7</sup> We are now investigating differences in the structure and function of atheroprotective and dysfunctional variants of HDL-C containing mutants of the main protein (apoA-I), including at high-field (1 GHz). Our overarching goal is to use the structural knowledge to develop improved diagnostic and preventative approaches to cardiovascular disease.



**Figure 1.** ssNMR can be used to detect site-specific structural features in aggregated medin amyloid fibrils and different HDL-C nanoparticle morphologies. Reproduced AAMD HDL model obtained from Pourmoussa *et al.*<sup>8</sup>

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