

Amyloid β -PeptidesSupramolecular Bait to Trigger Non-Equilibrium Co-Assembly and Clearance of A β 42Te-Haw Wu⁺, Rai-Hua Lai⁺, Chun-Nien Yao⁺, Jyh-Lyh Juang,^{*} and Shu-Yi Lin^{*}

Abstract: In living systems, non-equilibrium states that control the assembly-disassembly of cellular components underlie the gradual complexification of life, whereas in nonliving systems, most molecules follow the laws of thermodynamic equilibrium to sustain dynamic consistency. Little is known about the roles of non-equilibrium states of interactions between supramolecules in living systems. Here, a non-equilibrium state of interaction between supramolecular lipopolysaccharide (LPS) and A β 42, an aggregate-prone protein that causes Alzheimer's disease (AD), was identified. Structurally, A β 42 presents a specific groove that is recognized by the amphiphilicity of LPS bait in a non-equilibrium manner. Functionally, the transient complex elicits a cellular response to clear extracellular A β 42 deposits in neuronal cells. Since the impaired clearance of toxic A β 42 deposits correlates with AD pathology, the non-equilibrium LPS and A β 42 could represent a useful target for developing AD therapeutics.

Non-equilibrium states,^[1–3] or more specifically, far-from-equilibrium conditions have been found to be critical for the modulation of biomolecular assembly in living systems.^[4] One interesting example is the role of microtubules in the formation of cytoskeleton networks, an assembly process that exhibits a non-equilibrium behavior for its self-assembly and infrequent decay.^[5] Another example is the amyloid peptides (A β), which undergo self-aggregation to form amyloid plaques in triggering the neurodegenerative cascade of Alzheimer's disease (AD).^[6–8] Among the various A β isoforms, A β 40 and A β 42 are the two most abundant species. Although A β 40 is more abundant than A β 42 in cerebrospinal fluid, A β 42 has a higher self-aggregation potential than A β 40 in contributing to the amyloid deposits in AD brains.^[9] Intriguingly, in addition to its pathological role in AD, A β 42

has recently been implicated in antimicrobial activity against bacterial infection.^[10–12] The self-propagating A β 42 may kill bacteria by agglutinating the polysaccharide of bacterial outer membranes.^[13] From the perspective of chemical kinetics, the self-propagation of A β 42 represents a dynamic self-assembly process involving a transition state from the pre-organized protofibrils to a stable form of fibrils.^[14–19] On the other hand, LPS acts as an amphiphilic supramolecule with characteristics of spontaneous self-assembly in a concentration-dependent manner when dispersed in solution.^[20] One recent intriguing feature identified in AD brains is the co-localization of bacterial debris (i.e., LPS) with A β 42 deposits.^[21] The large amounts of LPS in the brain might elicit inflammatory responses in promoting AD development.^[22] However, one cannot exclude the possibility that there could be a physiological or protective function that has not been defined in normal or early AD brains. Inspired by this, we proposed that these two supramolecules might form an intermolecular interaction in a non-equilibrium state that could subsequently impact on the survival of neural cells (Figure 1).

To test this hypothesis, we investigated whether LPS could form a transient complex with A β 42 through a non-equilibrium co-assembly process that subsequently leads to dissociation. Bis-ANS, a specific fluorescent dye sensitive to A β 42 hydrophobicity,^[23] was used to assess the degree of A β 42 hydrophobicity during the process of amyloid polymerization upon the administration of LPS. The results revealed

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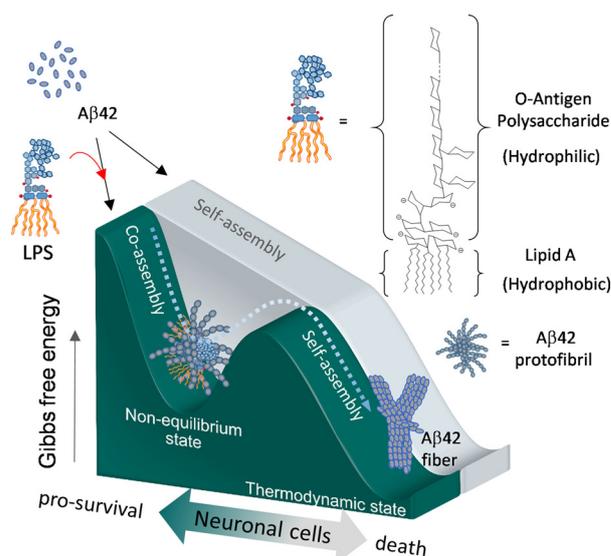


Figure 1. A hypothesized model describing that LPS might behave as a supramolecular bait to trigger non-equilibrium co-assembly with A β 42 protofibrils for pro-survival effect of neuronal cells.

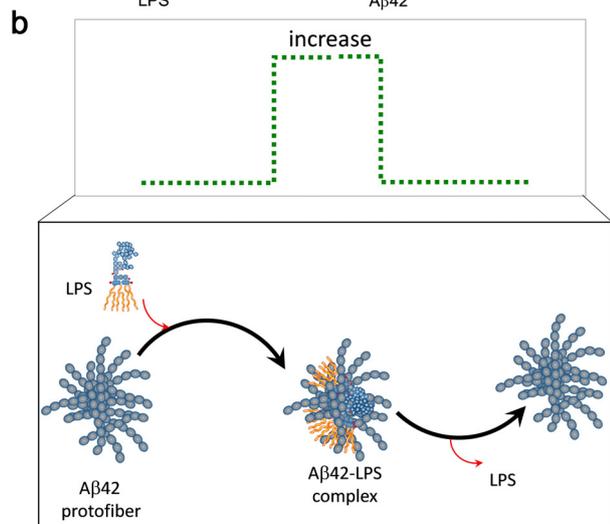
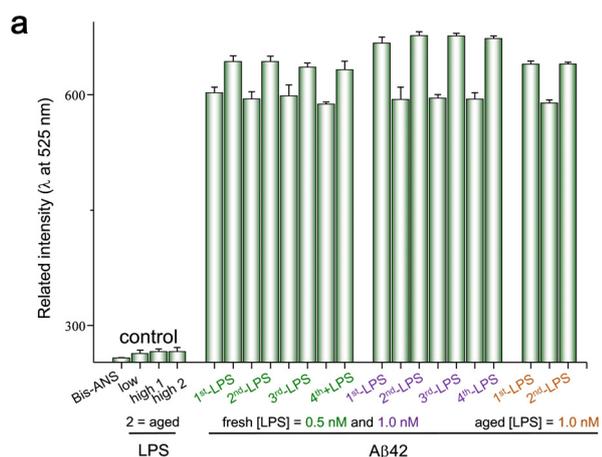


Figure 2. Confirmation of non-equilibrium interaction between LPS and A β 42 protofibrils. a) Oscillation of the hydrophobicity of A β 42 was noted by repeatedly adding freshly prepared LPS or aged LPS. The y-axis of the graph represents the relative fluorescence intensity of the bis-ANS emission wavelength at 525 nm. b) A simple illustration showing the hydrophobicity of A β 42 protofibrils could increase and decrease by LPS influx and efflux, respectively.

a repeated oscillating pattern of association and dissociation between the two molecules (Figure 2a). Specifically, the A β 42 hydrophobicity was noticeably increased and then rapidly reduced back to a baseline level within a 30 min incubation period after adding fresh LPS. This result suggested that the A β 42 protofibrils might induce transient LPS-A β 42 binding when they first encounter LPS in solution (Figure 2b). After the rapid growth of hydrophobicity, the continuing oscillation became well dampened if freshly prepared LPS was re-administered, with more pronounced effects being noted if LPS of higher concentration was re-administered. Similar effects were noted if an aged LPS (LPS being self-incubated overnight) was re-administered, but the dampening of A β 42 hydrophobicity became small. Since LPS is also an amphiphilic supramolecule that favors self-aggregation,^[24] we reasoned that aged LPS might become less competent than fresh LPS in forming LPS-A β 42 complex. Despite the amyloid fibrils being found to continuously form in the presence of LPS over the prolonged incubation time, the

formation of long fibers of A β 42 was clearly weakened (see Figure S1 in the Supporting Information). These results suggest a dissipative non-equilibrium state of self-aggregation for A β 42 might be induced by LPS.

Next, we investigated whether the transient association between LPS and A β 42 protofibrils impacts cytotoxicity. Surprisingly, cell viability assays suggested that SH-SY5Y neuronal cells retained more than 95 % cell viability after co-treatment of A β 42 with LPS. In contrast, less than 20 % cell viability was retained in cells treated with A β 42 alone (Figure S2, columns 2 and 3). To evaluate if the rescue effect was mediated through the LPS-A β 42 transient complex, we removed the unbound LPS from the solution and found that the rescue effect of LPS was completely lost (Figure S2, columns 3 and 4). However, when LPS were re-added back into the medium, the cell viability was returned to 85 % of the normal value in control cells (Figure S2, columns 1 and 5). Moreover, the rescue effect of LPS on the A β 42-induced apoptosis was found to occur in a dose-dependent manner with increasing LPS (Figure 3a). That being the case, how is this accomplished? Under normal conditions, both the oligomeric (protofibrils) and the monomeric forms of A β 42 can be internalized from the extracellular domains of brain cells for degradation.^[25,26] However, during the pathogenesis of AD, the process of A β 42 protofibril clearance via the endocytic pathway is interrupted, leading to an increased deposition of A β 42.^[27,28] Accordingly, we speculated that the

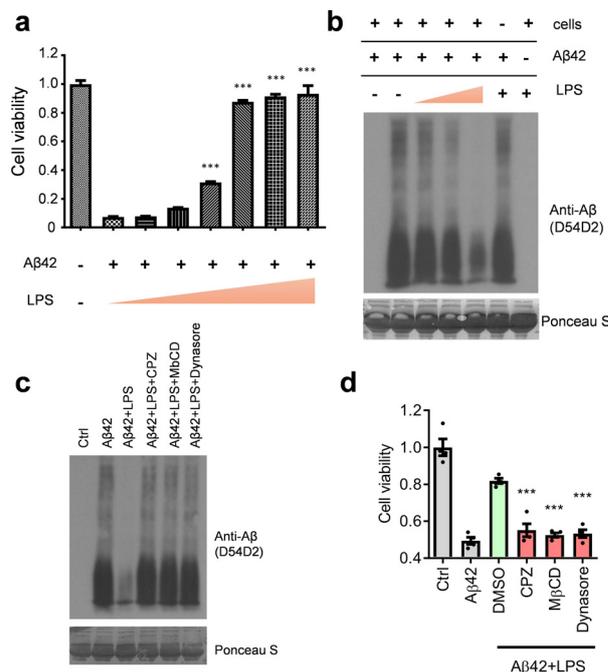


Figure 3. A β 42 peptides and neuronal toxicity are diminished in neuronal cells co-treated with LPS. a) The reduced cell viability effect of A β 42 on SH-SY5Y cells is significantly improved by co-treatment with LPS in a dose-dependent manner. b) Western blotting indicating A β 42 degradation in cells co-treated with or without LPS. c) The LPS-induced degradation of A β 42 peptides is suppressed by three endocytosis blockers: chlorpromazine (CPZ), methyl- β -cyclodextrin (M β CD), and dynasore. d) The rescue effect of LPS against A β 42 neuronal toxicity is blocked by endocytosis inhibitors. ***, $P < 0.001$.

LPS-A β 42 complex might restore the endocytic clearance of the A β 42 peptides. To test this, we set out to investigate whether the A β 42 levels and aggregation were both decreased. Western blot analysis of A β 42 peptides collected from the entire volume of the cell culture medium and total cell lysates suggested that the extracellular and intracellular proteins of A β 42 were markedly depleted upon co-treatment with LPS (Figure 3b, lanes 3–5). However, in the absence of cells, the A β 42 levels and oligomeric states were remained unchanged by the LPS (Figure 3b, lanes 2 and 6), suggesting that the formation of transient complex between LPS and A β 42 triggered a potent cellular response that resulted in the degradation of the A β 42 peptides. These data provide strong evidence to support a role for the non-equilibrium complex of A β 42-LPS in rescuing cells from death through promoting the clearance of A β 42 protofibrils.

To understand the underlying mechanisms, we first tested whether matrix metalloproteinases (MMPs) secreted from cultured cells might mediate the destruction of A β 42 peptides in the extracellular milieu. However, the treatment of neuronal cells with pan-MMP inhibitors showed no changes in A β 42 levels in cells co-treated with LPS (Figure S3). Accordingly, we turned our attention to the intracellular protein degradation pathways. We used pharmacological blockers to inhibit endocytosis mediated by the clathrin- and dynamin-dependent mechanisms. The results suggested that the blockade of the endocytic uptake of extracellular A β 42 effectively abolished the amyloid degradation (Figure 3c, lanes 4–6) as well as the pro-survival effect of LPS rescuing SH-SY5Y cells from the A β 42-induced apoptosis (Figure 3d). Since autophagy is also a known cellular mechanism for endolysosomal degradation,^[29] we inhibited autophagy with two typical inhibitors, methyladenine (3MA) and bafilomycin A1 (BA), and found very comparable effects to those observed for the endocytosis blockers (Figure S4). The results strongly suggest that an endolysosomal pathway mediates the degradation of A β 42 peptides in neuronal cells co-treated with LPS.

Furthermore, it is essential to clarify (i) why LPS can only bind with A β 42 but not with A β 40 (Figure S5) and (ii) how they bind together. To understand the binding preference, we speculated that A β 42 might be more prone to fast fibrillization that possibly constructs surfactant properties to increase the LPS-A β 42 interaction in comparison to the interaction with A β 40. As expected, A β 42 was found to still undergo rapid fibrillization in both the presence and absence of LPS (Figure S6), a phenomenon that was not observed for A β 40. Additionally, small-angle X-ray scattering analysis showed that A β 42 increased the critical aggregation concentration (CAC) of LPS from $3.49 \pm 0.052 \mu\text{g mL}^{-1}$ ^[30] to $15.91 \pm 0.13 \mu\text{g mL}^{-1}$ (Figure S7, left panel), whereas the CAC of LPS was found to not be obviously affected by the presence of A β 40 ($3.82 \pm 0.07 \mu\text{g mL}^{-1}$, Figure S7, right panel). The increased CAC of LPS by A β 42 suggested that a mutual interaction between LPS and soluble A β 42 protofibrils occurs to suppress the self-assembly process of LPS.

Prompted by the finding that A β 42 formed a complex with amphiphilic LPS, we speculated that the soluble A β 42 protofibrils might possess a specific groove that acts as a kinetic trap and enables the docking of amphiphilic LPS. To

test this possibility, we used a unique LPS sequester, an atomic sheetlike gold nanocluster (identified as SAuM) with a specific dock for the lipid A of the hydrophobic domain, that has been demonstrated in our previous work.^[30,31] Indeed, the binding efficiency of A β 42 protofibrils and LPS was found to be significantly decreased in the presence of the SAuM (Figure 4a, black line). Furthermore, we also showed that colistin, a cyclic peptide which is known to cap the hydrophilic domain (O-antigen) of LPS,^[32] exerted a similar effect in decreasing the binding efficiency of A β 42 protofibrils to LPS (Figure 4a, red line). Both results indicated that the A β 42 protofibrils possess LPS-specific amphiphilic grooves that allow docking with the hydrophobic and hydrophilic domains of LPS. Accordingly, we abolished the complex formation of LPS and A β 42 by blocking the lipid A or O-antigen binding sites in LPS with the above two inhibitors. The results clearly showed that the roles of LPS in promoting the clearance of A β 42 protofibrils (Figure 4b) or in attenuating the cytotoxicity of A β 42 were both compromised (Figure S8). We then integrated these results into a model of the structural-functional interaction between LPS and A β 42 in modulating the endocytic clearance of A β 42 in neuronal cells (Figure 4c).

In conclusion, we demonstrated that the specific groove of A β 42 protofibrils is important for complexing with amphiphilic LPS through a pattern of non-equilibrium behavior. We also speculate that LPS may actually act as a bait in attracting the A β 42 intermediates that sometimes deviate from their supposed thermodynamic self-assembly process. With respect

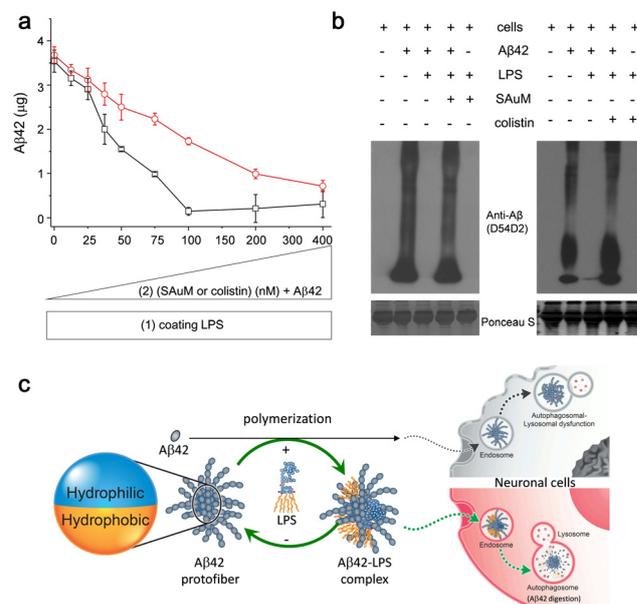


Figure 4. The amphiphilic groove of A β 42 protofibrils is required for the binding of LPS and the induction of neuronal clearance of A β 42. a) Two decay curves revealing that the complex formation of A β 42 protofibrils and LPS was impeded in the presence of antagonistic O-antigen (red circles, as colistin blocked the hydrophilic domain of LPS) or lipid A (black squares, as SAuM blocked the hydrophobic domain of LPS). b) LPS antagonists effectively suppressed the A β 42 degradation induced by LPS in cells. c) Pictorial illustration showing that LPS may induce an endo-lysosomal clearance of A β 42 in neuronal cells via the formation of a non-equilibrium complex with the A β 42 protofibrils.

to functionality, we provide evidence supporting the conclusion that such a transient supramolecule–supramolecule interaction can potently stimulate a strong cellular response toward autophagy-mediated protein degradation for A β 42 peptides in neuronal cells. Moreover, since the oscillation of the non-equilibrium state appears to be sustainably maintained, the extracellular A β 42 protofibrils were eventually diminished over a prolonged incubation time with the cells. Altered homeostasis between A β peptide production and clearance is defined as the pathological basis for the accumulated A β fibrils in AD brains. Since efforts aimed at blocking A β 42 production have not been successful, the current strategies of AD drug design have been shifted to target A β clearance and degradation. The current work is important because it provides a new strategy for structure-based drug design to discover therapeutics that promotes A β 42 clearance.

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Conflict of interest

The authors declare no conflict of interest.

Keywords: amyloid β -peptides · polysaccharides · proteins · non-equilibrium processes · supramolecular chemistry

- [1] A. Sorrenti, J. Leira-Iglesias, A. J. Markvoort, T. F. A. de Greef, T. M. Hermans, *Chem. Soc. Rev.* **2017**, *46*, 5476–5490.
- [2] S. Dhiman, A. Sarkar, S. J. George, *RSC Adv.* **2018**, *8*, 18913–18925.
- [3] C. Moberg, *Angew. Chem. Int. Ed.* **2020**, *59*, 2550–2553; *Angew. Chem.* **2020**, *132*, 2570–2573.
- [4] A. Pross, *What is life? How chemistry Becomes Biology*, 2nd ed., Oxford University Press, Oxford, **2016**.
- [5] A. Desai, T. J. Mitchison, *Annu. Rev. Cell Dev. Biol.* **1997**, *13*, 83–117.
- [6] E. N. Cline, M. A. Bicca, K. L. Viola, W. L. Klein, *J. Alzheimer's Dis.* **2018**, *64*, S567–S610.
- [7] A. K. Buell, *Biochem. J.* **2019**, *476*, 2677–2703.
- [8] J. Vaquer-Alicea, M. I. Diamond, *Annu. Rev. Biochem.* **2019**, *88*, 785–810.
- [9] N. Suzuki, T. T. Cheung, X. D. Cai, A. Odaka, L. Otvos, C. Eckman, T. E. Golde, S. G. Younkin, *Science* **1994**, *264*, 1336–1340.
- [10] B. L. Kagan, H. Jang, R. Capone, F. T. Arce, S. Ramachandran, R. Lal, R. Nussinov, *Mol. Pharm.* **2012**, *9*, 708–717.
- [11] P. Bergman, N. R. Roan, U. Romling, C. L. Bevens, J. Munch, *J. Intern. Med.* **2016**, *280*, 139–152.
- [12] M. L. Gosztyla, H. M. Brothers, S. R. Robinson, *J. Alzheimer's Dis.* **2018**, *62*, 1495–1506.
- [13] D. K. V. Kumar, S. H. Choi, K. J. Washicosky, W. A. Eimer, S. Tucker, J. Ghofrani, A. Lefkowitz, G. McColl, L. E. Goldstein, R. E. Tanzi, R. D. Moir, *Sci. Transl. Med.* **2016**, *8*, 340ra72.
- [14] M. Ahmed, J. Davis, D. Aucoin, T. Sato, S. Ahuja, S. Aimoto, J. I. Elliott, W. E. Van Nostrand, S. O. Smith, *Nat. Struct. Mol. Biol.* **2010**, *17*, 561–U556.
- [15] S. M. Butterfield, H. A. Lashuel, *Angew. Chem. Int. Ed.* **2010**, *49*, 5628–5654; *Angew. Chem.* **2010**, *122*, 5760–5788.
- [16] I. W. Hamley, *Chem. Rev.* **2012**, *112*, 5147–5192.
- [17] Z. Fu, D. Aucoin, J. Davis, W. E. Van Nostrand, S. Smith, *Biochemistry* **2015**, *54*, 4197–4207.
- [18] J. A. Luiken, P. G. Bolhuis, *J. Phys. Chem. B* **2015**, *119*, 12568–12579.
- [19] B. Morel, M. P. Carrasco, S. Jurado, C. Marco, F. Conejero-Lara, *Phys. Chem. Chem. Phys.* **2018**, *20*, 20597–20614.
- [20] X. Wang, P. J. Quinn, *Endotoxins: structure, function and recognition*, Vol. 53, Springer, Dordrecht, **2010**.
- [21] X. H. Zhan, B. Stamova, L. W. Jin, C. DeCarli, B. Phinney, F. R. Sharp, *Neurology* **2016**, *87*, 2324–2332.
- [22] K. M. Lucin, T. Wyss-Coray, *Neuron* **2009**, *64*, 110–122.
- [23] N. D. Younan, J. H. Viles, *Biochemistry* **2015**, *54*, 4297–4306.
- [24] K. Brandenburg, H. Mayer, M. H. J. Koch, J. Weckesser, E. T. Rietschel, U. Seydel, *Eur. J. Biochem.* **1993**, *218*, 555–563.
- [25] D. M. Walsh, B. P. Tseng, R. E. Rydel, M. B. Podlisny, D. J. Selkoe, *Biochemistry* **2000**, *39*, 10831–10839.
- [26] L. A. Welikovich, S. Do Carmo, Z. Magloczky, P. Szocsics, J. Loke, T. Freund, A. C. Cuello, *Acta Neuropathol.* **2018**, *136*, 901–917.
- [27] C. Yu, E. Nwabuisi-Heath, K. Laxton, M. J. Ladu, *Mol. Neurodegener.* **2010**, *5*, 19.
- [28] K. E. Marshall, D. M. Vadukul, K. Staras, L. C. Serpell, *Cell. Mol. Life Sci.* **2020**, <https://doi.org/10.1007/s00018-020-03464-4>.
- [29] N. Mizushima, T. Yoshimori, B. Levine, *Cell* **2010**, *140*, 313–326.
- [30] F. H. Liao, T. H. Wu, Y. T. Huang, W. J. Lin, C. J. Su, U. S. Jeng, S. C. Kuo, S. Y. Lin, *Nano Lett.* **2018**, *18*, 2864–2869.
- [31] F. H. Liao, T. H. Wu, C. N. Yao, S. C. Kuo, C. J. Su, U. S. Jeng, S. Y. Lin, *Angew. Chem. Int. Ed.* **2020**, *59*, 1430–1434; *Angew. Chem.* **2020**, *132*, 1446–1450.
- [32] P. Pristovšek, J. Kidrič, *J. Med. Chem.* **1999**, *42*, 4604–4613.

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