Amyloid-β fibrils assembled on ganglioside-enriched membranes contain both parallel β-sheets and turns

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Some protein and peptide aggregates, such as those of amyloid-β protein (Aβ), are neurotoxic and have been implicated in several neurodegenerative diseases. Aβ accumulates at nanoclusters enriched in neuronal lipids called gangliosides in the presynaptic neuronal membrane, and the resulting oligomeric and/or fibrous forms accelerate the development of Alzheimer’s disease. Although the presence of Aβ deposits at such nanoclusters is known, the mechanism of their assembly and the relation-ship between Aβ secondary structure and topography are still unclear. Here, we first confirmed by atomic force microscopy that Aβ140 fibrils can be obtained by incubating seed-free Aβ140 monomers with a membrane composed of sphingomyelin, cho-lesterol, and the ganglioside GM1. Using Fourier transform infrared (FTIR) reflection–absorption spectroscopy, we then found that these lipid-associated fibrils contained parallel β-sheets, whereas self-assembled Aβ140 molecules formed antiparallel β-sheets. We also found that the fibrils obtained at GM1-rich nanoclusters were generated from turn Aβ140. Our findings indicate that Aβ generally self-assembles into antiparallel β-structures but can also form protofibrils with parallel β-sheets by interacting with ganglioside-bound Aβ. We con-cluded that by promoting the formation of parallel β-sheets, highly ganglioside-enriched nanoclusters help accelerate the elongation of Aβ fibrils. These results advance our understanding of ganglioside-induced Aβ fibril formation in neuronal membranes and may help inform the development of additional therapies for Alzheimer’s disease.

Protein and peptide aggregates are neurotoxic and are thus implicated in neurodegenerative diseases (1–3). In particular, amyloid precursor protein is digested by secretases implicated in neurodegenerative diseases (1–3). In particular, atomic force microscopy (AFM) of a reconstituted lipid bilayer containing mouse synaptosomal lipids has suggested that Aβ-sensitive ganglioside nanoclusters promote Aβ40 assembly (29, 30). In addition, the chain length of ganglioside GD1b was found to influence Aβ42 assembly at the neuronal membrane in human pre-cuneus with amyloid deposits at such sites for the onset of Alzheimer’s disease. Aβ assembly at neuronal membranes (23, 24), in which a ganglioside-bound Aβ (GAβ) complex acts as an endog-enous seed, was first reported by Yanagisawa et al. (18) and was eventually demonstrated on ganglioside-containing liposomes using a thioflavin T assay, EM, and antibody assays (25, 26). The toxicity of the Aβ assembly resulting from the GAβ complex has been assessed using rat PC12 pheochromocytoma cells (27) and human neuroblastoma SH-SY5Y cells (28). These cells express gangliosides, including GM1, and nerve growth factor receptor–mediated neuronal cell death or cellular damage has been indicated. More recently, atomic force microscopy (AFM) of a reconstituted lipid bilayer containing mouse synaptosomal lipids has suggested that Aβ-sensitive ganglioside nanoclusters promote Aβ40 assembly (29, 30). In addition, the chain length of ganglioside GD1b was found to influence Aβ42 assembly at the neuronal membrane in human pre-cuneus with amyloids (31). Structural studies of Aβ polymerized at ganglioside-containing membranes are limited and contradictory. For example, Matsuzaki and Horikiri (32) found by CD that Aβ40 forms β-sheets at liposomes containing GM1, e.g. liposomes of sphingomyelin/cholesterol/GM1 (5:2:3). Similarly, FTIR attenuated total reflection spectroscopy indicated that Aβ40 forms antiparallel β-sheets at dry-cast films of egg yolk 1,α-phosphatidylcholine/GM1/Aβ40 (40:10:1) (32) or at liposomes of GM1/cho-

This article contains Figs. S1 and S2 and Table S1. To whom correspondence should be addressed. Tel.: 81-45-566-1771; E-mail: sato@bio.keio.ac.jp.

The abbreviations used are: Aβ, amyloid-β protein; AFM, atomic force microscopy; GAβ, ganglioside-bound Aβ; GlcCer, glucosylceramide; POPC, 1-palmitoyl-2-oleoyl-sn-glycero-3-phosphocholine; mN, millinewton.

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listerol/sphingomyelin (4:3:3) (33). On the other hand, NMR data collected by Utsumi et al. (34–36) suggest that Aβ₄₀ forms α-helices at membranes containing GM1 because of the hydrophobic environment. Recently, Hu et al. (37) also showed by Raman spectroscopy that α-helices and β-sheets of Aβ₄₀ were eventually observed with a planar lipid bilayer composed of GM1/sphingomyelin/cholesterol (5:55:40 and 20:40:40), although the topography of the fibrils was not investigated. Among these studies, spectroscopic evidence of ganglioside-induced fibrillar Aβ with parallel β-sheets has not yet been reported. In light of these results, a unifying model of Aβ assembly is needed to explain most of the structural data.

Previously, we found that Aβ-sensitive ganglioside nanoclusters in neuronal membranes induce a conformational change in Aβ₄₀ (29, 38). This and other characteristic properties of Aβ-sensitive ganglioside nanoclusters are mimicked by ganglioside-enriched planar membranes composed of ganglioside (GM1, GM2, GD1a, GD1b, or GT1b)/sphingomyelin/cholesterol (10:45:45) (30). These membranes are constructed by depositing a monolayer composed of ganglioside, sphingomyelin, and cholesterol with a lateral plasma membrane pressure (30 mN m⁻¹) on phospholipid-coated mica. This type of membrane is more stable than conventional liposomes (39) and supported lipid bilayers (37) with the same lipid composition, and AFM images can be obtained in a few days. In addition, various lipid compositions of the membrane are acceptable even if there are lipids incapable of forming liposomes.

Using FTIR reflection–absorption spectroscopy, we have now determined the secondary structure of Aβ₄₀ fibrils obtained in 15 min to 72 h on membranes containing GM1 at a 20% molar ratio (GM1/sphingomyelin/cholesterol, 20:40:40). The data indicate that the fibrils form turns and parallel β-sheets within 48 h. On the other hand, Aβ₄₀ assembled on membranes with 20% glucosylceramide (GlcCer), as well as self-assembled Aβ₄₀ forms antiparallel β-sheets. Solid-state NMR analyses by Tycko and co-workers (10, 40, 41) indicate that Aβ₄₀ predominantly forms cross-β-structures based on parallel β-sheets stabilized by intermolecular hydrogen bonds. Our results imply that by promoting the formation of parallel β-sheets, highly ganglioside-enriched nanoclusters also accelerate the elongation of Aβ fibrils.

**Results**

### Formation of Aβ fibrils on GM1-enriched membranes

As described previously (30), Aβ₄₀ fibrils were formed at ganglioside-enriched, planar, bilayer membranes, which were prepared by depositing 20:40:40 monolayers of GM1, sphingomyelin, and cholesterol onto 1-palmitoyl-2-oleoyl-sn-glycéro-3-phosphocholine (POPC)–coated mica (GM1-enriched membrane (Fig. 1A)). This composition mimics that of GM1-enriched microdomains and has frequently been used for ganglioside-induced Aβ assembly (39, 42). To confirm fibril formation, membranes were imaged by AFM in water after incubation with Aβ₄₀. GM1-enriched microdomains and Aβ assemblies were then visualized as areas higher than 4 nm on binarized AFM images. GM1-enriched microdomains before incubation with Aβ₄₀ had a diameter of 30–300 nm (>700 domains in a 5 μm × 5 μm area), and the apparent size of the domains increased after incubation with Aβ₄₀ for 15 min (60–500 nm, 280 domains) (Fig. S1). This topological change suggests that Aβ molecules are deposited on the GM1-enriched membrane to yield an Aβ layer (30). After 48 h, over a dozen Aβ₄₀ fibrils >1-μm long were clearly observed in a 5 × 5-μm area. These Aβ fibrils accumulated on round-shaped GM1-enriched microdomains, as reported previously (30).

To measure FTIR in air, after the formation of Aβ₄₀ fibrils, membranes were dried overnight and imaged in air (Fig. 1B, upper panel). The binarized AFM images of GM1-enriched membranes after incubation with Aβ₄₀ for 15 min, 48 h, and 72 h are shown in Fig. 1B (lower panel) at a height threshold of 2.0–3.0 nm. Drying slightly altered the shape of GM1-enriched microdomains, but Aβ₄₀ fibrils were still identifiable, and more Aβ₄₀ fibrils were observed after 72 h than after 48 h. On the other hand, one long fibril (>3 μm) and several short fibrils were observed on GlcCer-enriched membranes after 72 h (Fig. 2). These results indicate that GM1 generates and elongates Aβ fibrils more effectively than GlcCer.
**Immobilization of GM1-enriched membranes on gold-coated glass**

FTIR reflection–absorption spectra were collected to investigate the formation of lipid bilayers on gold-coated glass. Seven characteristic peaks in a monolayer of POPC were assigned according to the literature to \( \text{CH}_3, \text{CH}_2, \text{PO}_2^- \), and ester C–O stretching vibration (Fig. 3A and Table 1) (43, 44). A strong peak corresponding to \( \text{C} = \text{O} \) stretching vibration (\( \nu(\text{C} = \text{O}) \)) of 1742 cm\(^{-1}\) was also observed. On the other hand, amide I (1660 cm\(^{-1}\)) and amide II (1547 cm\(^{-1}\)) were observed in monolayers containing GM1 or GlcCer at a 20% molar ratio, as both lipids contain ceramide (44, 45). Two peaks, at 3342 and 1742 cm\(^{-1}\) (1660 cm\(^{-1}\)), in these membranes were assigned to N–H stretching and –CH\(_3\) scissoring vibration, respectively. The spectrum of a lipid bilayer composed of POPC as the first layer and GM1/sphingomyelin/cholesterol as the second layer is simply a superposition of the spectrum of each, clearly implying that a lipid bilayer with GM1 was formed on gold-coated glass.

**A\(\beta\) deposition on GM1- and GlcCer-enriched membranes**

To investigate the interaction between A\(\beta\) and a GM1-enriched membrane, FTIR reflection–absorption spectra were collected after 15 min, 24 h, 48 h, and 72 h. Incubation with A\(\beta_{40}\) shifted amide I and II peaks to 1663 and 1541 cm\(^{-1}\), respectively (Fig. 3B, upper panel, and Table S1). In addition, the height of these peaks significantly increased with time, as plotted in Fig. 4 along with peak shifts, indicating A\(\beta_{40}\) accumulation. The absorbance continued to increase even at 72 h, although peak shifts had nearly stabilized by that point.

The amide I and II bands in GlcCer-enriched membranes were at positions similar to those in GM1-enriched membranes (Fig. 3B, lower panel, and Fig. 4), implying comparable molecular structures in both membranes in light of the surface selection rule of reflection–absorption spectroscopy. In addition, the peaks were of similar relative intensity at 15 min and 72 h, implying comparable A\(\beta_{40}\) accumulation on both membranes, despite drastic differences in their ability to form fibrils (Figs. 1B and 2).

**Secondary structure of A\(\beta\) fibrils based on second-derivative reflection–absorption spectra**

The secondary structure of A\(\beta_{40}\) fibrils deposited on GM1-enriched membranes was determined from the reflection–absorption spectra of amide I (1700–1600 cm\(^{-1}\)) region and the corresponding second-derivative reflection spectra (46–49). The antiparallel \(\beta\)-sheet (pair of peaks, 1612–1640 and 1655–1675 cm\(^{-1}\)) and turn structures (1655–1675 cm\(^{-1}\) and/or 1680–1696 cm\(^{-1}\)) were determined on the basis of previous reports (50, 51).

The strongest peak of the amide I region from 15 min to 72 h shifted from 1676 to 1663 cm\(^{-1}\) observed in the raw spectrum of GM1-enriched membranes (Fig. 3B, upper panel, Fig. S2, and Table S1). The second-derivative reflection spectra indicated that two peaks (1661–1662 and 1691–1695 cm\(^{-1}\))
corresponding to the turns were appeared after 48 and 72 h (Fig. 5A, Table 1, and Fig. S2).

On the other hand, a strong peak at 1668 cm\(^{-1}\) with two shoulders was observed in the raw spectrum of GlcCer-enriched membranes after 72 h (Figs. S3B, lower panel, and S5B). The second-derivative spectra indicate that the two shoulders segregate into two peaks at 1631 and 1697 cm\(^{-1}\). Because the peak of GlcCer-enriched membranes after 72 h at 1631 cm\(^{-1}\) was distinguishable from that of GM1-enriched membranes at 1635 cm\(^{-1}\), these peaks at 1631 and 1635 cm\(^{-1}\) are assigned to antiparallel and parallel \(\beta\)-sheets, respectively (51). Two peaks, at 1631 and 1676 cm\(^{-1}\), of GlcCer-enriched membranes were assigned to a pair of peaks for antiparallel \(\beta\)-sheets. From these results, we concluded that GM1-enriched membranes induced \(\alpha_{40}\) fibrils with parallel \(\beta\)-sheets, in contrast to GlcCer-enriched membranes with antiparallel \(\beta\)-sheets (Fig. 5C and Table 2).

**Secondary structure of self-assembled \(\alpha_{40}\) based on attenuated total reflection spectra**

Seed-free \(\alpha_{40}\) was self-assembled for 15 min and 48 h (6), dropped on a suitable plate, and dried under nitrogen gas for 50 min. Three peaks at around 1630, 1670, and 1697 cm\(^{-1}\) in the attenuated total reflection spectrum at 15 min were again observed in the spectrum at 48 h (Fig. 6A). The pair of peaks at 1630 and 1666–1670 cm\(^{-1}\) in the second-derivative spectra are characteristic of antiparallel \(\beta\)-sheets because the major component at 1612–1640 cm\(^{-1}\) was accompanied by a minor component at 1670–1690 cm\(^{-1}\) (Table 2) (50). The peak at 1697 cm\(^{-1}\) is attributed to a turn structure.

**Structural features of residual \(\alpha_{40}\) in the supernatant incubated with GM1-enriched membranes**

Although \(\alpha_{40}\) fibrils formed on GM1-enriched membranes contain turns and parallel \(\beta\)-sheets (Fig. 5A), raw and second-derivative attenuated total reflection spectra (Fig. 6B) indicate that residual \(\alpha_{40}\) in the supernatant is structurally similar to self-assembled \(\alpha_{40}\) with major and minor components at 1630 and 1679 cm\(^{-1}\) corresponding to antiparallel \(\beta\)-sheets (Table 2). An AFM image of the residual \(\alpha_{40}\) in the supernatant after a 48-h incubation with a GM1-enriched membrane supports the FTIR spectra, short fibrils, 150 nm or less in length, were observed (Fig. 6C).

**Discussion**

The objective of this study was to investigate the secondary structure of \(\alpha_{40}\) assembled on GM1-enriched membranes with a view to clarify the mechanism of assembly. Often, \(\alpha_{40}\) assembly on ganglioside-containing membranes is investigated in solution by fluorescence thioflavin T assay and CD, because such membranes are often prepared as liposomes. However, seemingly contradictory results have been reported, preventing the formulation of a unified mechanism of ganglioside-induced \(\alpha_{40}\) assembly. For example, Fukunaga et al. (33) reported that GM1-induced \(\alpha_{40}\) contains antiparallel \(\beta\)-sheets based on FTIR but did not observe the \(\alpha\)-helices detected on NMR (36). Recently, \(\alpha\)-helices and \(\beta\)-sheets were detected by Raman spectroscopy of \(\alpha_{40}\) deposited for 24 h on supported lipid bilayers composed of GM1/sphingomyelin/cholesterol (5:55:40 and 20:40:40) (37). There is no evidence of ganglioside-induced fibrillar \(\alpha_{40}\) with parallel \(\beta\)-sheets that implies cross-\(\beta\) structures (10, 40, 41). Most unfortunately, however, the topography of \(\alpha_{40}\) assemblies are not always investigated when the secondary structure of \(\alpha_{40}\) assemblies is assessed.

We attempted to image \(\alpha_{40}\) assemblies directly with time via AFM using membranes containing GM1/sphingomyelin/...
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cholesterol (20:40:40) on mica (29, 30). The images confirmed that Aβ fibrils were formed on such membranes after 48–72 h but not on membranes containing GlcCer (Figs. 1B and 2). We have now determined from the secondary structures of Aβ assemblies from second-derivative reflection–absorption spectra (Fig. 5A) that Aβ fibrils deposited on GM1-enriched membranes for 48–72 h consist of turns (1661–1662 cm\(^{-1}\) and 1691–1695 cm\(^{-1}\)) and parallel β-sheets (1635–1637 cm\(^{-1}\)). The parallel β-sheets are clearly distinguishable from antiparallel β-sheets, typified by two characteristic peaks at 1631 and 1676 cm\(^{-1}\), formed on GlcCer-enriched membranes (Fig. 5B). In this case, the difference in secondary structure correlates with the AFM data (Figs. 1 and 2).

In light of our results, a model of ganglioside-induced Aβ assembly was proposed (Fig. 7) in which monomeric Aβ forms an initial layer not only on GM1-enriched nanoclusters but also on sphingomyelin/cholesterol area (step a) (29, 30). NMR (36) and Raman spectroscopic studies (37) indicated that Aβ at ganglioside nanoclusters then forms helices (step b, GAB formation); however, a prominent helical peak at 1650–1657 cm\(^{-1}\) was not observed (see Fig. 5A). Subsequently, Aβ molecules in contact with ganglioside-bound helical Aβ formed turn and antiparallel β-sheets and transited to parallel β-sheets (Fig. 7, steps c and d) (14). Previous FTIR findings of GM1-induced Aβ with antiparallel β-sheets, reported by Matsuzaki and Horikiri (32) and Fukunaga et al. (33), seem to resemble the present oligomeric Aβ assembly (Fig. 7, steps c and f). Finally, protofibrils with parallel β-sheets were formed and extended (Fig. 7, step e) as observed on reflection–absorption spectra at 48–72 h (Fig. 5A). The parallel β-sheet structure is stabilized by intermolecular interactions between Aβ molecules and β-sheet side chains to form cross-β units (9). Multiple molecular dynamics simulations support the binding of Aβ to ganglioside clusters through the combination of a CH–π/π interaction, a Lys\(^{28}\)–Neu5Ac interaction, and hydrophobic interactions at the C terminus and also support the involvement of two or three Aβ molecules of the GAβ complex in the formation of the parallel β-sheet (52).

Table 2

| Assignment | 20% GM1 on POPC | 20% GlcCer on POPC | Self-assembled |
|------------|-----------------|-------------------|----------------|
|            | 15 min Deposited | 24 h Oligomer/fibrils | 48 h (ATR) Supernatant | 15 min (ATR) Aggregates | 48 h (ATR) Aggregates |
| Parallel β | 1637            | 1635              | 1630/1679        | 1630/1666               | 1630/1670               |
| Antiparallel β | 1664\(^{a}\) | 1668\(^{a}\)/1691 | 1661/1691        | 1666\(^{a}\)            | 1667\(^{a}\)/1697        | 1697            | 1697            |
| Turn       | 1664\(^{a}\) | 1668\(^{a}\)/1691 | 1661/1691        | 1666\(^{a}\)            | 1667\(^{a}\)/1697        | 1697            | 1697            |

\(^{a}\) Weak peaks.

Figure 6. Formation of Aβ assemblies in solution. A and B, raw (blue lines) and second-derivative (red lines) attenuated total reflection amide I spectra of self-assembled Aβ\(_{40}\) (A) and residual Aβ\(_{40}\) in the supernatant of solution on GM1-enriched membranes. GM1/sphingomyelin/cholesterol (20:40:40) membrane was incubated with 10 μM Aβ\(_{40}\) at 37 °C for 48 h, and thereafter the supernatant was dropped on a mica plate for 15 min. After washing, Aβ\(_{40}\) deposited on the mica was imaged via AFM in water.
Based on attenuated total reflection spectra, the secondary structure of residual $\alpha\beta$ in the supernatant of the solution on GM1-enriched membranes is similar to that of $\alpha\beta$ deposited on GlcCer-rich membranes, with a major (1631 cm$^{-1}$) and a minor peak (1676 cm$^{-1}$) attributable to antiparallel $\beta$-sheets (Fig. 6B), which were also detected in $\alpha\beta$ self-assembled for 48 h (Fig. 6A). Indeed, most of the residual $\alpha\beta$ molecules in the supernatant may not interact at all with the GM1-enriched membrane and therefore will self-assemble in the same way as seed-free $\alpha\beta$ (Fig. 6C). $\alpha\beta$-oligomerized into antiparallel $\beta$-sheets (Fig. 7, step g), as described previously by Stroud et al. (13) and Fu et al. (14) and confirmed by reflection-absorption spectroscopy of self-assembled $\alpha\beta$ and residual $\alpha\beta$ in the supernatant of the solution on GM1-enriched membranes after 48 h (Fig. 6). That $\alpha\beta$ deposited on GlcCer/sphingomyelin/cholesterol also formed antiparallel $\beta$-sheets (Fig. 5B), confirming that a GM1-enriched nanocluster is required to form fibrils with parallel $\beta$-sheets.

We noted that a mixture of GM1/sphingomyelin/cholesterol (20:40:40) formed a lipid bilayer with POPC on gold-coated glass as well as on mica (Fig. 1A) (30), with characteristic peaks of POPC (2ν(C=O), 1742 cm$^{-1}$) and GM1 (amide I and II, 1660 and 1547 cm$^{-1}$, respectively) observable on reflection-absorption spectra (Fig. 3A and Table 1). The transfer ratio of the 20% GM1 monolayer onto the POPC-coated slide was almost 1.0 (see “Experimental Procedures”), implying the formation of a suitable bilayer of 20% GM1 and POPC. The intensity (absorbance) of amides I and II in GM1- and GlcCer-containing membranes increased with time (Fig. 3), suggesting an accumulation of $\alpha\beta$. This result also suggests a comparable accumulation of an $\alpha\beta$ layer on both membranes, although AFM indicated that $\alpha\beta$ fibrils were selectively deposited on GM1-enriched membranes only (Figs. 1B and 2), as reported previously (30).

In conclusion, the data indicate that $\alpha\beta$ generally self-assembles into antiparallel $\beta$-structures but is competent to form protofibrils with parallel $\beta$-sheets, in this case by interaction with GAβ. This model is based on data from AFM and FTIR of a ganglioside-enriched planar membrane. The $\alpha\beta$ topography obtained by AFM was eventually linked to secondary structures obtained by FTIR. In addition, an initial $\alpha\beta$ layer was also detected by both methods, suggesting that the formation of this layer may explain the seemingly contradictory data in the literature. Our data also highlight the growing significance of molecular dynamics simulation in investigating the interaction between $\alpha\beta$ and neural membranes. Finally, these data advance our understanding of ganglioside-induced $\alpha\beta$ fibril formation on neuronal membranes, which may accelerate the development of novel therapies against Alzheimer’s disease.

**Experimental procedures**

**Lipids**

Monosialoganglioside GM1 from bovine brain and GlcCer from human (Gaucher disease) spleen were purchased from Sigma or Wako Pure Chemical Industries, Ltd. (Osaka, Japan). Sphingomyelin from bovine brain and synthetic POPC were from Matreya LLC (State College, PA) and Sigma, respectively.

**Preparation of seed-free soluble $\alpha\beta$**

Seed-free soluble $\alpha\beta$ was prepared as described previously (29, 30). In brief, synthetic $\alpha\beta$ (human, 1–40; code 4379-v, Peptide Institute Inc., Osaka, Japan) was dissolved in ice-cold 0.02% ammonia and ultracentrifuged at 560,000 × g for 3 h at 4 °C to remove undissolved peptide aggregates. The seed-free fraction (40–110 μM) was stored in aliquots at −80 °C until used. Prior to use, aliquots were diluted in Dulbecco’s PBS(−), pH 7.4 (Nissui Pharmaceutical Co., Ltd., Tokyo, Japan).

**Preparation of GM1-enriched membranes and AFM**

GM1-enriched lipid bilayers were prepared on mica as described previously (29, 30). Briefly, a POPC monolayer was prepared at 25 °C at the air–water interface of a Langmuir–Blodgett trough (FSD-220, USI Corp., Fukuoka, Japan), with...
water as the subphase, and deposited horizontally on freshly cut 1 × 1-cm mica at a surface pressure of 35 mN m⁻¹ (Fig. 1A, POPC-coated mica). To form the bilayer, a second monolayer consisting of GM1/sphingomyelin/cholesterol (20:40:40, molar ratio) at a surface pressure of 30 mN m⁻¹ was loaded horizontally onto POPC-coated mica by deposition.

The GM1-enriched membrane was incubated with 10 μM seed-free soluble Aβ40 in PBS for 15 min to 72 h at 37 °C (30, 31). After washing three times with PBS, the membrane was imaged at 25 °C in water using an SPM-9600 atomic force microscope (Shimadzu Corp., Kyoto, Japan) and a 38-μm soft cantilever (BL-AC40TS-C2, Olympus, Tokyo, Japan) with integrated pyramidal silicon nitride tips with spring constant 0.1 N m⁻¹. Multiple topographic images (2 × 2 μm, n = 3) were acquired in dynamic mode at 1–2 Hz, and representative images were used in further analyses.

To estimate the Aβ-coated areas, AFM images were binarized based on height, and pixels were counted using the GNU Image Manipulation Program. In particular, areas higher than around 4 nm on a binarized image were considered Aβ-coated layers. Fibril length and domain size were measured in ImageJ (National Institutes of Health, Bethesda, MD) using a line drawn along a fibril. We estimated the lengths of the long and short axes of an Aβ assembly and defined them as fibrils when the long/short axis aspect ratio was >3.

**Immobilization of GM1-enriched membranes onto gold-coated glass**

Glass slides (40 × 20 × 1.1 mm) coated with an evaporated gold layer 300 nm thick and a stabilizing chromium layer 50 nm thick were purchased from Geomatec (Yokohama, Japan) and gold layer 300 nm thick and a stabilizing chromium layer 50 nm thick. Finally, the GM1-enriched lipid bilayer was incubated from changes in the lipid area and the area of the slide that images were used in further analyses.

Attenuated total reflection spectroscopy

Attenuated total reflection spectra were collected as described previously (53). Briefly, 84 μM seed-free soluble Aβ40 in MilliQ water was incubated at 25 °C for 15 min or 48 h to induce self-assembly. About 15 μl of the resulting solution was dropped on a germanium plate and dried under nitrogen gas for 50 min. Spectra were collected using a single-reflection accessory (Spectra-Tech Foundation Performer, Thermo Fisher Scientific) and a germanium prism, with number of accumulations 1000. To compare these with the reflection–absorption spectra, attenuated total reflection spectra were transformed into absorbance (α) spectra according to α = 4πk/λ, where k and λ are the imaginary parts of the complex refractive index (1.5) and the wavelength, respectively.

**Second-derivative analysis of FTIR spectra**

FTIR spectra were analyzed in OMNIC, version 7.3. The second derivative (54) of each spectrum was calculated by the Savitzky–Golay method (55).

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