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1 Early-life stress facilitates the development of Alzheimer's disease pathology via angiopathy

2 Running title: Early-life stress-associated Alzheimer's disease

3

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4

5 Keyword: Alzheimer's disease, angiopathy, immune system, early life stress, amyloid-beta,

6 cognitive function

7

3

1 Abstract

2 Background: Alzheimer's disease (AD), a progressive neurodegenerative disorder, is a serious
3 social problem. Recently, several early-life factors have been associated with an increased risk
4 of a clinical diagnosis of AD.

5 Methods: We investigated the involvement of early-life stress in AD pathogenesis using
6 heterozygous the amyloid precursor protein (APP) mutant mice ($App^{NL-G-F/wt}$) and wild-type
7 ($App^{wt/wt}$) mice. Maternal separation was used as an animal paradigm for early-life stress. Object
8 location and fear conditioning tests were performed to measure cognitive functions, in addition
9 to biochemical tests. Immunohistochemical analyses were performed after the behavioral tests.

10 Results: We found that maternal-separated $App^{wt/wt}$ mice showed narrowing of vessels and
11 decreased pericyte coverage of capillaries in prefrontal cortex, while maternal-separated App^{NL-}
12 $G-F/wt$ mice additionally showed impairment of cognitive function, and earlier formation of A β
13 plaques and disruption of the blood–brain barrier. Severe activation of microglia was detected in
14 the maternal-separated $App^{NL-G-F/wt}$ mice and maternal-separated $App^{wt/wt}$ mice. At the early stage,
15 morphological changes and inflammatory responses were observed in the microglia of the

4

- 1 maternal-separated *App*^{NL-G-F/wt} mice and maternal-separated *App*^{w^t/w^t} mice, as well as
- 2 morphological changes in the microglia of the non-maternal-separated *App*^{NL-G-F/wt} mice.
- 3 Conclusions: Microglia activation induced by maternal separation in combination with the APP
- 4 mutation may impair the vascular system, leading to AD progression. These findings therefore
- 5 suggest that maternal separation causes early induction of AD pathology via angiopathy.

1 Introduction

2 The development of abnormal physiology in association with increased age is observed in the
3 progression of many disorders. Because daily habits and environmental stresses such as alcohol
4 intake and diet have the major influence on life-span, rather than genetic factors (Christensen et
5 al., 2006), acceleration of the aging process by such environmental factors is thought to increase
6 both the risk of developing a disease and mortality from a disease. Alzheimer's disease (AD) is
7 one such disease that is thought to be influenced by environmental factors; it is characterized by
8 progressive impairment of cognitive functions such as memory. It is considered that the
9 acquired predisposition to AD increases with age, and that trauma during adulthood increases
10 the likelihood of developing AD. Moreover, recent epidemiological research has suggested that
11 stressful environmental factors during the developmental period, including mistreatment,
12 neglect, and the loss of a parent, can also increase the likelihood of developing AD (Norton et
13 al., 2011; Radford et al., 2017; Seifan et al., 2015). However, as the research behind these
14 findings is generally the result of retrospective cohort studies, it is unclear whether early-life
15 stress affects the onset and progression of AD.

16 Recently, vascular impairment, a postmortem pathological finding in the brain, was reported

1 to occur before the onset of AD. Therefore, vascular impairment is considered as an initial
2 symptom of AD (Iadecola, 2017; Montagne et al., 2016; Wen et al., 2004). The neurovascular
3 unit (NVU) consists of neurons, endothelial cells, pericytes, microglia, and astrocytes. As the
4 NVU is important for the extrusion of amyloid beta, an impairment in extraction can cause
5 sporadic AD and cerebral amyloid angiopathy (Mawuenyega et al., 2010). In particular, the
6 capillary vessels in the brain play an important role in the extrusion of amyloid beta.
7 Dysfunction of the blood–brain barrier (BBB) and a decrease in amyloid beta clearance through
8 the cerebral vascular system is observed in AD (Zlokovic, 2011). Degeneration of cells making
9 up the BBB, such as pericytes, causes dysfunction of the BBB (Sweeney et al., 2016). However,
10 it remains unclear whether early-life stress results in dysfunction of the vascular system.

11 In this study, we investigated whether maternal separation, which can be used as an early-life
12 stress in rodents, facilitates the development of AD, with the purpose of understanding the
13 causal association between the development of AD and early-life stress. After finding
14 indications that maternal separation affects the development of AD, we next investigated the
15 changes induced by maternal separation, paying particular attention to capillary vessel
16 morphology in heterozygous the amyloid precursor protein (APP) mutant mice ($App^{NL-G-F/wt}$) and

7

1 wild-type (*App*^{wt/wt}) mice. As a result, we established that maternal separation facilitates the
2 development of AD. Furthermore, we found that impairment of capillaries, probably caused by
3 abnormalities in microglia, leads to dysfunction of the BBB, thereby facilitating the
4 development of clinical AD.

5

8

1 Materials and methods

2 All experimental procedures were approved by the Animal Experimentation Ethics Committee
3 of the Tokyo Metropolitan Institute of Medical Science (49040).

4

5 *Animals*

6 The original lines of App mutant mice ($App^{NL-G-F/NL-G-F}$) were established as a C57BL/6J
7 congenic line (a genetic background strain) by repeated back-crosses, as described previously,
8 and were obtained from RIKEN Center for Brain Science (Wako, Japan)(Saito et al., 2014). All
9 experiments were performed with $App^{NL-G-F/wt}$ and $App^{wt/wt}$ that were born from $App^{NL-G-F/wt}$ mice
10 x $App^{NL-G-F/wt}$ mice or $App^{NL-G-F/wt}$ mice x $App^{wt/wt}$ mice. $App^{NL-G-F/wt}$ mice with slow onset of
11 pathology were used because it was important to investigate the process of decline in function.
12 All mice were maintained under a 12:12 h light/dark cycle (lights on at 8:00 AM). All efforts
13 were made to minimize the number of animals used and their suffering. All experiments, except
14 for maternal separation, were performed in adolescence (postnatal day 45 (P 45)) or adulthood
15 (P90–P120).

16

1 *Maternal separation*

2 Maternal separation is usually used as a form of early-life stress in mice. Pups were separated
3 from their mother for 3 hours (AM 9:00-12:00) daily from P2 to P15. The separated pups were
4 taken to a different room and placed on a heat pat to maintain their body temperature. The
5 control mice had no experience of maternal separation. The body weights were not significantly
6 different between the non-maternal-separated and maternal-separated mice at several stages of
7 development.

8

9 *Behavioral assessment*

10 Male mice were allowed to habituate in the behavioral room for at least 1 hour before
11 commencement of the behavioral test. Mice showing outlier behavior > 2 standard deviations
12 from the mean or < 2 standard deviations from the mean were not analyzed by the behavioral
13 test.

14 The object location test consisted of three phases. All phases were performed under a light
15 intensity of 10 lux. On day 1, the animals were placed in an empty square box (50 cm x 50 cm)
16 for 10 minutes. On day 2, the animals were placed in the same box for 10 minutes with two

1 same identical unfamiliar objects. On day 3, the animals were placed in the same box, with one
2 of the two items being displaced to a novel location in the area. The discrimination index was
3 calculated as the ratio of the time spent exploring the novel place to that spent exploring the
4 familiar place: discrimination index = (novel place exploration time – familiar place exploration
5 time)/(novel place exploration time + familiar place exploration time). On each day, the
6 exploration time in each place was calculated automatically using a DVTrack video tracking
7 system (Muromachi, Tokyo, Japan).

8 The fear conditioning test consisted of three phases. On day 1, the animals were placed in a
9 triangular box for 5 minutes under a light intensity of 30 lux. On day 2, the animals were placed
10 into a square box with a stainless-steel grid floor and were allowed to explore the box freely for
11 2 minutes. Subsequently, a tone, which served as the conditioned stimulus (CS), was presented
12 for 30 sec. During the last 1 sec of CS presentation, a 0.75 mA electric shock was applied,
13 which served as the un-conditioning stimulus (US). Two more CS-US pairings were presented
14 with a 1 minutes inter-stimulus interval under a light intensity of 100 lux. On day 3, the animals
15 were placed into the same square box as used on day2 for 5 minutes under a light intensity of
16 100 lux. On day 4, the animals were placed in the same triangular box as used on day 1 for 10

11

1 minutes and were allowed to explore the box freely for 2 minutes. Subsequently, CS was
2 presented for 30 sec. Seven more CS were presented with a 1 minutes inter-stimulus interval
3 under a light intensity of 100 lux. In each test, the percentage of freezing time and distance
4 travelled (cm) were calculated automatically using ImageFZ software (O'Hara & Co., Tokyo,
5 Japan). Freezing was taken to indicate that the mice remembered the box (context) or tone (cue)
6 in which they were exposed to the electrical shock.

7

8 Immunohistochemistry

9 At P45 or P120, male and female mice were anaesthetized with isoflurane and then
10 sequentially perfused through the heart with PBS followed by 4% paraformaldehyde (PFA) in
11 PBS. Their brains were post-fixed, cryoprotected in 20% sucrose at 4°C, and embedded in OCT
12 compound and frozen. Coronal sections of mouse brain were then cut using a cryostat (20 µm).
13 Free-floating sections containing PFC were activated by HistVT one (Nakalai., Kyoto, Japan) at
14 70°C for 20 minutes, incubated in 0.4% Block-ase (DS Pharma Biomedical., Osaka, Japan) in
15 PBS at room temperature for 20 minutes, and then incubated with primary antibody
16 (Supplemental figure 1). All primary antibodies were diluted 1:500 in PBS containing 0.3%

12

1 Triton X-100. The sections were then incubated with secondary antibody (Jackson., Maine,
2 United State) (diluted 1:500 in PBS containing 0.3% Triton X-100) for 2 hours at room
3 temperature. All sections types were incubated with DAPI (Nacalai Tesque., Kyoto, Japan) for
4 nuclear staining. Fluorescence micrographs of the sections subjected to immunostaining
5 procedures were captured and digitized with a FluoView® FV3000 confocal laser scanning
6 microscope (Olympus., Tokyo, Japan). With the aim of quantifying the morphological change of
7 microglia, skeleton analysis and particle analysis were performed by ImageJ(Fernández-Arjona
8 et al., 2017; Xu et al., 2018).

9

10 Luminex analysis assay

11 Following the sacrifice of male and female mice, mouse PFC fractions were dissociated with
12 a tissue homogenizer in 100 µl of 0.5 × cell lysis buffer 2 diluted in PBS (R&D Systems.,
13 Minnesota, United State). Tissues were lysed at room temperature for 30 minutes, and debris
14 was then removed by centrifugation. Magnetic luminex assays were performed using a Mouse
15 Premixed Multi-Analysis Kit (R&D Systems., Minnesota, United State). Assay procedures were
16 carried out following manufacturer's instructions. Microplates were run on a Bio-Plex 200

1 system (Bio-Rad., California, United State).

2

3 Corticosterone assay

4 Male and female mice were subjected to a novel cage stress (Ratray et al., 2009), in which

5 each mouse was placed in isolation in a new cage with a smooth floor for 30 minutes at P45.

6 Blood samples were taken before stress and immediately after stress, and centrifuged at 3,300

7 rpm to separate serum. Corticosterone concentrations were measured using an enzyme-linked

8 immunosorbent assay (ELISA) (AssayPro., Missouri, United State).

9

10 Statistical data analysis

11 Group data are presented as the mean \pm SEM. The statistical significance of between-group

12 differences was assessed by the Tukey–Kramer test, Dunnett’s test, or the paired t-test, using

13 JMP software (SUS Institute., North Carolina, United State). The interaction between area and

14 distance was assessed using the polynomial regression model. The statistical significance of

15 differences between groups and over time was assessed by two-way repeated ANOVA followed

16 by a *post-hoc* Holm’s test, using R software.

1 Results

2 *Experimental design*

3 Figure 1A shows the experimental schedule. To investigate whether stress in the early
4 developmental stage is involved in the onset and/or progression of AD, we performed maternal
5 separation for 3 hours daily during the period from P2 to P15 in $App^{wt/wt}$ and $App^{NL-G-F/wt}$ mice.
6 The $App^{NL-G-F/wt}$ mice show an AD-like phenotype in the late-aged stage (Saito et al., 2014). We
7 then performed behavioral tests and tissue sampling around P45 (adolescence) and P120 (adult).

8

9 *Maternal separation induced impairment of cognitive function and facilitated formation of*
10 *amyloid beta plaque in the adult $App^{NL-G-F/wt}$ mice*

11 Figure 1B illustrates the effect of maternal separation on object location test which evaluate
12 spatial memory (Denninger et al., 2018). Object location memory in the $App^{wt/wt}$ mice was not
13 altered by maternal separation, as indicated by the fact that the discrimination index was not
14 changed by maternal separation. By contrast, object location memory in the $App^{NL-G-F/wt}$ mice
15 was impaired by maternal separation, as indicated by a discrimination index close to zero. The
16 non-maternal-separated $App^{NL-G-F/wt}$ mice showed intact object location memory. These results

1 indicate that the combination of APP mutation and maternal separation influence object location
2 memory. Figure 1C indicates the effect of maternal separation on contextual and cued fear
3 memory. Contextual fear memory in the $App^{wt/wt}$ mice was not altered by maternal separation,
4 while that in the $App^{NL-G-F/wt}$ mice was impaired by maternal separation, as indicated by the
5 lower freezing time in the maternal-separated $App^{NL-G-F/wt}$ mice in comparison with the non-
6 maternal-separated $App^{NL-G-F/wt}$ and non-maternal separated $App^{wt/wt}$ mice. By contrast, tone-cued
7 fear memory was not impaired in any of the four groups, indicating that contextual memory,
8 which is related to prefrontal cortex–hippocampus signaling and is known to be necessary for
9 the encoding of spatial cues, was impaired (Kitamura et al., 2017). Figure 1D indicates the
10 effect of maternal separation on the formation of amyloid beta plaque. The formation of
11 amyloid beta plaque in the $App^{NL-G-F/wt}$ mice was facilitated in mice that underwent maternal
12 separation than in non-maternal-separated $App^{NL-G-F/wt}$ mice.

13

14 *Maternal separation induces vascular system impairment in adult mice*

15 Phenotypes showing decreased pericyte coverage, a narrowing of capillary diameter, and
16 impairment of the BBB are frequently observed in the postmortem brain of AD patients

1 (Halliday et al., 2016). Figure 2 illustrates the effect of maternal separation on the vascular
2 system. The maternal-separated *App*^{wt/wt} and maternal-separated *App*^{NL-G-F/wt} mice showed lower
3 pericyte coverage of capillaries in the prefrontal cortex (PFC) than the non-maternal-separated
4 controls in adult. Furthermore, maternal separation induced a narrowing of capillary diameter in
5 both the *App*^{wt/wt} mice and *App*^{NL-G-F/wt} mice (Figure 2A). The more vessel-associated microglia
6 were detected in the maternal-separated *App*^{NL-G-F/wt} mice (Figure 2B). The result suggests
7 strengthening of the interaction between microglia and vessels in the maternal-separated *App*^{NL-}
8 *G-F/wt* mice. The vessel-associated microglia migrate towards cerebral vessel and cause increased
9 BBB permeability; Microglia, which can acquire a reactive phenotype that phagocytoses BBB
10 components, initiate leakage of systemic substances into the parenchyma (Haruwaka et al.,
11 2019). In our study, extra-vascular accumulation of fibrinogen deposits in the PFC was detected
12 only in the maternal-separated *App*^{NL-G-F/wt} mice (Figure 2C). We speculate that this
13 accumulation was caused by leakage of fibrinogen due to impairment of the vascular system by
14 more vessel-associated phagocytotic microglia only in the maternal-separated *App*^{NL-G-F/wt} mice.
15 Figure 2D shows the relationship between the sizes of the amyloid beta plaques and the
16 distances from the amyloid beta plaques to vessels in the maternal-separated and non-maternal-

1 separated $App^{NL-G-F/wt}$ mice that had mild amyloid beta plaque symptoms. The correlations
2 between the sizes of amyloid beta plaques and the distances from the amyloid beta plaques to
3 vessels were significant, which suggests the possibility that amyloid beta plaques were
4 generated around the vessels and caused disappearance of the vessels after disruption. We also
5 detected amoeboid microglia surrounding amyloid beta plaques stained with tomato-lectin
6 (Acarin et al., 1994).

7

8 *Either maternal separation or APP mutation caused activation of and morphological change in*
9 *microglia in adult mice*

10 As the vascular system is affected by microglia, we examined the activity and morphology of
11 microglia. Figure 3 shows the effects of maternal separation on microglia. Maternal separation
12 caused activation of the microglia in the PFC of both $App^{wt/wt}$ mice and $App^{NL-G-F/wt}$ mice (Figure
13 3A). Although the numbers of microglia were not significantly different between any of the
14 groups, except for the microglia surrounded by amyloid beta plaque, the proportions of soma
15 and dendrites stained by Iba1 were lower in both $App^{wt/wt}$ mice and $App^{NL-G-F/wt}$ mice subjected to
16 maternal separation (Figure 3A) than in the corresponding controls. Maternal separation may

18

1 have induced morphological changes in both genotypes. More detailed analysis of the
2 morphological changes revealed differences between non-maternal-separated $App^{wt/wt}$ mice and
3 other groups (Figure 3B). Lower numbers of dendrite terminals and a shortening of the total
4 branch length were detected in the maternal-separated $App^{NL-G-F/wt}$ mice than in the non-
5 maternal-separated $App^{wt/wt}$ mice. On the other hand, the number of dendrite terminals was
6 lower in the non-maternal-separated $App^{NL-G-F/wt}$ mice than in the non-maternal-separated
7 $App^{wt/wt}$ mice (Figure 3B). The non-maternal-separated $App^{NL-G-F/wt}$ mice may have milder
8 microglial symptoms than the maternal-separated animals. Furthermore, the circularity and
9 solidity of microglia increased in only the maternal-separated $App^{NL-G-F/wt}$ mice. We speculated
10 that maternal separation in combination with the APP mutation induced ameboid-like severe
11 morphological change.

12

13 *The maternal-separated $App^{wt/wt}$ and $App^{NL-G-F/wt}$ mice showed intact object location memory,*
14 *absence of amyloid beta plaque, and normal vascular morphology and pericyte in adolescence*

15 To detect initial changes in the maternal-separated and/or mutated mice, we examined the
16 phenotypes at an earlier stage (P90), that is, in adolescence. Figure 4A shows the effect of

1 maternal separation on object location memory in adolescence. In all groups, the discrimination
2 index for object location memory indicated object location memory to be intact. Figure 4B
3 shows that amyloid beta plaque was absent in all groups. Figure 4C shows no impairment of
4 vascular morphology or pericytes in any of the groups. We detect no changes in cognitive
5 function, plaque formation and vascular system in the maternal-separated and/or mutated mice
6 at adolescence, which allow us to rule out that the maternal-separated and/or mutated mice are
7 model mice for developmental disorders.

8

9 *Maternal separation and APP mutation cause functional and morphological change to*
10 *microglia in adolescence*

11 Figure 5 indicates the effect of maternal separation on microglia in adolescence. The numbers
12 of activated microglia were not significantly different between each group (Figure 5A). A more
13 detailed analysis of morphological differences revealed lower numbers of dendrite terminals in
14 the maternal-separated $App^{NL-G-F/wt}$ mice than in the non-maternal-separated $App^{wt/wt}$ mice. In the
15 $App^{wt/wt}$ mice, the maternally separated mice also showed lower numbers of dendrite terminals.
16 Furthermore, lower numbers of dendrite terminals were detected in the non-maternal-separated

1 *App*^{NL-G-F/wt} mice than in the non-maternal-separated *App*^{wt/wt} mice (Figure 5B). On the other
2 hand, the numbers of vessel-associated microglia did not differ between the groups (Figure 5C).
3 To evaluate the pro-inflammatory properties of microglia, we measured the level of TNF-R1
4 protein in the PFC. Maternal-separated *App*^{wt/wt} and *App*^{NL-G-F/wt} mice showed higher levels of
5 TNF-R1 protein than the non-maternal-separated mice (Figure 5D). We detected changes
6 associated with both maternal separation and heterozygotic mutation in adolescence, indicating
7 the possibility that a change in microglia caused by both maternal separation and APP mutation
8 induces an AD-like phenotype. To assess the effect of maternal separation on the hypothalamus-
9 pituitary-adrenal axis in adolescence, we measured serum corticosterone levels induced by a
10 novel cage stress in maternal-separated and non-separated mice. Maternal-separated *App*^{NL-G-F/wt}
11 and *App*^{wt/wt} mice subjected to a novel cage stress in adolescence mice showed more
12 corticosterone release than non-maternal-separated *App*^{NL-G-F/wt} and *App*^{wt/wt} mice (Figure 5E).
13

1 Discussion

2 In this study, we investigated whether maternal separation facilitates the development of AD,
3 with the aim of understanding the causal association between early-life stress and the
4 development and progression of AD. We also investigated for changes induced by maternal
5 separation, particularly in capillary vessels. We established that maternal separation facilitates
6 the impairment of spatial cognitive function and the formation of amyloid beta plaque in *App*^{NL-}
7 *G-F/wt* mice, with disruption of micro-capillaries, and we verified that early-life stress constitutes
8 a risk factor for AD. Furthermore, we found that morphological and functional changes to
9 microglia are early symptoms in our experimental model, and suggest the possibility that
10 impairment of the cerebral vascular system through microglia caused by combination of
11 maternal separation and APP mutation induces dysfunction in the BBB, thereby facilitating the
12 clinical condition of AD.

13 Research using the Childhood Trauma Questionnaire to assess childhood stress has suggested
14 that early-life stress may be associated with increased risk of AD and dementia (Radford et al.,
15 2017). However, because most previous reports, including the above research, were based on
16 retrospective cohort studies, it remains unclear whether early-life stress affects the onset and

1 progression of AD. Although it has been reported that maternal separation facilitates the
2 formation of amyloid beta plaque using an AD mouse model different from the one we used
3 (Hui et al., 2017), the pathway by which maternal separation induces increased amyloid beta
4 plaque remains unclear. We found that maternal separation facilitates the impairment of spatial
5 cognitive function and the formation of amyloid beta plaque through impairment of the cerebral
6 vascular system in *App*^{NL-G-F/wt} mice. We propose that these maternally separated *App*^{NL-G-F/wt}
7 mice can be used as a stress-induced AD mouse model. As it is known that progression of AD in
8 humans is exacerbated by stress, such as a change in abode and hospital admission in old age,
9 our AD mouse model may contribute to understanding the mechanism of stress-induced AD and
10 how it can be prevented.

11 We found that maternal separation mildly impaired the cerebral capillary vascular system.
12 However, dysfunction of the BBB, which is a more serious symptom, was detected in the
13 maternal-separated *App*^{NL-G-F/wt} mice and was related to both maternal separation and genetic
14 factors. Several mechanisms are involved in clear amyloid beta clearance (Chen et al., 2017).
15 One mechanism involves transcytosis through cells of the BBB. Another mechanism involves
16 phagocytosis by microglia. The reduction in amyloid beta clearance due to impairment of the

1 vascular system and impaired microglial phagocytoses of amyloid beta may induce
2 inflammation and increase fibrin generation due to coagulation factor XII (FXII) activation by
3 amyloid beta. The interaction of amyloid beta with fibrin (or fibrinogen) can lead to increased
4 fibrin deposition in cerebral blood vessels, and these accumulated fibrin deposits may induce
5 microinfarcts, which result in hemorrhage, inflammation, and BBB disruption, all of which are
6 commonly observed in AD (Ahn et al., 2017). Our result that disappearance of vessel around
7 larger amyloid plaque (Fig. 2D) may be caused by the interaction of amyloid beta with fibrin.

8 We report that maternal separation induces impairments to the cerebral vascular system, with
9 impairments to pericytes and narrowing of vessels in adulthood. Both of these phenotypes are
10 detected in the postmortem brains of human patients with AD (Halliday et al., 2016). Therefore,
11 these findings indicate a relationship between early-life stress and AD pathology. It has been
12 demonstrated that both antenatal and postnatal stress delay the development of the BBB during
13 weaning (Gomez-Gonzalez and Escobar, 2009). We detected no impairment in vascular
14 morphology or in pericytes in adolescence. Our findings that early-life stress induces
15 impairments in adult mice is in accord with the Development Origins of Health and Disease
16 (DOHaD) hypothesis, which purports that exposure to certain environmental conditions during

1 critical periods of development and growth may have significant adverse effects on short- and
2 long-term health. As the ultrastructural breakdown of capillaries and a decrease in blood flow
3 are detected in AD patients and AD mouse models (de la Torre, 2004), it is likely that
4 impairment of the cerebral vascular system develops before the onset of AD pathogeny in
5 *App*^{NL-G-F/wt} mice.

6 We detected morphological changes in microglia and changes in their pro-inflammatory
7 properties (which are mediated by the activation of TNF-R1) at an early stage, while we did not
8 detect amyloid plaque formation, changes to the vascular system, or changes in behavior.

9 Microglia have been reported to show morphological changes in response to the clinical
10 condition of AD (Yamada and Jinno, 2013). An increase in corticosterone in response to stress
11 has been reported to be related to a change in microglia (Rincon-Cortes and Sullivan, 2014). In
12 our study, maternal-separated *App*^{NL-G-F/wt} and maternal-separated *App*^{wt/wt} mice showed
13 increased corticosterone release in response to a novel cage stress at adolescence, which
14 suggests that maternal separation affects the stress response. In fact, the production of
15 corticosterone in response to stress facilitates the release of monocytes into the circulation from
16 bone marrow, which in turn facilitates monocyte adherence to vessels (Niraula et al., 2018). The

1 release of monocytes upregulates IL-1 β and stimulates endothelial cells (McKim et al., 2018).
2 Haruwaka et al. hypothesized that endothelial cells may respond to systemic inflammation and
3 release molecules to trigger microglia migration and phenotype changes. They also reported that
4 chemokine (C-C motif) ligand 5 (CCL5), which is secreted by endothelial cells, attracts
5 microglia to vessels. Recently, plasma from aged mice was shown to affect the activity of
6 microglia, referred to as neuroinflammation, and increase VCAM1 expression on endothelial
7 cell, a major cell type of the BBB, in young mice (Yousef et al., 2019). The authors suggested
8 that circulating cytokines and chemokines with detrimental effects on the brain give rise to these
9 changes in microglia and endothelial cells. Considering that increased age is crucial for the
10 development of AD, increases in inflammatory signals in the plasma may induce AD. On the
11 other hand, the non-maternal-separated *App*^{NL-G-F/wt} mice also showed morphological changes in
12 adolescence. Our data support the hypothesis that impairment of the vascular system by amyloid
13 beta oligomers induces activation of microglia (Nortley et al., 2019). As microglia and the
14 vascular system are deeply linked, the possibility that an undetectable alteration of the vascular
15 system led to a change in the microglia, which then led to the development of AD via an
16 interaction between microglia and the vascular system, cannot be ruled out. The changes in

1 microglia induced by maternal separation and those induced by APP mutation may occur via
2 different mechanisms. Microglia activation induced by maternal separation in combination with
3 *App*^{NL-G-F/wt} mice may impairs the vascular system, leading to AD progression.

4 In the present study, we established that maternal separation facilitates the development of
5 AD. Furthermore, we identified changes in microglia induced by maternal separation and/or
6 genetic factors as an early phenotype, which probably preceded the impairment of the vascular
7 system in the stress-induced AD mouse model. These results indicate that microglia could have
8 potential use as biomarkers and as targets for new therapies against AD.

9

10

11

1 Author contributions

2 T.T. and H.O. designed the research. T.T. performed and analyzed all experiments. T.S. and T.S.
3 generated and provided App mutant mice (*App*^{NL-G-F/NL-G-F}). S.H., M.H., H.S., and M.H. helped
4 with this study and made important suggestions. T.T. generated all the figures and tables and
5 wrote the manuscript. H.O. edited the manuscript and supervised the study.

6

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12

13 Competing interests

14 The authors declare that the research was conducted in the absence of any commercial or
15 financial relationships that could be construed as a potential conflict of interest.

16

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6

7

1 Figure Legends

2

3 Figure. 1 Maternal separation induces Alzheimer's disease-like behavior and amyloid beta
4 plaque formation.

5 (A) Time schedule for maternal separation and other investigations. (B) Spatial memory was

6 measured using an object location test. Behavioral layout of the object recognition test for the

7 four experimental groups. Maternal-separated $App^{NL-G-F/wt}$ mice showed impairment of spatial

8 memory. * $p < 0.05$ Maternal-separated $App^{NL-G-F/wt}$ mice vs non-maternal-separated $App^{wt/wt}$

9 mice; $n = 10$ / Non- maternal-separated $App^{wt/wt}$ mice, $n = 13$ / Non-maternal-separated App^{NL-G-

10 F/wt mice, $n = 8$ / Maternal-separated $App^{wt/wt}$ mice, $n = 11$ / Maternal-separated $App^{NL-G-F/wt}$ mice

11 (Dunnett's test). (C) Contextual memory and cued memory were measured using a fear-

12 conditioning test. Behavioral layout of the fear conditioning test for the four experimental

13 groups. Maternal-separated $App^{NL-G-F/wt}$ mice showed impairment of contextual memory.

14 Contextual test: Main effect of group, $F_{(3,68)} = 8.4507$, $p = 0.0001$; main effect of time, $F_{(4,272)} =$

15 5.3835 , $p = 0.0003$; interaction of group \times time, $F_{(12,272)} = 1.0939$, $p = 0.3651$; ** $p < 0.01$

16 Maternal-separated $App^{NL-G-F/wt}$ mice vs non-maternal-separated $App^{wt/wt}$ mice, *** $p < 0.001$

33

- 1 Maternal-separated $App^{NL-G-F/wt}$ mice vs non-maternal-separated $App^{NL-G-F/wt}$ mice; n = 19/ Non-
- 2 maternal-separated $App^{wt/wt}$ mice , n = 24/ Non-maternal-separated $App^{NL-G-F/wt}$ mice , n = 10/
- 3 Maternal-separated $App^{wt/wt}$ mice , n = 22/ Maternal-separated $App^{NL-G-F/wt}$ mice (two-way
- 4 repeated ANOVA followed by Holm's test). (D) The formation of amyloid beta plaque detected
- 5 by 82E1-biotin (cyan). The formation of amyloid beta plaque was facilitated by maternal
- 6 separation in maternal-separated $App^{NL-G-F/wt}$ mice. *** p < 0.001 Maternal-separated $App^{NL-G-F/wt}$
- 7 mice vs other groups n = 4/each group (Tukey-Kramer test). Scale bar = 100 μ m.
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- 9
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1 Figure. 2 Maternal separation induced impairment to the vascular system.

2 (A) The health of the vascular system was evaluated by tomato-lectin labelling of endothelial

3 cells (green) and CD13 labelling of pericytes (magenta). Maternal separation induced lower

4 pericyte coverage in $App^{wt/wt}$ mice and $App^{NL-G-F/wt}$ mice. * $p < 0.05$, Maternal-separated $App^{wt/wt}$

5 vs non-maternal-separated $App^{wt/wt}$ mice, ** $p < 0.01$ Maternal-separated $App^{NL-G-F/wt}$ vs non-

6 maternal-separated $App^{wt/wt}$ mice, ** $p < 0.01$, Maternal-separated $App^{wt/wt}$ vs non-maternal-

7 separated $App^{NL-G-F/wt}$ mice, *** $p < 0.001$ Maternal-separated $App^{NL-G-F/wt}$ vs non-maternal-

8 separated $App^{NL-G-F/wt}$ mice. Furthermore, maternal separation also induced a narrowing of

9 vascular diameter in both phenotypes. * $p < 0.05$ Maternal-separated $App^{wt/wt}$ mice vs non-

10 maternal-separated $App^{wt/wt}$ mice, *** $p < 0.001$ Maternal-separated $App^{NL-G-F/wt}$ mice vs non-

11 maternal-separated $App^{wt/wt}$ mice, ** $p < 0.01$ Maternal-separated $App^{NL-G-F/wt}$ mice vs non-

12 maternal-separated $App^{NL-G-F/wt}$ mice; $n = 3$ /each group (Tukey-Kramer test). Scale bar = 20 μ

13 m. (B) Microglia (White arrow in Left image) were detected by microglial labelling with Iba1

14 (magenta) and DAPI (cyan). Vessel-associated microglia (Grey square in Right image) were

15 detected by microglial labelling with Iba1 (magenta) and endothelial cell labelling with tomato-

16 lectin (green). In maternal-separated $App^{NL-G-F/wt}$ mice, vessel associated microglia were

1 increased. * $p < 0.05$ Maternal-separated $App^{NL-G-F/wt}$ mice vs non- maternal-separated $App^{wt/wt}$
2 mice; n = 3/ Non-maternal-separated $App^{wt/wt}$ mice, n = 3/ Non-maternal-separated $App^{NL-G-F/wt}$
3 mice, n =3/ Maternal-separated $App^{wt/wt}$ mice, n =4/ Maternal-separated $App^{NL-G-F/wt}$ mice
4 (Dunnett's test). (C) The integrity of the BBB was evaluated by tomato-lectin labelling of
5 endothelial cells (green) and fibrinogen (magenta). Maternal separation caused accumulation of
6 fibrinogen outside of vessels in maternal-separated $App^{NL-G-F/wt}$ mice. * $p < 0.05$ Maternal-
7 separated $App^{NL-G-F/wt}$ mice vs non-maternal-separated $App^{wt/wt}$ mice; n = 3/each group (Dunnett
8 test). Scale bar = 20 μ m. (D) The distance from the amyloid beta plaques to vessels and the
9 area of amyloid beta plaque were determined using 82E1-biotin labelling of amyloid beta
10 (magenta) and tomato-lectin labelling of endothelial cells (green). The area of amyloid beta
11 plaque showed a significant association with the distance from the amyloid beta plaque to
12 vessels in non-maternal-separated and maternal-separated $App^{NL-G-F/wt}$ mice. Small amyloid beta
13 plaques are indicated with cyan squares, and amyloid beta plaques with a size midway between
14 small and big amyloid beta plaques are indicated by grey squares. $r^2 = 0.987186$ $p < 0.001$; n =
15 12 (polynomial regression). Scale bar = 20 μ m.
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1 Figure. 3 Maternal separation and genetic factors activated and changed the morphology of
2 microglia.

3 (A) Microglial morphology and activation were detected by labelling microglia with Iba1
4 (magenta) and labelling activation with CD68 (cyan). Maternal separation increased in activated
5 microglia in $App^{wt/wt}$ and $App^{NL-G-F/wt}$ mice. * $p < 0.05$ Maternal-separated $App^{wt/wt}$ mice vs non-
6 maternal-separated $App^{wt/wt}$ mice, * $p < 0.05$ Maternal-separated $App^{NL-G-F/wt}$ mice or non-
7 maternal-separated $App^{wt/wt}$ mice. * $p < 0.05$ Maternal-separated $App^{NL-G-F/wt}$ mice or non-
8 maternal-separated $App^{NL-G-F/wt}$ mice (Tukey-Kramer test). Furthermore, occupancy of soma and
9 dendrites in maternal-separated $App^{wt/wt}$ and maternal-separated $App^{NL-G-F/wt}$ mice was lower. $n =$
10 3/ Non-maternal-separated $App^{wt/wt}$ mice, $n = 3$ / Non-maternal-separated $App^{NL-G-F/wt}$ mice, n
11 =3/ Maternal-separated $App^{wt/wt}$ mice, $n = 4$ / Maternal-separated $App^{NL-G-F/wt}$ mice (Tukey-
12 Kramer test). Scale bar = 10 μ m. (B) Detailed morphological analysis was performed using
13 the ImageJ plugin AnalyzeSkeleton. Images show examples of the outlines of microglia.

14 Maternal separation and APP mutation decreased dendritic terminals in $App^{wt/wt}$ and $App^{NL-G-F/wt}$
15 mice. * $p < 0.05$ Non-maternal-separated $App^{NL-G-F/wt}$ mice vs non-maternal-separated $App^{wt/wt}$
16 mice, ** $p < 0.01$ Maternal-separated $App^{wt/wt}$ mice vs non-maternal-separated $App^{wt/wt}$ mice,

37

1 *** $p < 0.001$ Maternal-separated $App^{NL-G-F/wt}$ mice vs non-maternal-separated $App^{wt/wt}$ mice
2 (Dunnett test). Furthermore, the total branch length in maternal-separated $App^{NL-G-F/wt}$ mice and
3 maternal-separated $App^{wt/wt}$ mice were shorter. * $p < 0.05$ Maternal-separated $App^{wt/wt}$ mice vs
4 non-maternal-separated $App^{wt/wt}$ mice, ** $p < 0.01$ Maternal-separated $App^{NL-G-F/wt}$ mice vs non-
5 maternal-separated $App^{wt/wt}$ mice. The circularity of microglia increased in only the maternal-
6 separated $App^{NL-G-F/wt}$ mice. ** $p < 0.01$ Maternal-separated $App^{NL-G-F/wt}$ mice vs non-maternal-
7 separated $App^{wt/wt}$ mice. The solidity of microglia increased in only the maternal-separated
8 $App^{NL-G-F/wt}$ mice. ** $p < 0.01$ Maternal-separated $App^{NL-G-F/wt}$ mice vs non-maternal-separated
9 $App^{wt/wt}$ mice; $n = 4$ /each group (Dunnett test).

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1 Figure. 4. Alzheimer's disease-like phenotypes did not begin during adolescence in maternal-
2 separated $App^{NL-G-F/wt}$ mice, and impairment of the vascular system did not begin during
3 adolescence in all groups.

4 (A) Spatial memory was measured using an object location test. Maternal-separated $App^{NL-G-F/wt}$
5 mice showed normal cognitive function. n = 9/ Non-maternal-separated $App^{wt/wt}$ mice, n = 11/
6 Non-maternal-separated $App^{NL-G-F/wt}$ mice, n = 12/ Maternal-separated $App^{wt/wt}$ mice, n = 11/
7 Maternal-separated $App^{NL-G-F/wt}$ mice (B) The formation of amyloid beta plaque detected by
8 82E1-biotin labelling of amyloid beta. No amyloid beta plaque was detected in maternal-
9 separated $App^{NL-G-F/wt}$ mice. n = 3/each group (C) The health of the vascular system was
10 evaluated by tomato-lectin labelling of endothelial cells (green) and CD13 labelling of pericytes
11 (magenta). All groups showed normal pericyte coverage and a healthy vascular diameter. n =
12 3/each group. Scale bar = 20 μ m.

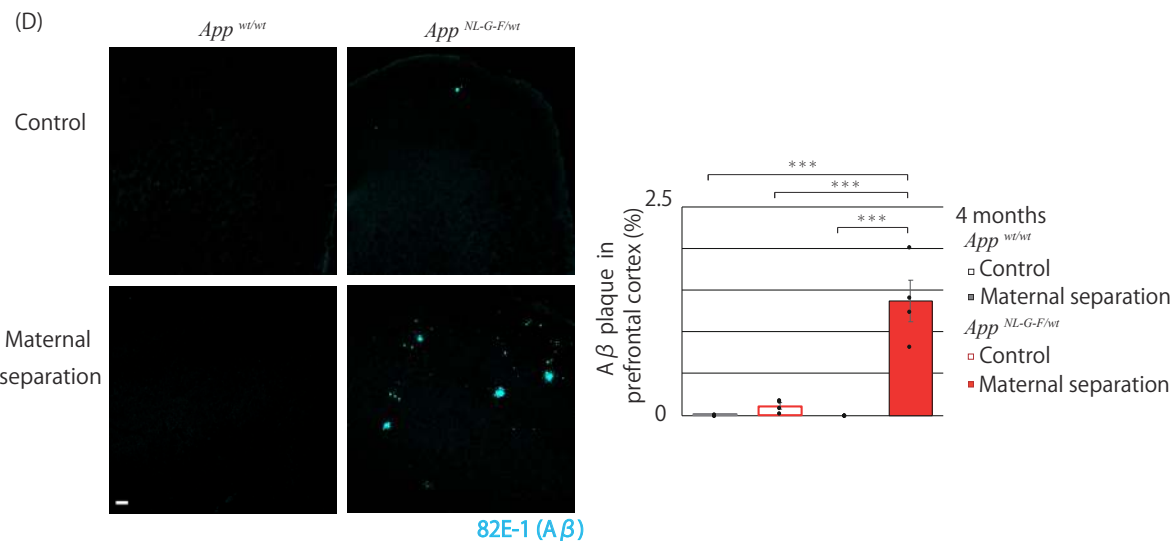
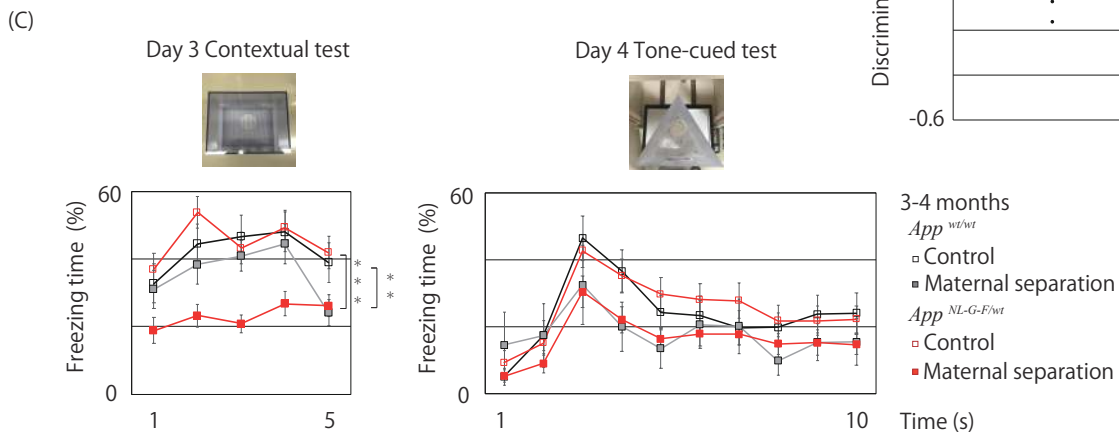
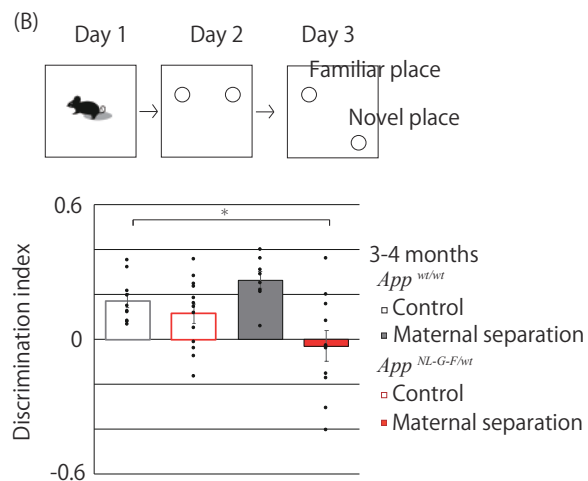
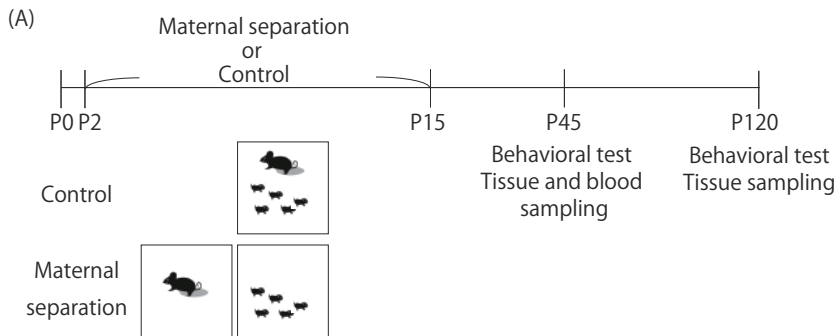
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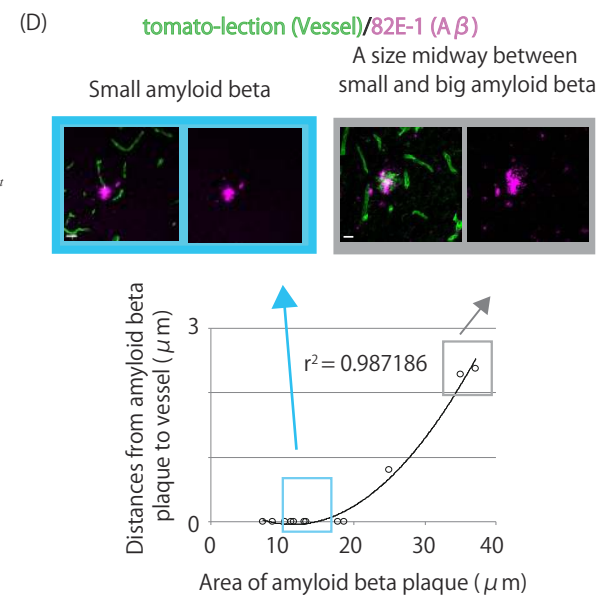
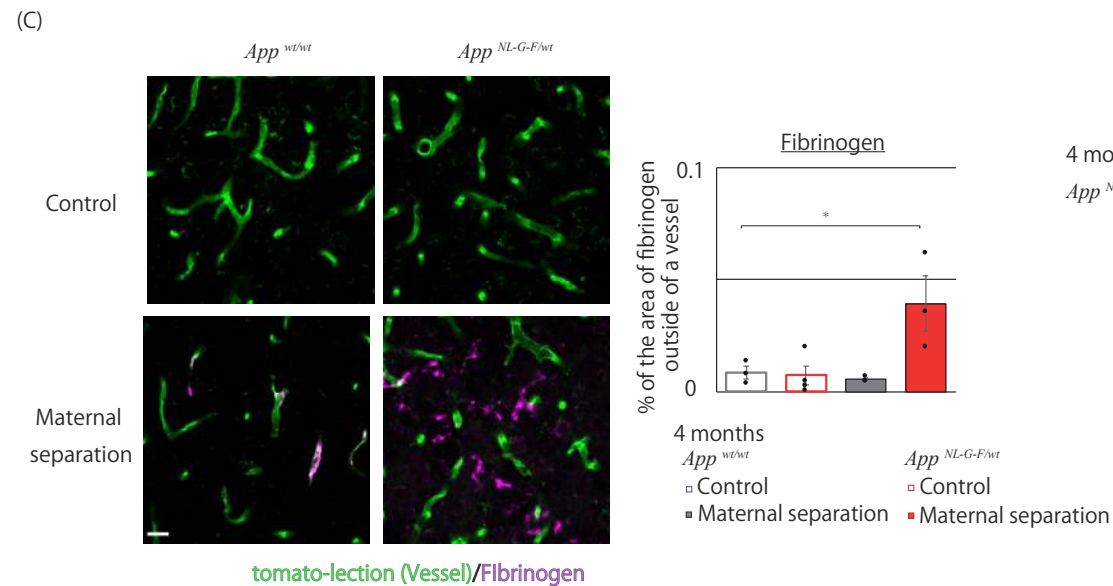
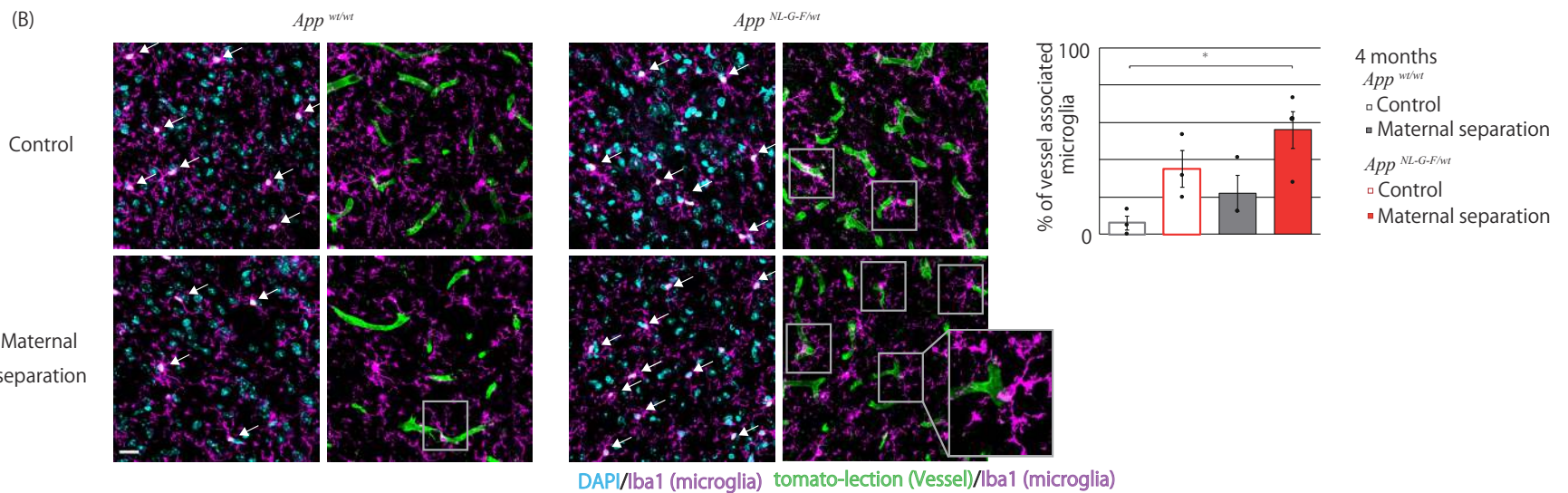
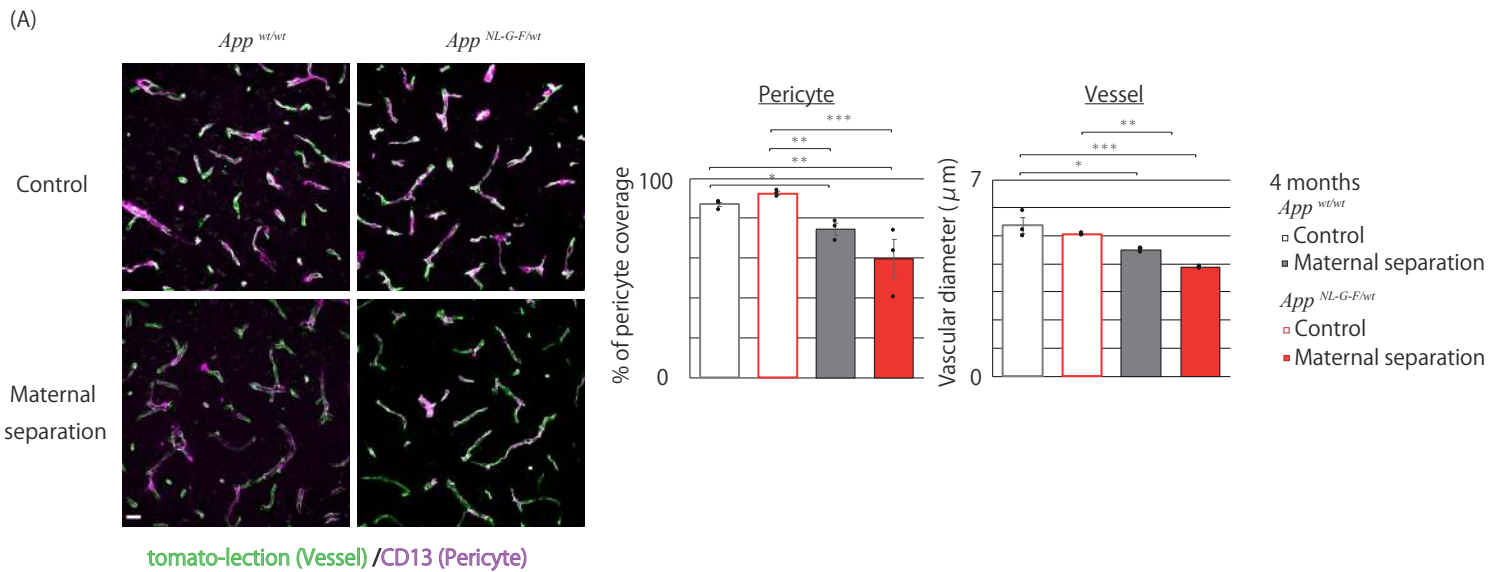
1 Figure 5. Maternal separation and genetic factors induced morphological changes in microglia
2 and maternal separation in combination with genetic factors led to abnormality in the stress
3 response during adolescence.

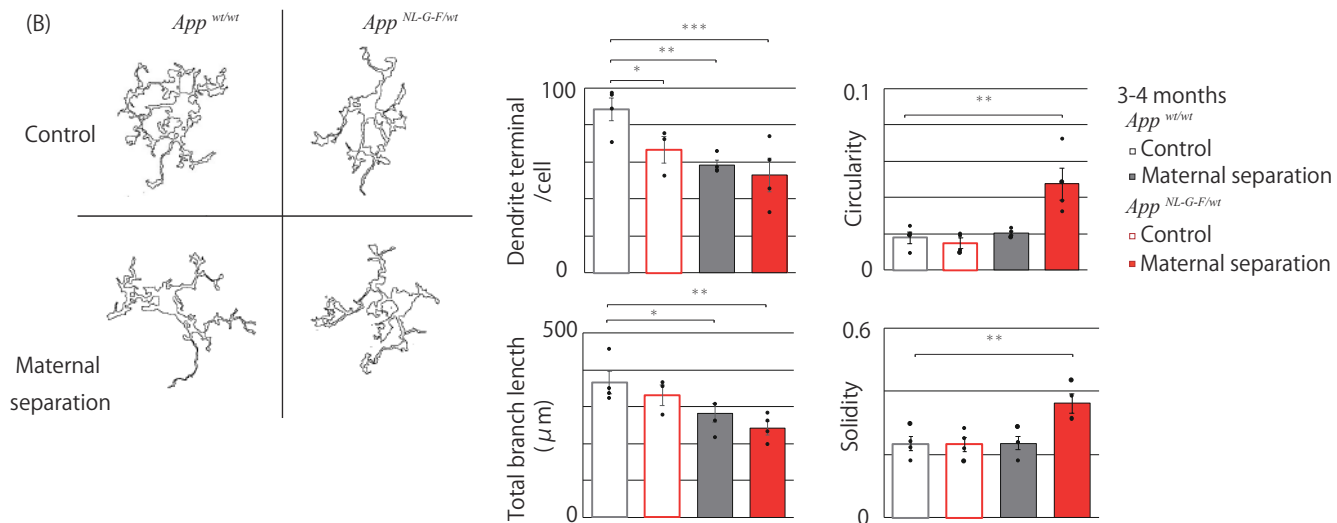
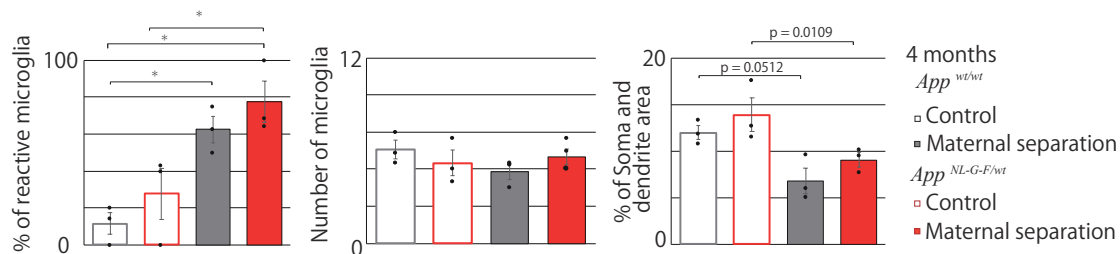
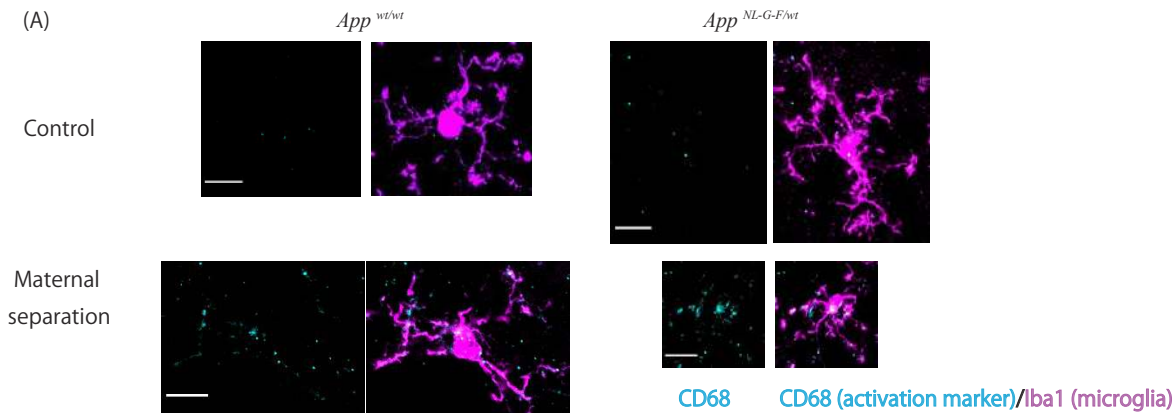
4 (A) Microglial morphology and activation were detected by labelling with Iba1 (magenta) and
5 CD68 (cyan), respectively. Increased activation of microglia was not observed in maternal-
6 separated *App*^{wt/wt} and maternal-separated *App*^{NL-G-F/wt} mice. n = 3/each group. Scale bar = 10 μ
7 m. (B) Detailed morphological analysis was performed using the ImageJ. Maternal separation
8 and APP mutation decreased dendritic terminals in *App*^{wt/wt} and *App*^{NL-G-F/wt} mice. *** p < 0.001
9 Non-maternal-separated *App*^{NL-G-F/wt} mice vs non-maternal-separated *App*^{wt/wt} mice, ** p < 0.01
10 Maternal-separated *App*^{wt/wt} mice vs non-maternal-separated *App*^{wt/wt} mice, ** p < 0.01
11 Maternal-separated *App*^{NL-G-F/wt} mice vs non-maternal-separated *App*^{wt/wt} mice (Dunnett test). (C)
12 Microglia (white arrow in left image) were detected by microglial labelling with Iba1 (magenta)
13 and DAPI (cyan). Vessel-associated microglia (Grey square in right image) were detected by
14 microglial labelling with Iba1 (magenta) and endothelial cell labelling with tomato-lectin
15 (green). The numbers of vessel-associated microglia were similar across the four experimental
16 groups. n = 3/each group. Scale bar = 20 μ m. (D) The protein level of TNF-R1 was higher in

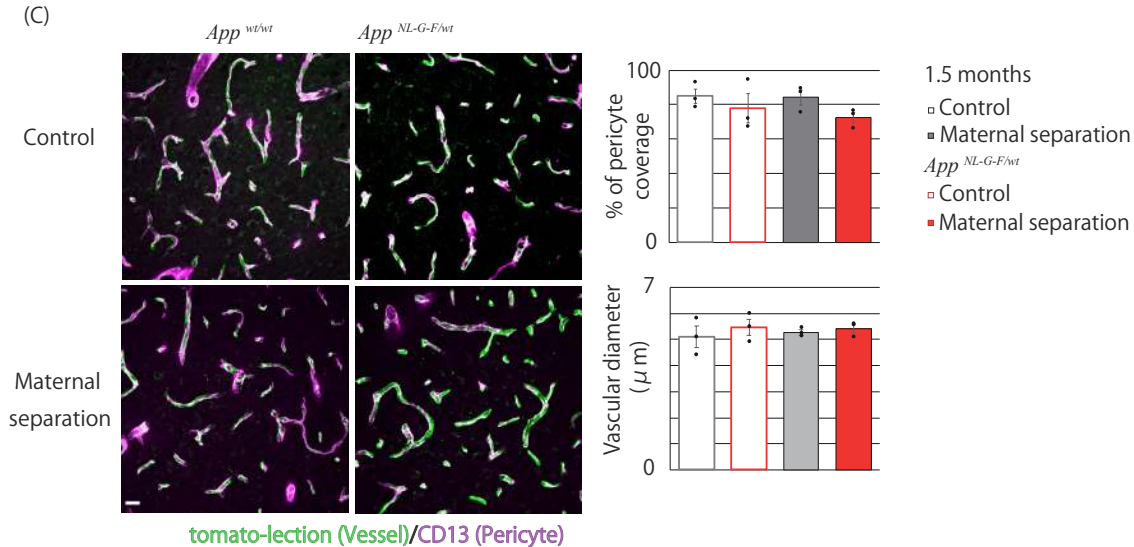
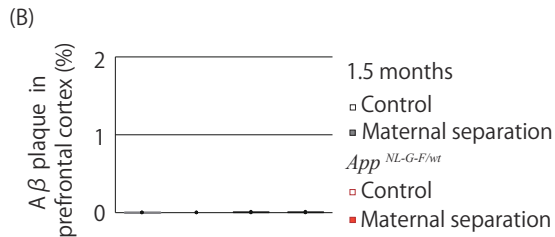
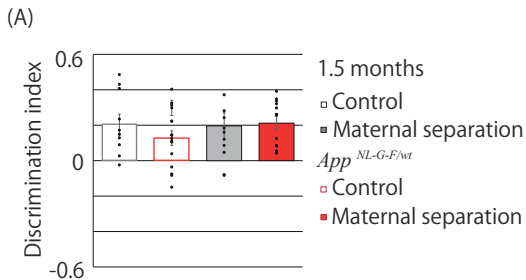
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- 1 maternal-separated $App^{wt/wt}$ and maternal-separated $App^{NL-G-F/wt}$ mice. $p = 0.058$ Maternal-
- 2 separated $App^{wt/wt}$ mice vs non-maternal-separated $App^{wt/wt}$ mice, $p = 0.054$ Maternal-separated
- 3 $App^{NL-G-F/wt}$ mice vs non-maternal-separated $App^{wt/wt}$ mice $n = 3$ /each group (Dunnett test). (E)
- 4 The response to novel cage stress was measured to assess an effect of maternal separation on the
- 5 HPA axis. Not only $App^{NL-G-F/wt}$ mice but also $App^{wt/wt}$ mice showed that increased stress caused
- 6 by maternal separation induced corticosterone release. Maternal-separated $App^{NL-G-F/wt}$ mice: F
- 7 $(1,2) = 75.414$, $p = 0.0130$; Maternal-separated $App^{wt/wt}$ mice: $F (1,2) = 22.035$, $p = 0.0425$. * $p <$
- 8 0.05 vs corticosterone before novel cage stress; $n = 3$ /each group (Paired t-test).

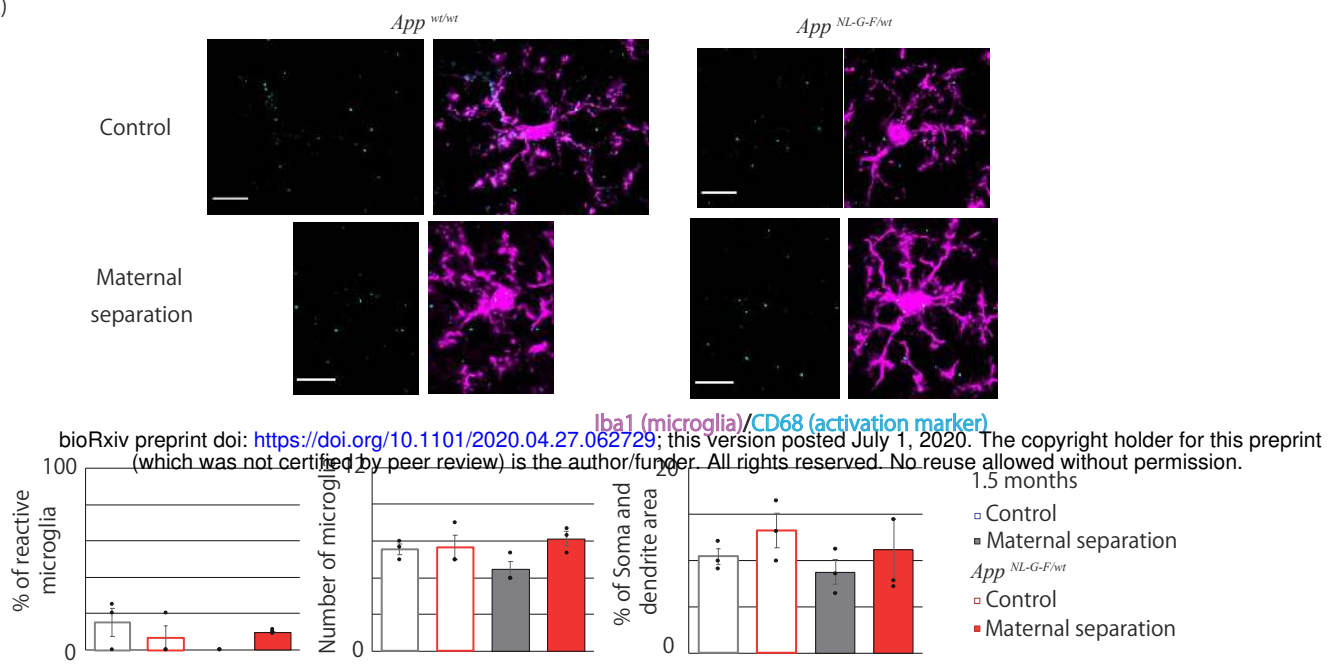




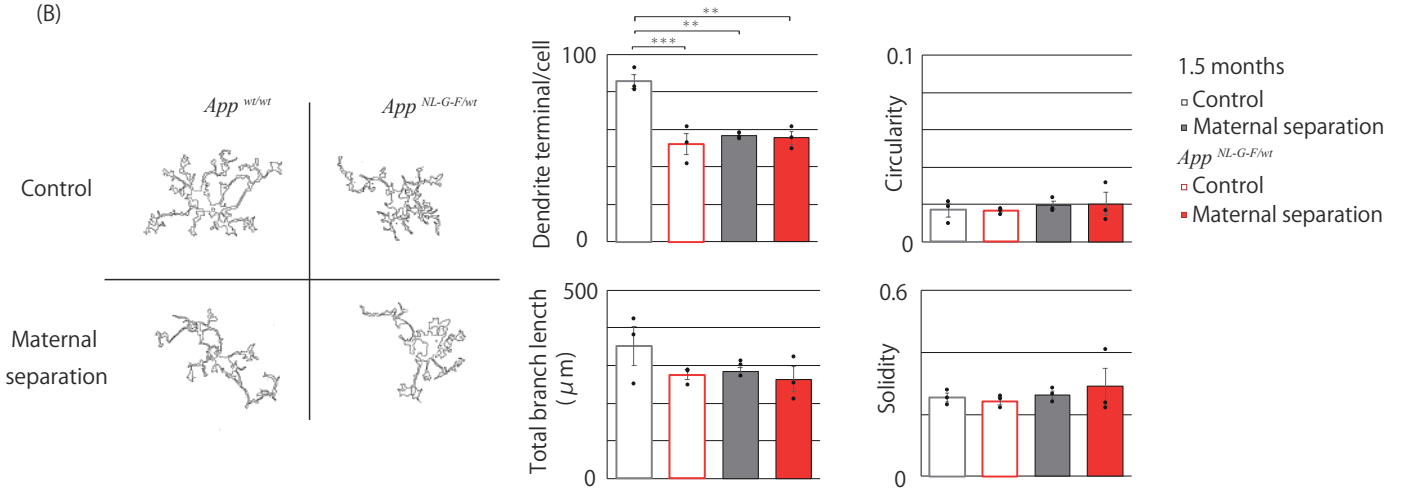




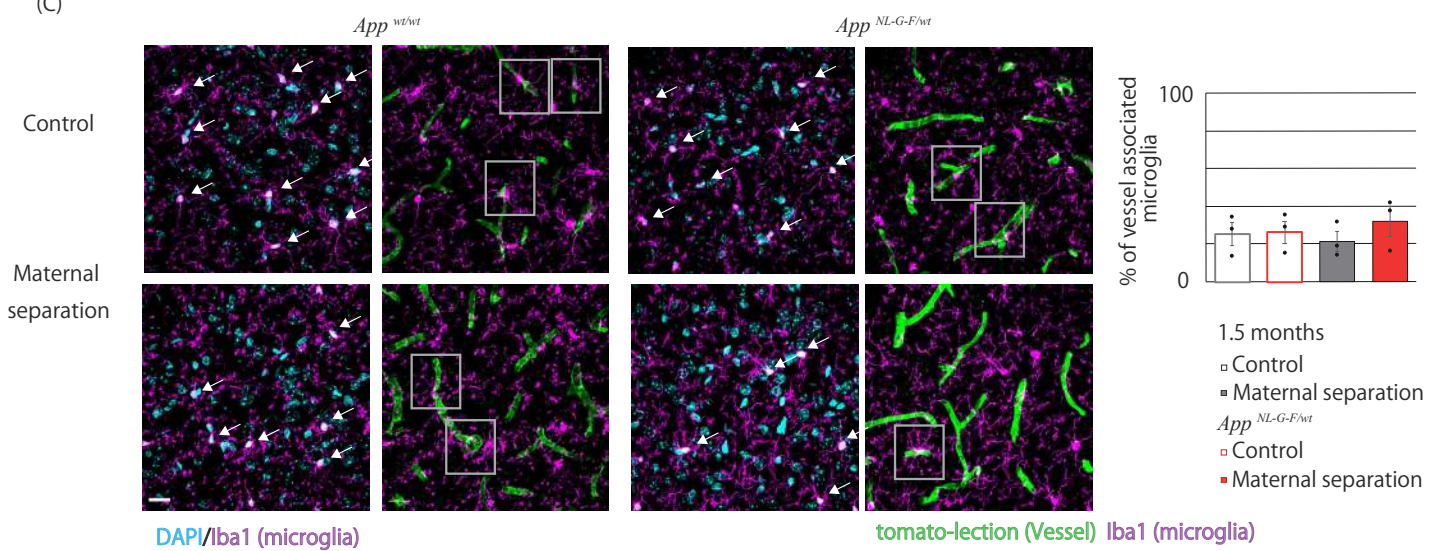
(A)



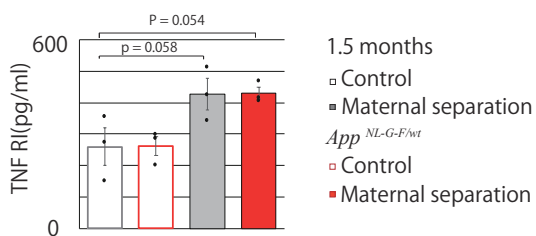
(B)



(C)



(D)



(E)

