

Purification and characterization of a better-behaved Alzheimer's disease protein

Alzheimer's Disease (AD) is an incurable disease that causes memory, reasoning and thinking to deteriorate over time. AD is marked by brain atrophy and plaque-like deposits of a small peptide called amyloid beta ($A\beta$) outside of brain cells and tangles of a protein called tau inside brain cells. Although plaques are no longer considered to cause AD, small aggregates of toxic $A\beta$ (oligomers) are currently thought to play a role in the AD process. To study how the $A\beta$ peptide may contribute to AD, one needs pure $A\beta$. Unfortunately, the chemically made (synthetic) $A\beta$ peptide ($A\beta$ syn) is very difficult to use as it needs extremely strong solvents to dissolve. To overcome this problem, we created a gene for $A\beta$ and added a solubility tag consisting of 6 extra lysine amino acids (6K), at the start (N-amino terminal) or the end (C-carboxyl terminal) of $A\beta$ to determine which addition made $A\beta$ more water soluble. We then engineered bacteria (*Escherichia coli*) to 'express' the gene and produce the modified $A\beta$ peptide in large quantities.

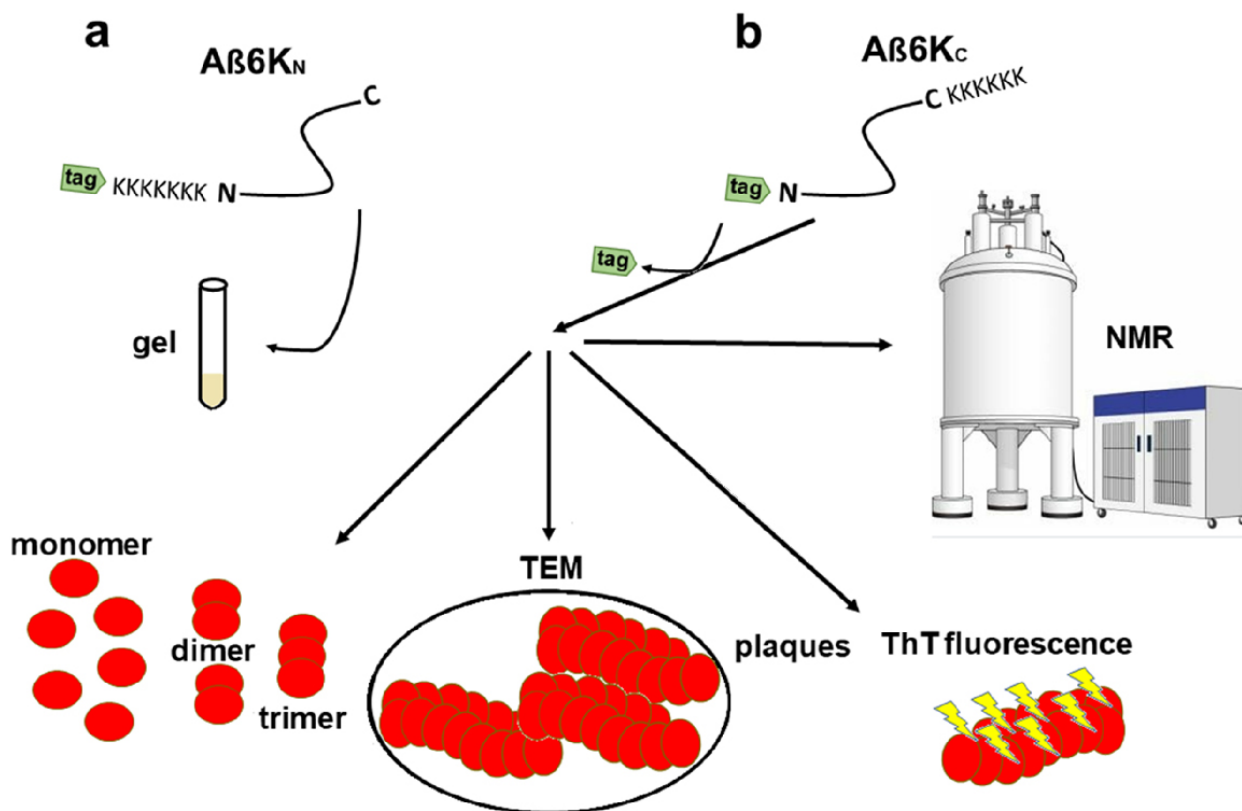


Fig. 1. Generation and characterization of a better behaved $A\beta$ peptide. We added a water solubility tag (6 lysines-KKKKKK) to the start (N-amino-terminal) or end (C-carboxyl-terminal) of the $A\beta$ peptide to determine which addition increased $A\beta$ solubility. (a) Adding KKKKKK to the start of $A\beta$ ($A\beta 6K_N$) caused a gel to form during purification while (b) adding KKKKKK to the end of $A\beta$

(A β 6K_C) increased expression in bacteria. The A β 6K_C was easier to purify. We used three different chromatography steps to purify A β 6K_C with the affinity tag (Histidine 6X tag) being removed before the last step. A β 6K_C behaved like the synthetic A β , forming oligomers (dimers, trimers) seen on protein gels, forming plaques that bound thioflavin T (ThT) and fluoresced. It also formed plaques that were seen by transmission electron microscopy (TEM). A β 6K_C had the expected sequence and shape of other A β proteins as seen by nuclear magnetic resonance (NMR) spectroscopy.

When the 6K was added to the start (A β 6K_N), the peptide was poorly expressed in bacteria and was difficult to purify. It formed a gel which prevented us from purifying it further. When the 6K tag was added to the end (A β 6K_C), the peptide was well expressed and easy to purify. We used three different separation techniques (affinity, ion exchange and reverse phase chromatography) to purify the A β 6K_C. We then tested if the A β 6K_C behaved similarly to A β syn. Using a cross-linking agent, we chemically trapped A β syn or A β 6K_C oligomers and visualized them on a gel that separates molecules according to size. We expected to see A β syn or A β 6K_C clusters: dimers, trimers, tetramers and larger. Our A β 6K_C aggregated similarly to the A β syn except it formed more dimers and trimers. We tested if A β 6K_C could form plaques using a marker called thioflavin T (ThT) which binds to plaques and fluoresces when more plaques form. Both A β 6K_C and A β syn could form plaques that fluoresced with ThT to the same extent and on a similar time frame, with A β 6K_C forming plaques a few minutes faster. This was expected as previously reports demonstrated that bacterial purified A β aggregated faster than A β syn. We then examined A β syn and A β 6K_C before and after plaque formation by transmission electron microscopy (TEM). The A β 6K_C before aggregation was more homogenous (all peptide aggregates being the same size) compared to A β syn that had different sized aggregates. After aggregation and plaque formation, A β syn plaques were more distinct than the A β 6K_C plaques. We then tested if the individual amino acids in A β 6K_C were similarly organized and that A β 6K_C had the same shape as other A β using nuclear magnetic resonance (NMR) spectroscopy. For NMR, we need a peptide with specific nitrogen and carbon isotopes that are 'visible' in NMR. We accomplished this by growing the bacteria with isotopically labeled nitrogen (ammonium chloride) and carbon (glucose) and purified the isotopically labeled A β 6K_C. The labelled A β 6K_C dissolved easily in water and was used in specially designed NMR experiments that allowed us to map the structural and atomic details of this peptide. This well-behaved, water-soluble, conveniently expressible version of A β forms plaques and can be induced to form oligomers, which are thought to contribute to the AD process. We plan to conduct more NMR experiments with purified oligomers to see how they interact with each other and to discover potential drugs to interrupt this oligomerization process and halt AD progression.

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