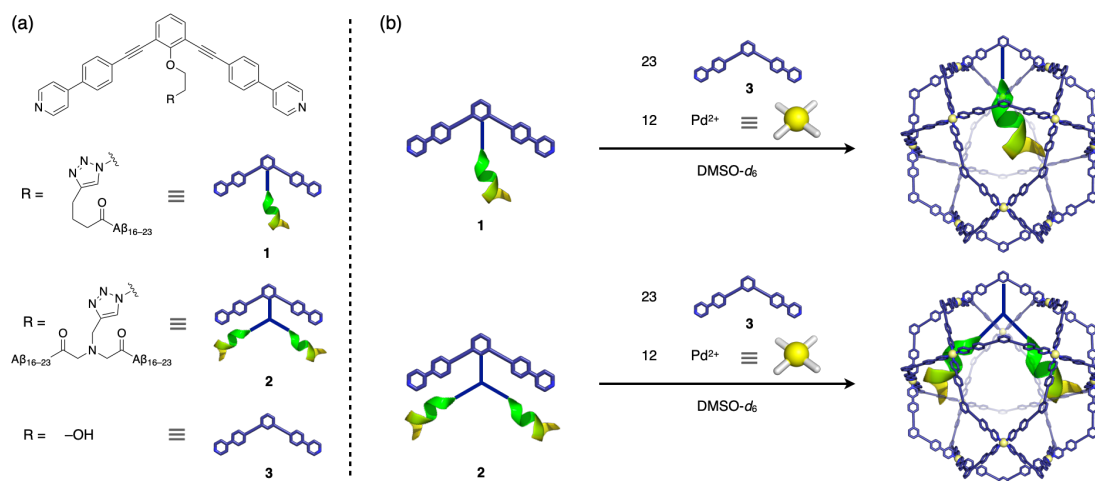


Structural analysis of amyloid β hydrophobic core in the oligomeric state by encapsulation within an $M_{12}L_{24}$ cage

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Amyloid β ($A\beta$) is an aggregative peptide consisting of around 40 amino acids residues. $A\beta$ is the main component of amyloid plaques often found in the brains of Alzheimer's disease patients. Recent studies show the soluble oligomeric state of $A\beta$ exhibits significant neurotoxicity, but the nature of the $A\beta$ oligomer is largely unexplored due to the rapid formation of insoluble $A\beta$ fibril through the uncontrollable aggregation.¹ We previously found that the encapsulation within an $M_{12}L_{24}$ cage, self-assembled from palladium(II) ions (M) and organic bidentate ligands (L), effectively prevents undesired aggregation of proteins including $A\beta$ proteins.² In this research, we sought to analyze the oligomeric structure of $A\beta$ by encapsulating a specified number of $A\beta$ fragments into an $M_{12}L_{24}$ cage.

The $A\beta_{16-23}$ fragment (H-Lys-Leu-Val-Phe-Phe-Ala-Glu-Asp-NH₂), which is the core of the hydrophobic β -strand, was studied to elucidate the $A\beta$ oligomeric structure. One or two molecules of the $A\beta_{16-23}$ fragments were tethered to bis(pyridine) ligands to afford **1** or **2**, respectively (**figure a**). $M_{12}L_{24}$ cages containing only one or two $A\beta_{16-23}$ fragments were self-assembled from **1** or **2** with ligand **3** and palladium(II) ions in DMSO-*d*₆ (**figure b**). The structures of encapsulated peptides were analyzed by 3D NMR measurements.



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