

1 **Amyloid-associated hyperconnectivity drives tau spread across**
2 **connected brain regions in Alzheimer’s disease**

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Formatiert: Links

1 **OVERLINE: NEURODEGENERATION**

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4 **One sentence summary:** Aβ induces neuronal hyperconnectivity, which promotes tau
5 spreading from temporal lobe epicenters to connected brain regions in Alzheimer's disease

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8 **Editor's Summary**

9
10 **ABSTRACT**

11 In Alzheimer's disease, Aβ triggers the aggregation and spreading of tau pathology which
12 drives neurodegeneration and cognitive decline. However, the pathophysiological link between
13 Aβ and tau remains unclear, which hinders therapeutic efforts to attenuate Aβ-related tau
14 accumulation. Aβ has been found to trigger neuronal hyperactivity and hyperconnectivity, and
15 preclinical research has shown that tau spreads across connected neurons in an activity-
16 dependent manner. Therefore, Aβ may induce neuronal hyperactivity and hyperconnectivity
17 thereby promoting tau spread across connected brain regions. By combining Aβ-PET, resting-
18 state fMRI and longitudinal tau-PET in 69 cognitively normal controls devoid of amyloid
19 pathology and 140 patients with a positive amyloid-PET scan covering the AD spectrum, we
20 confirmed that Aβ induces hyperconnectivity of temporal lobe tau epicenters to posterior brain
21 regions that are vulnerable to tau accumulation in AD. This was replicated in an independent
22 sample of 55 controls and 345 individuals with preclinical AD and low cortical tau-PET uptake,
23 suggesting that the emergence of Aβ-related hyperconnectivity precedes neocortical tau
24 spreading typically observed in clinical stages of AD. Lastly, using longitudinal tau-PET, we
25 confirmed that Aβ-related connectivity increases of tau epicenters to other brain regions
26 mediated the effect of Aβ on faster tau accumulation. Together, these findings suggest that Aβ
27 promotes tau spreading by eliciting neuronal hyperconnectivity and that targeting Aβ-related
28 neuronal hyperconnectivity may attenuate tau spreading in AD.

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hat gelöscht: Supporting the notion that fMRI-detected hyperconnectivity may mirror Aβ-associated neuronal hyperactivity, we found that compared to controls, neurons in a small AD post-mortem dataset expressed higher c-Fos, a Calcium-sensitive marker of ante-mortem neuronal activity.

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1 INTRODUCTION

2 Alzheimer's disease (AD) is initiated by brain-wide amyloid- β ($A\beta$) accumulation, followed by
3 the spread of neurofibrillary tau pathology from the temporal lobe across connected brain regions,
4 ensuing neurodegeneration and cognitive decline.(1-4) This well-established amyloid-cascade
5 model of AD is strongly supported by genetics and in vivo biomarker studies,(5-7) showing that
6 mutations in $A\beta$ -associated genes *APP*, *PSEN1* and *PSEN2*(δ) but not in tau-associated genes such
7 as *MAPT* (9) can cause full AD pathology in humans, including $A\beta$ plaque and tau tangle
8 formation. Furthermore, neuropathological and neuroimaging studies have demonstrated tau
9 pathology to arise in the medial temporal lobe in regions belonging to Braak stages I and II, in a
10 large proportion of cognitively normal elderly as well as AD patients (10-12), whereas the
11 dispersion of tau pathology to temporo-parietal brain regions of Braak stages III and IV is strongly
12 dependent on concomitant $A\beta$ pathology (11, 12). Thus, $A\beta$ is assumed to trigger the spreading of
13 tau from the medial temporal lobe to the neocortex, which constitutes the major driver of
14 neurodegeneration and cognitive decline in AD(13, 14) and it is therefore of pivotal clinical interest
15 to understand the link between $A\beta$ and tau to therapeutically target the $A\beta$ -tau axis.

16 Preclinical studies have shown that $A\beta$ -oligomers can directly trigger tau phosphorylation in
17 cultured hippocampal neurons(15) and that $A\beta$ promotes the local aggregation of soluble tau in
18 combined $A\beta$ and tau transgenic mice.(16) However, post-mortem and PET studies in AD patients
19 have consistently shown that $A\beta$ and tau accumulation patterns are spatially distinct, especially in
20 early stages of AD, with $A\beta$ accumulating rather diffusely in the neocortex(17-19) whereas tau
21 typically emerges rather focally in the medial temporal lobe with subsequent spread to connected
22 neocortical regions.(2, 20, 21) Due to the spatial mismatch between $A\beta$ and tau deposition patterns,
23 it is unlikely that tau accumulation in AD is entirely induced by local $A\beta$. In addition, age-related
24 medial temporal lobe tau pathology also occurs in the absence of cortical $A\beta$, suggesting that $A\beta$
25 and tau accumulation can start independently of each other.(22, 23)

26 Our lead hypothesis that cortical $A\beta$ indirectly triggers the spreading of tau pathology(22, 23) from
27 local epicenters defined as regions with highest tau pathology at a given point in time, across
28 connected brain regions by inducing neuronal hyperactivity and hyperconnectivity offers an
29 alternative explanation for the link between $A\beta$ and tau.(24, 25) This hypothesis is based on in vitro
30 studies which have shown that tau is actively released from neurons and spreads across synaptic
31 connections and that the rate of neuronal tau release and trans-synaptic spread is dependent on
32 neuronal activity.(24, 26, 27) Congruently, PET-assessed tau pathology expands from local
33 epicenters to regions that are functionally connected and show correlated neuronal activity on MEG

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1 and resting-state fMRI.(2, 28-35) A β has previously been described as a potent trigger of neuronal
2 hyperexcitability, increased neuronal firing and hypersynchronicity in cell and animal
3 models,(36-41) explained by increased Glutamate release(42), reduced GABA sensitivity(43),
4 and increased postsynaptic Ca²⁺-influx.(44) Further, functional MRI (fMRI) studies have
5 reported A β -related functional connectivity increases as measured by higher inter-regional
6 synchrony of neuronal activity(45) in A β -transgenic mice and AD patients (46-50). In fact, changes
7 in functional connectivity in AD patients are considered to occur early in AD when A β is the
8 dominant pathology, where a cascading network failure is associated with the progression of AD
9 pathophysiology (50-54). In addition, patients suffering from autosomal dominant and sporadic
10 AD show an increased prevalence of subclinical epileptiform neuronal activity and seizures at
11 early disease stages, together supporting the notion that A β induces an increase in neuronal
12 activity and connectivity.(55-59)

13 We hypothesize that an A β -related increase in neuronal activity and connectivity is critical for
14 triggering the trans-neuronal spread of tau, since enhanced neuronal activity amplifies synaptic tau
15 release(60) and exacerbates trans-synaptic tau spread.(24) Supporting this, recent work in
16 combined A β and tau transgenic mice has specifically shown that A β -associated neuronal
17 hyperactivity triggers tau spread from the entorhinal cortex to connected regions.(61) However, it
18 is not known whether A β -induced increases in neuronal activity and connectivity actively
19 contribute to the spreading of tau pathology across interconnected brain regions in AD patients.
20 Establishing this link bears a high potential to therapeutically attenuate tau spreading in AD and
21 thus halt neurodegeneration and cognitive decline(25).

22 To address this, we included 209 participants from the Alzheimer's disease Neuroimaging
23 Initiative (ADNI), encompassing 69 A β -negative (A β -) cognitively normal (CN) controls and 140
24 A β -positive (A β +) patients across the preclinical to dementia spectrum of AD, with available
25 amyloid-PET and 3T resting-state fMRI to model functional connectivity and A β -related
26 connectivity increases, as well as longitudinal [¹⁸F]Flortaucipir tau-PET with approximately 2.7
27 years of follow-up to model the accumulation and spread of tau pathology. In this dataset, we tested
28 i) whether regional A β deposition induces connectivity increases to epicenters of earliest tau
29 deposition, ii) whether stronger connectivity of tau epicenters to a given brain region is linked to
30 faster tau accumulation in that region and iii) whether accelerated spread of tau across connected
31 regions is mediated by A β -associated connectivity increases. Cross-sectional analyses between A β
32 and connectivity increases of tau epicenters were replicated using baseline amyloid-PET, tau-PET

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1 and resting-state fMRI data of 345 preclinical AD patients (cognitively normal, A β +) and 55 A β -
2 controls of the A4 study.↓
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hat gelöscht: Lastly, we aimed to support the notion of a potential role of A β in triggering neuronal hyperactivity in AD patients, by exploratorily assessing the proportion of c-Fos (i.e., an established marker of neuronal activity) expressing neurons in a small AD post-mortem brain tissue dataset compared to non-demented controls. (62)

1 **RESULTS**

2 Our primary neuroimaging sample from the ADNI database included 209 subjects, with 69
3 controls **devoid of amyloid pathology** (CN-A β -) and 140 A β + subjects across the preclinical
4 to dementia spectrum of AD (68 CN-A β +, 47 Mild Cognitive Impairment [MCI]-A β +, 25
5 Dementia-A β +). All ADNI subjects had available baseline Flortbetaben (n=71) or Flortbetapir
6 (n=138) amyloid-PET, 3T resting-state fMRI and longitudinal Flortaucipir tau-PET with a
7 mean follow-up time of 2.74 years. Clinical characteristics and patient demographics are shown
8 in Table 1. Amyloid and tau-PET SUVRs were parcellated into 200 cortical ROIs using the
9 Schaefer atlas, excluding typical subcortical sites of Flortaucipir off-target binding.(62, 63) For
10 amyloid-PET data, regional and global amyloid-PET **data were** intensity normalized to the
11 whole cerebellum, **and** transformed to centiloid to harmonize Flortbetaben and Flortbetapir
12 amyloid-PET tracers.(64) Longitudinal change rates in tau-PET SUVRs **obtained using** **an** the
13 inferior cerebellar grey **reference** were computed for each ROI using a previously established
14 approach, employing linear-mixed effects models controlling for random slope and
15 intercept.(29) Surface renderings of baseline tau-PET SUVRs are shown Figure 1A, illustrating
16 increasing tau-PET load across the AD spectrum. Longitudinal tau-PET change rates are shown
17 in Figure 1B, showing overall faster tau-PET increase in AD spectrum patients vs. CN A β -
18 subject. For validation of cross-sectional analyses regarding the association between A β and
19 connectivity increases, we included data from 400 CN participants of the A4 study (55 CN-
20 A β -, 345 CN-A β +), with available cross-sectional Flortbetapir amyloid-PET, Flortaucipir tau-
21 PET and 3T resting-state fMRI (see Table 1 for demographics).

22

23 ***Amyloid is associated with increased connectivity of tau epicenters to posterior brain regions***

24 Our first aim was to determine whether regional A β is associated with functional connectivity
25 increases of tau epicenters. To address this, we first determined subject-specific tau epicenters
26 in A β + subjects, defined as 5% **of** cortical brain regions with highest baseline tau-PET SUVRs
27 (Fig. 2A), following our previously established protocols.(2, 31, 65) Subject-specific tau
28 epicenters were subsequently used as seed regions for assessing subject-level resting-state
29 functional connectivity to the remaining 95% non-epicenter brain regions (Fig. 2A-D). A
30 group-average mapping of epicenter location and epicenter connectivity of the ADNI A β +
31 subjects is shown in Figure 2E, illustrating that tau epicenters are typically located in the
32 inferior temporal lobe and show strong connectivity to posterior brain regions (Fig. 2F). We
33 then assessed whether A β is linked to a strengthening of the connectivity between the tau

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1 epicenters and target regions. To this end, we determined ROI-wise linear regression models
2 with functional connectivity between the tau epicenter and the target ROI as the dependent
3 variable, and A β load defined as centiloid averaged across the tau epicenter and target ROI to
4 account for A β at both ends of the connection as a predictor, controlling for age, sex, ethnicity,
5 diagnosis, MRI scanner type and average motion during the resting-state fMRI scan as
6 determined by mean framewise displacement. Supporting our hypothesis, we found that higher
7 A β was associated with stronger connectivity of tau epicenters to widespread posterior
8 temporo-parietal brain regions that typically accumulate tau relatively early in the course of
9 AD ((12, 21, 66) Fig.3A). To illustrate this point further, the average centiloid values vs. tau
10 epicenter connectivity of significant regions described in Fig. 3A is shown in Fig. 3B,
11 confirming that higher A β is related to stronger connectivity. In addition, we determined the
12 overall distribution of the regression derived beta-values of the association between centiloid
13 and connectivity (Fig. 3C), showing that beta-values were significantly greater than zero
14 (95%CI=[0.10;0.14], T=11.693, p<0.001). This result pattern supports our hypothesis that A β
15 is linked to an overall connectivity increase of the tau epicenter to the rest of the brain.
16 Supporting the robustness of these findings, these results were replicated in 345 A β + subjects
17 with preclinical AD of the A4 sample, showing a highly congruent spatial pattern of A β -related
18 connectivity increases of tau epicenters to posterior brain regions (Fig. 3D). This result was
19 also congruent when reclassifying A β -positivity in A4 subjects at a slightly more restrictive
20 amyloid-PET SUVR threshold of 1.15 as previously suggested by the A4 imaging core (see
21 Fig.S1).(67) Similar results were obtained when restricting the ADNI cohort to cognitively
22 normal A β + subjects (n=68) to match the A4 cohort of preclinical AD subjects (Fig.S2), overall
23 supporting the view that A β -related connectivity changes occur early in AD. All results were
24 specific for A β + subjects, as no associations between A β and tau epicenter connectivity were
25 detected in A β - subjects of ADNI or A4.

26 To investigate further if A β -related connectivity increases were specific to tau epicenters, we
27 examined the correlation of the connectivity between two ROIs and their average amyloid-PET
28 centiloids for each A β + subject. We created a 200x200 amyloid load matrix for each subject,
29 representing cumulative amyloid levels across ROI pairs, which we then correlated with the
30 subject-specific functional connectivity matrix. The results showed positive correlations
31 between amyloid load and connectivity across ROI pairs within A β + subjects of ADNI
32 (T=4.571, p<0.001) and A4 (T=7.793, p<0.001), indicating that ROI pairs with higher amyloid
33 loads tend to have higher connectivity, regardless of being a tau epicenter or not (see Fig.S3).

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1 Using an alternative metric of neuronal activity (FDG-PET to measure cortical glucose
2 consumption intensity normalized to the Pons) that was available in a subset of A β + ADNI
3 patients (n=61) confirmed that higher regional amyloid-PET was related to higher glucose
4 metabolism (mean/SD subject-level correlation between regional amyloid-PET and FDG-
5 PET=0.33/0.16; 95%CI=[0.29,0.38], T=15.91, p<0.001). Together, these results suggest that
6 A β is linked to increased functional connectivity potentially driven by neuronal hyperactivity.

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8 ***Regions that are more strongly connected to the tau epicenter show faster tau accumulation***

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9 Next, we addressed our second aim, whether stronger connectivity of the tau epicenter to other
10 brain regions is associated with faster tau accumulation in those regions, indicative of faster
11 tau spread across connections. This analysis was performed using ADNI data only, as no
12 longitudinal tau-PET data were available in A4. We performed ROI-wise regression models in
13 A β + subjects, assessing whether higher subject-specific connectivity of the tau epicenter to a
14 given region predicted faster tau-PET change rates in that region, controlling for age, sex,
15 ethnicity, diagnosis, MRI scanner type and motion during the fMRI scan. We found widespread
16 associations (p<0.05, FDR-corrected) between tau epicenter connectivity and faster tau
17 accumulation, predominantly for temporo-parietal, occipital and superior frontal brain regions
18 (see Fig. 4A) showing wide overlap with regions in which we detected A β -related connectivity
19 increases (see black outline in Fig. 4A). When assessing the distribution of region-specific
20 associations between epicenter connectivity and tau-PET increase over time (regression
21 derived beta-values displayed in Fig. 4A), beta values were greater than zero
22 (95%CI=[0.16;0.2], T=20.012, p<0.001, Fig. 4B), suggesting that higher epicenter
23 connectivity is overall related to faster tau accumulation. Importantly these results were
24 obtained using subject-level connectivity data, showing that patient-specific connectivity
25 patterns and strength are relevant for tau spreading. Together, these results support the
26 hypothesis that brain regions that are more strongly connected to the tau epicenters on the
27 individual level show faster tau accumulation over time, in line with the concept of trans-
28 neuronal tau spreading.

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30 ***The association between A β and tau accumulation is mediated by A β -related connectivity***
31 ***increases***

32 For our third aim, we assessed A β -induced connectivity changes as a putative
33 pathophysiological link between A β and tau spreading. Again, this analysis was restricted to

1 the ADNI dataset with available longitudinal tau-PET. In the ADNI A β + subjects, we
2 determined ROI-wise bootstrapped mediation models with 1000 iterations, testing whether
3 higher A β (centiloids averaged across the tau epicenter and any given target ROI) was linked
4 to faster tau accumulation in a given target ROI, and whether this effect was mediated by a A β -
5 related functional connectivity increases between the target ROI and the tau epicenter (concept
6 illustrated in Fig. 4C). All mediation models were controlled for age, sex, ethnicity, diagnosis,
7 MRI scanner type and motion during the fMRI scan and FDR-corrected for multiple
8 comparisons ($p < 0.05$). Supporting our hypothesis, we found that faster tau-PET accumulation
9 in temporo-parietal regions were mediated by A β -related connectivity increases of these brain
10 regions to the tau epicenter (Fig. 4C and D). Taken together, these results favor a
11 pathophysiological disease model in which regional A β induces stronger functional
12 connectivity to the tau epicenter, which in turn facilitates the spreading of tau from the
13 epicenters to connected brain regions.

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DISCUSSION

A β deposition has been shown to trigger neuronal hyperactivity, hypersynchronicity and hyperconnectivity in animal and cell models(39-41, 68) and to be associated with epileptiform brain activity in AD mouse models and AD patients.(37, 56-58) We hypothesized that A β -induced neuronal hyperactivity manifests in functional connectivity increases (52-54, 69) and thereby accelerates the activity-dependent spread of tau pathology across connected brain regions in AD patients.(24, 70) Prior studies have identified functional connectivity increases in AD patients as one of the earliest pathological findings in AD, which has led to the formation of the cascading network hypothesis (52-54), which claims a key role of early network dysfunction in promoting AD pathophysiology. In the current study we aimed to integrate the insights on early changes of functional connectivity in AD with our previous findings on connectivity-based tau spreading to test whether A β -induced hyperconnectivity promotes tau spreading throughout connected brain regions.

Supporting this model, our first finding was that higher regional A β was linked to stronger resting-state fMRI-assessed functional connectivity of posterior temporal, parietal and occipital brain regions to temporal-lobe tau epicenters in 140 patients across the preclinical to clinical AD spectrum. Importantly, this result could be replicated in an independent sample of 345 patients with preclinical AD, suggesting that A β -related connectivity increases are an early event in AD that precedes brain-wide tau deposition. Second, we could confirm that stronger patient-level functional connectivity of tau epicenters to tempo-parietal, frontal and occipital brain regions was associated with faster tau accumulation in these regions, indicative of faster connectivity-mediated tau spread.(2, 28, 29, 71) Thirdly, we tested an integrative pathophysiological disease progression model, showing that the association between temporo-parietal A β deposition and faster tau accumulation in these regions is mediated by A β -driven connectivity increases to the tau epicenter. Together, our results suggest that A β induces neuronal hyperactivity and functional connectivity increases, in line with previous evidence from the cascading network failure model, showing A β -associated connectivity increases as a key feature of AD pathophysiology.(53) Most importantly, we could show that A β -associated connectivity increases facilitate the spreading of tau from epicenters to connected regions in AD. This finding embeds A β -associated connectivity increases as a key link between A β -deposition and tau spreading, thereby rendering A β -induced changes in neuronal activity and connectivity as a potential therapeutic target to attenuate tau spreading and subsequent neurodegeneration and cognitive decline in AD.

hat gelöscht: Further supporting a role of A β in triggering neuronal hyperactivity, we included a small number of post-mortem samples from AD patients and controls, showing a higher proportion of c-Fos+ neurons in AD vs. controls in the visual cortex that typically shows high A β but low tau, indicative of elevated ante-mortem neuronal activity.(71)

1
2 Our first main finding in patients across the AD spectrum showed that regional A β deposition
3 is associated with increased functional connectivity of temporo-parietal and occipital regions
4 to tau epicenters typically located in the temporal lobe. This result pattern was replicated in an
5 independent cohort of preclinical AD patients, in which neocortical tau deposition is typically
6 low,(72) supporting a sequence of events in which A β -related connectivity increases precede
7 and potentially accelerate the spread of tau across the neocortex.(25) Similarly, previous
8 resting-state fMRI studies in A β -transgenic mice and preclinical AD patients reported A β -
9 related functional connectivity increases within and between large scale brain networks,(41,
10 45, 46, 73), as also suggested by the cascading network failure model of AD (52, 53). More
11 recently, a task-fMRI study in a small cohort of cognitively normal older individuals could
12 further show A β -associated hyperconnectivity between the Default Mode Network and the
13 medial temporal lobe during cognitive demands, suggesting that A β -related network changes
14 interfere with neuronal processing very early in the disease process (50). Together, these
15 findings provide converging evidence that A β can induce early-stage functional connectivity
16 increases.

17
18 Consistent across both neuroimaging samples, the A β -associated connectivity increases were
19 found primarily between **temporal lobe** tau epicenters and posterior brain regions which are
20 also physiologically strongly connected to temporal lobe sites of earliest tau deposition.(2)
21 Nevertheless, we also found small but brain-wide associations between A β -deposition and
22 higher connectivity, as well as an association between higher A β and higher FDG-PET in a
23 smaller subset of AD patients, supporting the view that A β is associated with hyperconnectivity
24 and **neuronal** hyperactivity. Congruently, data from rodent models show that A β induces
25 existing synapses to become hyperactive, rather than triggering synaptogenesis,(40, 74)
26 together suggesting that A β leads to an upregulation and strengthened co-activation of pre-
27 existing connections rather than the formation of new connections. From a mechanistic point
28 of view, electrophysiological studies in A β -transgenic mice and AD patients have shown that
29 A β -related hyperactivity and therefore connectivity is explained by a loss of inhibition and an
30 excitatory shift in neuronal activity,(75-77) potentially due to A β -related alterations in Calcium
31 clearance, as well as altered neurotransmitter release/sensitivity and synaptic vesicle
32 release.(42, 43, 75) Since the BOLD signal that is used to determine functional connectivity
33 has been shown to increase as a function of Calcium-dependent neuronal excitation,(78) the

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hat gelöscht: Along the same lines, our exploratory post-mortem data analysis, despite a limited sample size, indicated that AD patients show an increased proportion of c-Fos positive neurons than controls. C-Fos expression is upregulated in response to Calcium influx and typically higher in hyperactive neurons,(75, 76) hence the elevated number of c-Fos positive neurons in the AD brain tissue samples potentially reflect elevated ante-mortem neuronal activity. This finding is in line with previous studies, showing that elevated c-Fos expression in AD patients in the medial temporal lobe, where tau typically emerges in very early AD stages.(77, 78) Importantly, we detected c-Fos positive neurons in the primary visual cortex, i.e. a brain region that typically has high A β but is typically less affected by tau pathology and tau associated neurodegeneration.(20) Supporting this, we did not find differing neuron counts between AD patients and controls in the examined samples, suggesting limited or absent tau-related neurodegeneration. Therefore, bearing in mind the relatively small sample size, our results suggest that the increased c-Fos signal relates to A β -associated processes, in line with animal data showing hyperactive neurons in the vicinity of A β plaques, yet these results require further confirmation in larger post-mortem cohorts.(40)

1 Aβ-related connectivity increases observed in our study may indeed reflect an excitatory shift
2 in neuronal activity ensuing stronger inter-regional synchronicity and connectivity. However,
3 the exact relationships between neuronal activity, Aβ and connectivity between brain regions
4 remain unclear until specifically tested by combining markers of connectivity,
5 electrophysiological activity, and biomarkers of Aβ. Nevertheless, our hypothesis-driven and
6 translational findings provide robust evidence for Aβ to induce increased connectivity of the
7 tau epicenter to connected brain regions.

8 For our second main finding, we could show that stronger patient-specific connectivity of large
9 parts of the temporo-parietal, occipital and frontal cortex to tau epicenters is linked to faster
10 tau accumulation in these brain regions. This finding strongly supports the concept of trans-
11 neuronal tau spreading(79, 80) and robustly recapitulates our previous results, showing that the
12 seed-based connectivity pattern of tau epicenters closely aligns with the spatial pattern of tau
13 accumulation in AD patients.(2, 3, 30) Yet, a key novelty of our study is the combination of
14 longitudinal tau-PET with subject-specific resting-state fMRI connectomics, compared to the
15 previous usage of group average connectivity templates derived from healthy controls.(2, 3,
16 30) More specifically, our findings suggest that Aβ-associated changes of the patient-specific
17 connectome influence the rate of tau spreading in AD.(81) This is closely in line with the results
18 of a previous study in cognitively normal older adults, showing that Aβ-associated
19 hyperconnectivity between the default mode network and the medial temporal lobe during
20 cognitive demands is predictive of faster medial temporal lobe tau accumulation in these early
21 tau vulnerable brain regions (50). Thus, the inter-individual heterogeneity in tau spreading
22 patterns observed in AD and its clinical variants(2, 34, 82) is potentially not only determined
23 by spatial heterogeneous locations of tau epicenters,(2, 29, 82) but also by inter-individual
24 differences in the strength of functional connectivity and in particular by Aβ-induced
25 connectivity changes.(25, 50, 83) Therefore, factors that have been related to altered functional
26 brain network architecture and connectivity strength such as vascular health or
27 neuroinflammation(84, 85) may also alter tau trajectories in AD.

28 Our third finding was that the influence of Aβ on faster tau accumulation is mediated by an
29 Aβ-related functional connectivity increase of the tau epicenter to other brain regions.
30 Importantly, this mediation effect was detected for typical tau vulnerable regions in the
31 temporo-parietal cortex,(20, 21) suggesting that the spreading of tau pathology from epicenters
32 to regions predominantly belonging to Braak stage regions III/IV is specifically facilitated by
33 Aβ-related connectivity increases. From a biological point of view, Aβ-related neuronal

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1 hyperactivity may trigger a stronger activity-dependent secretion of seeding-competent tau
2 species from tau harboring neurons (24, 79) thereby increasing the likelihood that secreted and
3 misfolded tau seeds taken up by a synaptically connected neuron where they induce template-
4 based tau misfolding and aggregation. Supporting this, we have previously reported that
5 elevated p-tau species that are newly synthesized and actively secreted into the CSF (86) drive
6 the spreading of tau across connected brain regions. (30) This interpretation is also supported
7 by previous studies in tau transgenic mice and cell models, showing that i) tau spreads
8 specifically across synaptic connections and that ii) higher neuronal activity (triggered by opto-
9 genetic stimulation of neuronal activity) induces faster neuronal tau secretion and trans-
10 synaptic tau spreading. (24, 79) Overall, these results support a disease model, in which A β -
11 related changes in neuronal activity and connectivity may play a key role in initiating the A β -
12 related spread of tau across connected brain regions. Previous work has further suggested that
13 the initial and relatively slow spreading of tau from the medial temporal lobe to the temporal
14 cortex may be triggered by remote effects of A β in regions connected to initial medial temporal
15 lobe tau epicenters (“pull-effect”), whereas the local convergence of A β and tau in the inferior
16 temporal lobe leads to an acceleration of tau spread from the A β -tau convergence zones to
17 connected brain regions (“push-effect”). (69) It is possible that remote A β in regions not
18 harboring any tau pathology triggers hyperactivity and hyperconnectivity to tau epicenters (50),
19 thereby initiating the spread of local tau seeds from the medial temporal lobe to A β -harboring
20 connected brain regions. Second, the regional convergence of A β and pathological tau seeds
21 may foster local interactions between both pathologies, promoting stronger tau
22 hyperphosphorylation and the generation of tau seeds, which may then lead to an acceleration
23 of tau spreading (87). Therefore, disentangling potential remote and local interactions between
24 A β and tau in the context of connectivity-associated tau spread and neuronal
25 hyperactivity/connectivity will be an important field for future investigation, to better
26 characterize the acceleration phase of A β -dependent tau accumulation and spread.

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28 When interpreting the results of our study, several limitations should be considered. First, we
29 used resting-state fMRI, which is based on the BOLD signal and therefore an indirect proxy
30 measure of neuronal activity. Connectivity is based on the quantification of co-fluctuations of
31 the BOLD signal, hence our neuroimaging results do not directly provide evidence for A β -
32 associated neuronal hyperactivity in AD patients but rather emphasize an increase of
33 synchronicity between two regions harboring A β . However, since preclinical studies have

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1 shown concomitant A β -associated neuronal hyperactivity and hyper-synchronicity **which is**
2 equivalent to the definition of functional connectivity, we deemed resting-state fMRI an
3 adequate measure to assess these changes by proxy.(40) In addition, we preformed exploratory
4 sub-analyses in the ADNI cohort, showing that higher amyloid-PET **are** associated with higher
5 **regional glucose consumption** in a subsample of A β + subjects with available FDG-PET data
6 (n=61), suggesting that higher amyloid **pathology is indeed** linked to higher neuronal activity.
7 **In addition,** it will be a key next step to combine our current analysis framework with
8 electrophysiological measures of neuronal activity **such as EEG or MEG** that allow a more
9 direct quantification of the neuronal excitatory/inhibitory balance, as well as additional markers
10 of neuronal activity or synaptic density. In addition, we employed resting-state fMRI to
11 determine the existence of a connection between any set of brain regions, **whereas it remains**
12 unclear whether a “functional” connection is enabled by a direct underlying “structural” axonal
13 connection along which tau spreads.(81, 88) Rather, a functional connection may reflect both
14 direct and indirect “multi-synaptic” connections.(89) Addressing this limitation by including
15 individual structural connectomes would have required high-quality diffusion MRI to
16 determine subject-level structural connectivity which would have drastically limited the sample
17 size. Thus, we refrained from additionally including structural connectivity to constrain
18 functional connections in the current study. Lastly, the cross validation in the A4 sample was
19 restricted to cross-sectional analyses on A β vs. connectivity changes, since longitudinal tau-
20 PET data are not yet available in this dataset. Therefore, the longitudinal associations between
21 A β -related connectivity changes and faster tau spreading remain to be independently replicated
22 once these data become available. **Beyond the limitations, our** study features several clear
23 strengths. **as** we employed a fully hypothesis-driven approach to determine the role of
24 connectivity changes in cross-linking A β accumulation and tau spreading. **In addition, the**
25 study design **included** an independent replication sample illustrated the robustness of the
26 association between A β and connectivity increases.

27
28 In conclusion, our findings provide evidence that A β -associated neuronal activity and
29 connectivity changes may be a key missing link between the accumulation of A β and the
30 subsequent spreading of tau pathology in AD. These findings are of high clinical importance,
31 since modulating neuronal activity may be a promising target for attenuating A β -related tau
32 accumulation and spreading. Previous trials repurposing anti-epileptic drugs to lower neuronal
33 hyperexcitability and hyperactivity in AD patients have been tested, yet over relatively short

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hat gelöscht: In addition, we implemented a translational study design, including exploratory post-mortem analyses which further substantiated the notion that A β is related to neuronal hyperactivity using issue samples from AD patient brains. ...

1 timespans of several weeks, focusing on clinical endpoints (56, 58, 90). However, we reason
2 that surrogate biomarker endpoints should be considered, such as the activity-dependent
3 secretion of phosphorylated tau species detectable in CSF or plasma (86), or the long-term
4 spreading and accumulation of tau pathology, which would require longer intervention periods
5 in subjects at the early phase of A β -induced tauopathy. Such a treatment approach may be
6 particularly promising in early-stage AD patients in addition to targeting A β , to target fibrillar
7 A β deposition directly, as well as its consequences on tau spreading, to therefore maximize the
8 likelihood to attenuate tau-related neurodegeneration and cognitive decline.(13, 14, 91)
9 Together, our results should encourage others to further investigate neuronal activity and
10 connectivity as key links between A β and tau, to help specifically target the A β -tau axis in AD.

11

12

1 Material and METHODS

2 Study design

3 The overall objective of this observational study was to determine whether amyloid deposition
4 induces neuronal hyperconnectivity and hyperactivity, thereby driving tau spreading across
5 interconnected brain regions. To this end, we used observational neuroimaging and biomarker
6 data from the ADNI and A4 study for independent replication. No treatments or interventions
7 were administered for the included individuals and data and no randomization was performed.
8 We included the largest possible sample size that matched the inclusion criteria specified
9 below, hence no a priori study sized calculations were performed. The authors were not blinded
10 to the diagnosis or biomarker status of the included individuals.

11

12 **ADNI Participants**

13 209 Subjects were included from the Alzheimer's disease Neuroimaging Initiative (ADNI)
14 database, based on availability of clinical data, baseline 18F-Florbetapir/Florbetaben amyloid-
15 PET, 3T resting-state fMRI and longitudinal 18F-Flortaucipir tau-PET data. All baseline data
16 had to be obtained within a timeframe of 6 months. Participants diagnostic status was
17 determined by ADNI as cognitively normal (CN; Mini Mental State Examination [MMSE]≥24,
18 Clinical Dementia Rating [CDR]=0, non-depressed), mildly cognitively impaired (MCI;
19 MMSE≥24, CDR=0.5, objective memory-impairment on education-adjusted Wechsler
20 Memory Scale II, preserved activities of daily living) and demented (MMSE=20–26,
21 CDR>0.5, National Institute of Neurological and Communicative Disorders and
22 Stroke/Alzheimer's Disease and Related Disorders Association criteria for probable AD).
23 Amyloid status was determined on global amyloid-PET SUVRs using tracer-specific cut-offs
24 at 1.11/1.08 for Florbetapir/Florbetaben, as previously established in the ADNI cohort (92, 93)
25 All study procedures were conducted in accordance with the declaration of Helsinki, ethical
26 approval was obtained by ADNI investigators. All study participants provided written informed
27 consent.

28

29 **A4 participants**

30 For replication of cross-sectional analyses, we included baseline data from 400 participants of
31 the A4 study, based on availability of clinical data, 18F-Florbetapir amyloid-PET, resting-state
32 fMRI and [18F-Flortaucipir tau-PET obtained at the baseline study visit. All subjects were
33 classified as CN (MMSE>25, CDR=0, Wechsler Logical Memory score of 6 to 18), as defined

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1 by the inclusion criteria of the A4 trial (<https://clinicaltrials.gov/study/NCT02008357>).
2 Amyloid status was determined using global amyloid-PET SUVRs and the 1.11 global SUVR
3 cut-off defined in the ADNI cohort. All study procedures were conducted in accordance with
4 the declaration of Helsinki, ethical approval was obtained by A4 investigators. All study
5 participants provided written informed consent.

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7 *Neuroimaging acquisition*

8 Structural and functional MRI were acquired using 3T Siemens, GE and Philips scanners. T1-
9 weighted structural scans were collected using an MPRAGE sequence (TR=2300ms; Voxel
10 size=1x1x1mm; for parameter details see: [https://adni.loni.usc.edu/wp-](https://adni.loni.usc.edu/wp-content/uploads/2017/07/ADNI3-MRI-protocols.pdf)
11 [content/uploads/2017/07/ADNI3-MRI-protocols.pdf](https://adni.loni.usc.edu/wp-content/uploads/2017/07/ADNI3-MRI-protocols.pdf)). Resting-state-fMRI was obtained using
12 a 3D echo-planar imaging (EPI) sequence with 200 fMRI volumes per subject
13 (TR/TE=3000/30ms; flip angle=90°; Voxel size=3.4x3.4x3.4mm).

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14 PET data was assessed post intravenous injection of ¹⁸F-labeled tracers (Flortaucipir: 6x5min
15 time-frames, 75-105min post-injection; Florbetapir: 4x5min time-frames, 50-70min post-
16 injection; Florbetaben: 4x5min time-frames, 90-110min post-injection; for more information
17 see <http://adni.loni.usc.edu/methods/pet-analysis-method/pet-analysis/>). In a subset of ADNI
18 participants, we also included FDG-PET data, recorded 6x5min frames 30-60min post-
19 injection. A4 and ADNI imaging data were recorded using congruent imaging protocols.

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21 *Image processing*

22 All images were screened for artifacts before preprocessing, and processing was conducted
23 independently for [the ADNI and A4 sample](#). T1-weighted structural MRI scans were bias-
24 corrected, segmented, and non-linearly warped to Montreal Neurological Institute (MNI) space
25 using the CAT12 toolbox (<https://neuro-jena.github.io/cat12-help/>). Dynamically acquired
26 PET images were realigned and averaged to obtain single Flortaucipir/Florbetapir images
27 which were rigidly registered to the T1-weighted MRI scan. [As reference regions, we used the](#)
28 [inferior cerebellar grey for Flortaucipir, the whole cerebellum for Florbetapir/Florbetaben, and](#)
29 [the Pons for FDG-PET \(94, 95\) Reference regions](#) and the cortical Schaefer atlas including 200
30 regions of interest (ROIs) were warped from MNI to T1-native space using the CAT12-derived
31 non-linear normalization parameters, masked with subject-specific grey matter and applied to
32 PET data to determine standardized uptake value ratios (SUVRs) for each region of the
33 Schaefer 200 atlas. (62) Global and regional Florbetapir/Florbetaben SUVRs were converted to
34 centiloid using equations provided by ADNI. To determine longitudinal tau-PET change, we

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1 employed linear mixed models controlling for random slope and intercept as described
2 previously.(2, 3)
3 Resting-state fMRI images were slice-time corrected and realigned to the first volume for
4 motion correction, followed by co-registration to the respective T1-weighted images. Using
5 rigid-transformation parameters, T1-derived grey-matter, eroded white matter and eroded
6 cerebrospinal fluid (CSF) segments were transformed to EPI space. To denoise EPI images,
7 we regressed out nuisance covariates, including the timeseries of the eroded white matter and
8 eroded CSF plus six motion parameters as well as their time and dispersion derivatives,
9 followed by detrending and band-pass filtering (0.01-0.08Hz) in native space. To further reduce
10 movement artifacts which may compromise connectivity assessment, (96) we performed
11 motion scrubbing in which volumes exceeding a 0.5mm frame-wise displacement threshold
12 were removed, as well as one prior and two subsequent volumes. All included subjects had at
13 least five minutes of resting-state fMRI remaining after scrubbing (97). Spatial smoothing was
14 not carried out to avoid artificially enhancing functional connectivity caused by signal spilling
15 between adjacent brain regions. Pre-processed resting-state-fMRI images were subsequently
16 warped to MNI space using the CAT12-derived spatial normalization parameters.

18 *Assessment of tau epicenters and functional connectivity*

19 Tau epicenters were determined for each subject as 5% of the 200 cortical Schaefer ROIs with
20 highest baseline tau-PET SUVRs which is equivalent to 10 ROIs per individual. Subject-
21 specific functional connectivity matrices were determined across the 200 ROIs of the Schaefer
22 atlas as Fisher-z-transformed Pearson moment correlations between ROI-specific time-series.
23 Negative correlations were eliminated, and autocorrelations were set to zero. Subject-specific
24 tau epicenter masks were applied to the subject-specific functional connectivity matrices to
25 determine average seed-based connectivity patterns of the tau epicenter to the rest of the brain.
26 The tau epicenter to ROI connectivity per subject can be found in supplementary data files
27 S1&S2 for ADNI and A4.

29 *Statistics*

30 Sample demographics for ADNI and A4 were compared using ANOVAs for continuous
31 variables and Chi-squared tests for categorical variables. To determine the effect of A β on
32 seed-based connectivity of the tau epicenter, we used linear regressions to determine the effect
33 of regional A β (defined as the average centiloid across the tau epicenter and the target ROI) on

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1 epicenter connectivity, controlling for age, sex, ethnicity, MRI scanner type, diagnosis and
2 framewise displacement. This analysis was conducted for ADNI and A4. ROI-wise p-values
3 were FDR-corrected for multiple comparisons. In ADNI, we also applied a t-test against zero
4 on the overall distribution of beta values, to determine whether there is a global association
5 between A β and increased epicenter connectivity.

6 Next, we tested whether higher epicenter connectivity is linked to faster tau accumulation in
7 the ADNI dataset with available longitudinal tau-PET data. To this end, we performed linear
8 regression per ROI, testing the effect of tau epicenter connectivity at baseline on longitudinal
9 tau-PET change rates, controlling for age, sex, ethnicity, diagnosis, MRI scanner type and
10 average motion during the resting-state fMRI scan **defined as the** mean framewise
11 displacement. ROI-wise p-values were FDR-corrected for multiple comparisons. Again, we
12 applied a one-sample t-test to the beta-values to determine the global pattern of associations.
13 Lastly, we employed bootstrapped mediation analyses with 1000 iterations per ROI, to assess
14 whether the effect of A β on tau accumulation rates was mediated by stronger connectivity to
15 the tau epicenter. Mediation models were again controlled for age, sex, ethnicity, diagnosis,
16 MRI scanner type and average motion during the resting-state fMRI scan. All analyses were
17 conducted for epicenter connectivity to all cortical ROIs and resulting beta and p-values were
18 FDR-corrected for multiple comparisons. All statistical analyses were conducted using R
19 statistical software (Version 4.0.4). All plots have been created using ggplot2. Error bars in
20 scatterplots indicate 95% confidence intervals of regression lines. In boxplots, the boxes
21 indicate median \pm interquartile range, whiskers extend to values ± 2.5 of the interquartile range
22 from the median.

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7 [content/uploads/how_to_apply/ADNI_Acknowledgement_List.pdf](http://adni.loni.usc.edu/wp-content/uploads/how_to_apply/ADNI_Acknowledgement_List.pdf).

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20 manuscript

21

22 **Competing interests:** None

23

24 **Data and materials availability:** ADNI and A4 data are publicly available after registration
25 ~~on the study websites (ADNI: <https://adni.loni.usc.edu/about/adni4/>; A4:~~
26 ~~<https://www.a4studydata.org>), Subject-level connectivity of tau epicenters can be found in the~~
27 ~~supplementary.~~

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Kommentiert [DN82]: Please make sure that all authors' contributions are accounted for, with every author's initials appearing at least once. If authors share initials, please use first initial and full surname. Where possible, please indicate which authors performed each experiment. Please provide author contributions in narrative form using sentences rather than list.

Kommentiert [DN83]: All consulting, whether paid or unpaid, and whether related to the present work or not, needs to be declared here. Any patents related to this work need to be stated here (cite patent title and filing #).

Kommentiert [I84R83]: Can this be generated from the COI forms? I do not have access to the forms of the individual authors, and each of them must have indicated this there

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Table 1: Neuroimaging sample characteristics

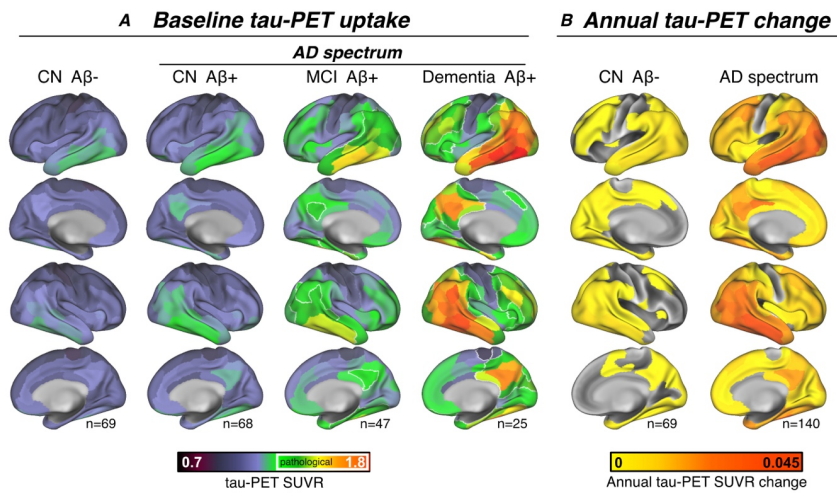
ADNI (n=209)	CN Aβ- (n=69)	CN Aβ+ (n=68)	MCI Aβ+ (n=47)	Dementia Aβ+ (n=25)	p-value
Age	71.7±6.16 ⁴	73.6±6.22	74.4±6.12	75.8±8.02 ¹	0.027
Sex (f/m)	38/31	46/22	21/26	7/18	0.004
MMSE	29.0±1.08 ^{3,4}	29.1±1.30 ^{3,4}	27.4±1.91 ^{1,2,4}	23.4±3.49 ^{1,2,3}	0.001
ADAS13 total	7.67±4.36 ^{3,4}	8.25±4.73 ^{3,4}	16.7±6.49 ^{1,2,4}	27.8±8.63 ^{1,2,3}	0.001
Centiloid	7.94±7.37 ^{2,3,4}	64.3±35.3 ^{1,4}	77.4±32.5 ^{1,4}	96.0±30.1 ^{1,2,3}	0.001
Amyloid tracer (FBB/FMM)	52/17	45/23	29/18	12/13	0.873
Tau-PET global SUVR	1.06±0.080 ^{3,4}	1.09±0.077 ^{3,4}	1.19±0.212 ^{1,2,4}	1.35±0.399 ^{1,2,3}	0.001
Tau-PET follow-up years	3.42±1.12 ^{2,3,4}	2.80±1.21 ^{1,3,4}	2.21±1.02 ^{1,2}	1.68±0.60 ^{1,2}	0.001
Number of tau- PET visits	2.39±0.65	2.68±0.74	2.66±0.67	2.40±0.50	0.034
Mean Framewise Displacement in resting-state fMRI	0.099±0.059	0.128±0.074	0.117±0.082	0.137±0.068	0.043
A4 (n=400)	CN Aβ- (n=55)	CN Aβ+ (n=345)			p-value
Age	70.0±4.15	72.2±4.91			<0.001
Sex (m/f)	28/27	150/195			0.377
Centiloid	11.1±7.82	69.5±31.4			<0.001
Global tau-PET SUVR	1.06±0.076	1.10±0.08			<0.001
MMSE	28.8±1.25	28.6±1.13			0.204
Mean Framewise Displacement in resting-state fMRI	0.106±0.067	0.110±0.066			0.666

Post-Hoc Tukey Test: ¹p<0.05 vs. CN Aβ-, ²p<0.05 vs. CN Aβ+, ³p<0.05 vs. MCI Aβ+, ⁴p<0.05 vs. Dementia Aβ+. Means plus/minus standard deviations are displayed for continuous measures. Absolute numbers are displayed for categorical measures.

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Figure legends:

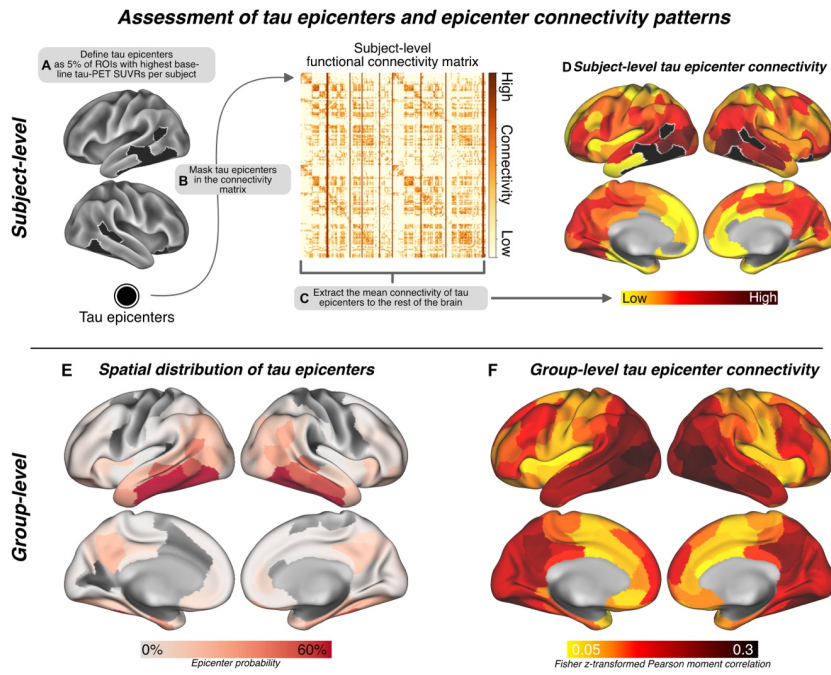
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Figure 1: (A) Baseline tau-PET SUVRs values stratified by diagnostic group and Aβ status, showing increasing tau load across the AD spectrum. SUVRs surpassing an abnormal cut-off of 1.3 are outlined in white. (B) Annual tau-PET SUVR change rates, as calculated using ROI-wise linear mixed models, stratified by amyloid status, illustrating faster tau accumulation in Aβ+ subjects, particularly in temporoparietal brain regions.

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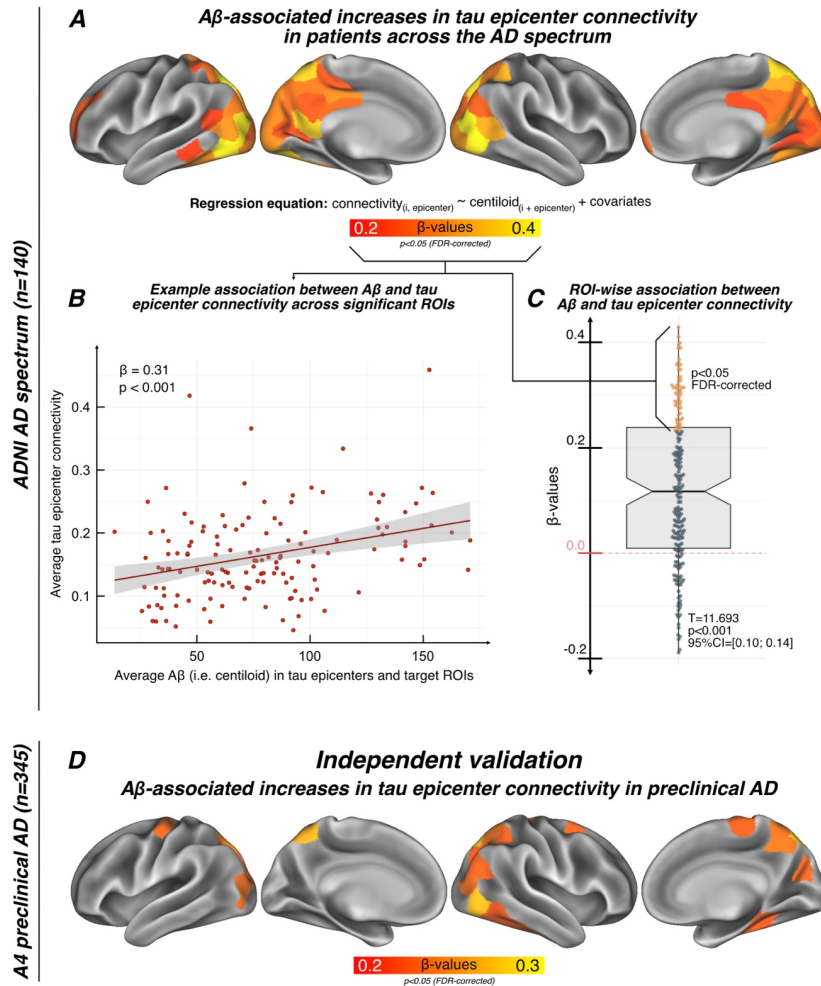
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3 **Figure 2:** (A) For each participant, 5% of brain regions with the highest baseline tau-PET SUVRs were
4 defined as tau epicenters. (B) Epicenter masks were applied to (C) subject-specific connectivity
5 matrices to (D) extract epicenter connectivity patterns. (E) Mapping of group-average epicenter
6 probability and (F) epicenter connectivity.

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3 **Figure 3:** (A) Association between A β and epicenter connectivity in ADNI. (B) Average association4 between A β and connectivity for significant ROIs ($p < 0.05$, FDR-corrected) displayed in panel (A).

5 (C) Unthresholded distribution of beta values displayed in panel (A) showing that beta-values are

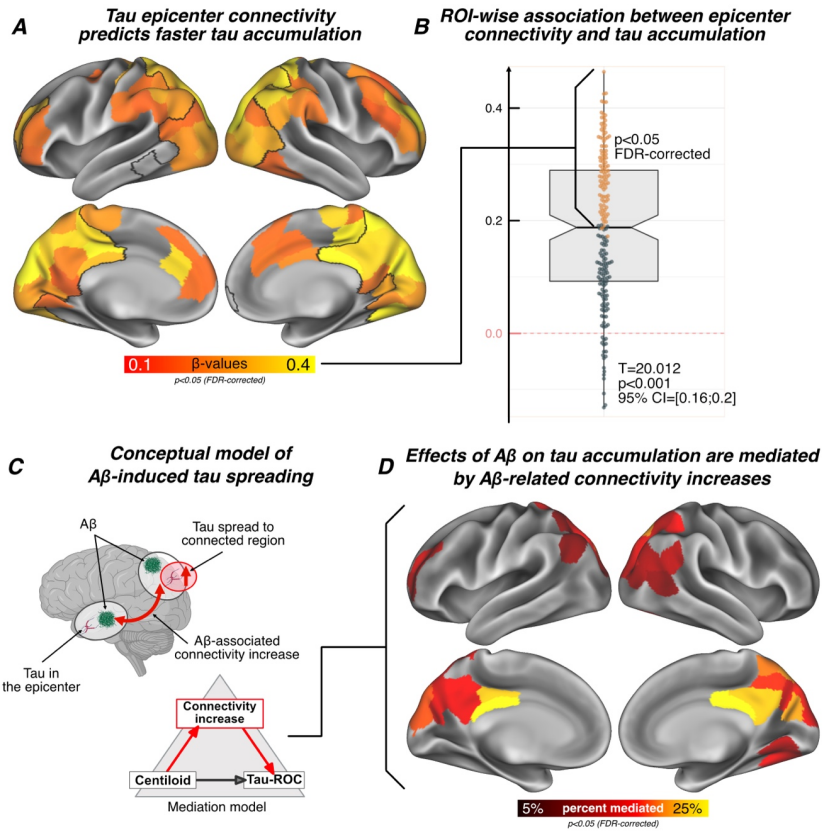
6 overall significantly larger than zero ($p < 0.001$). (D) Validation of the association between A β and7 epicenter connectivity in A4. Regression-based associations were corrected for age, sex, ethnicity,8 scanner type, diagnosis and motion during the fMRI scan.

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Kommentiert [DN89]: Please use this word only to report statistical significance and always accompanied by a p value directly in the text, for example: ($p=x.xx$) or ($p<x.xx$). Any other uses of this word should be removed or replaced throughout the text.

Kommentiert [DN90]: Please use this word only to report statistical significance and always accompanied by a p value directly in the text, for example: ($p=x.xx$) or ($p<x.xx$). Any other uses of this word should be removed or replaced throughout the text.

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Figure 4: (A) Association between stronger epicenter connectivity and faster tau accumulation in ADNI, including (B) the unthresholded distribution of beta-values, showing an overall positive association between epicenter connectivity and faster tau accumulation. (C) Conceptual model of Aβ-related tau spreading and (D) bootstrapped mediation showing that effects of Aβ on tau accumulation are mediated by increased epicenter connectivity. *Regression-based associations and mediations were corrected for age, sex, ethnicity, scanner type, diagnosis and motion during the fMRI scan.*

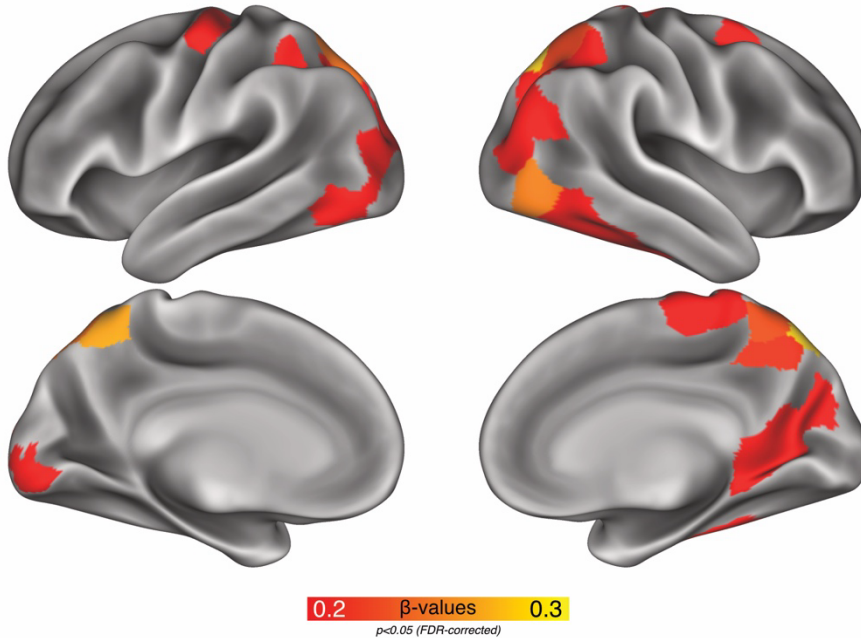
hat gelöscht: **Figure 4:** Post-mortem analyses of AD vs. control (CTRL) brains. (A) Overview of an example tissue staining. The red rectangle on the brain surface rendering highlights the anatomical location of the extracted probe in the primary visual cortex. The red rectangle on the microscopic image illustrates the location of the zoom-in images displayed in panel B. (B) Merged images of Dapi, NeuN, Aβ and c-Fos in an example control and AD subject. Group differences in (C) Aβ-plaque area, (D) neuron count (i.e. NeuN), and (E) the proportion of neurons with c-Fos positive signal. [5]

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Supplementary Figures

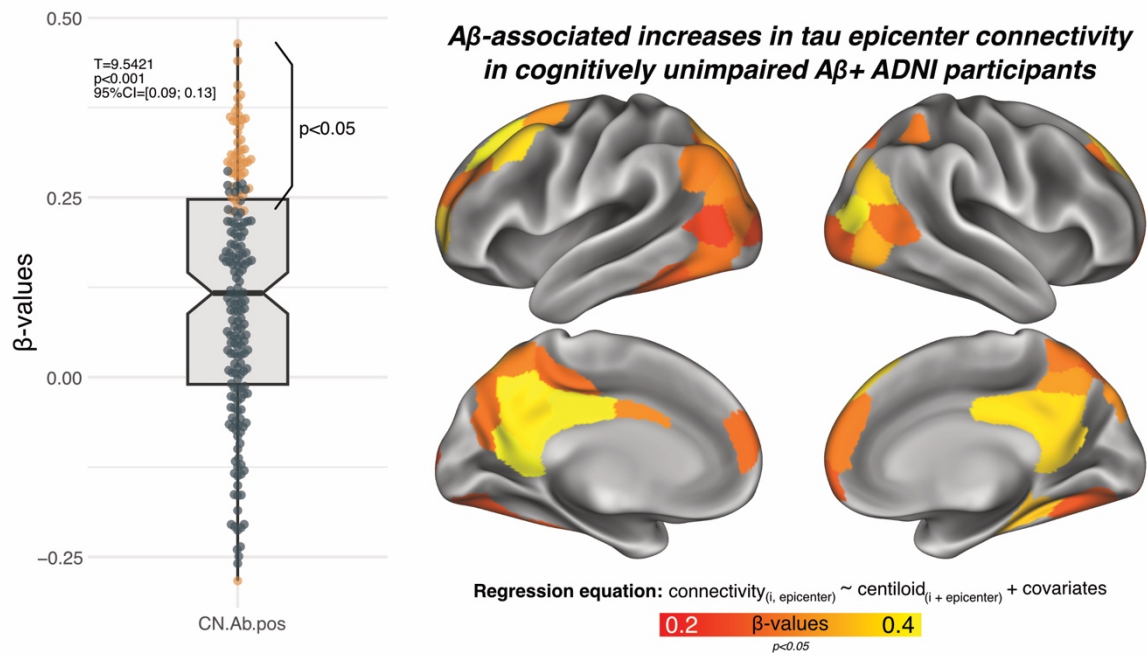
Fig. S1:

A β -associated increases in tau epicenter connectivity in cognitively unimpaired A β + A4 participants



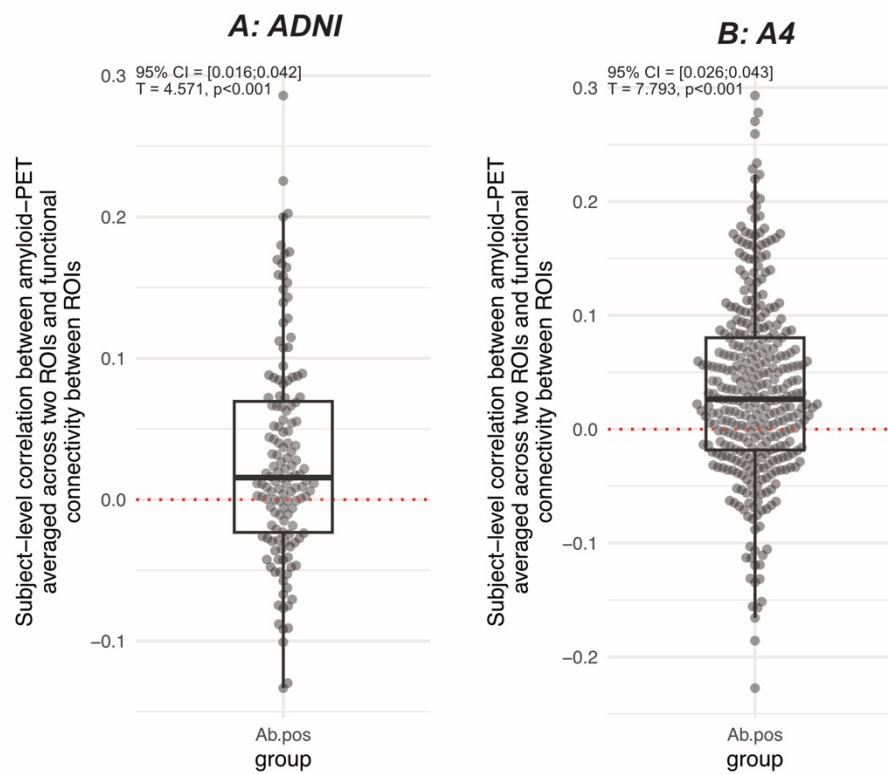
Exploratory sub-analysis showing the association between A β and epicenter connectivity in cognitively unimpaired A β + subjects of the A4 cohort, reclassified as A β + at a global amyloid-PET SUVR threshold of 1.15 ($n=322$) as recommended by the A4 imaging core. The surface rendering shows the spatial pattern of significant ($p < 0.05$, FDR-corrected) associations between amyloid-PET levels in the tau epicenter and target ROI and the connectivity between the epicenter and the target ROI.

Fig. S2:



Exploratory sub-analysis showing the association between $A\beta$ and epicenter connectivity in cognitively unimpaired $A\beta^+$ subjects of the ADNI cohort (n=68). The surface rendering shows the spatial pattern of significant ($p < 0.05$) associations between amyloid-PET levels in the tau epicenter and target ROI and the connectivity between the epicenter and the target ROI. The beeswarm plot shows that the overall distribution of beta values is significantly greater than zero as shown by a one sample t-test, with 95% Confidence intervals not overlapping with zero. Yellow dots in the beeswarm plot reflect significant ROIs projected on the brain surface.

Fig. S3:



Beeswarm plots, illustrating the distribution of subject-level correlation coefficients between amyloid-PET summarized across any given ROI pair and subject-level functional connectivity matrices for $A\beta^+$ subjects of ADNI and A4. Results show an overall positive distribution of correlation coefficients as indicated by a significant t-test against zero, and 95% confidence intervals of the distributions not including zero. These data suggest that ROI pairs with a combined high amyloid load tend to have higher functional connectivity within individuals.