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Post-mortem AT-8 reactive tau species correlate with non-

² plaque Aβ levels in the frontal cortex of non-AD and AD

3 brains

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28 Abstract

29 The amyloid cascade hypothesis states that A β and its aggregates induce pathological changes in tau, leading to formation of neurofibrillary tangles (NFTs) and cell death. A caveat with this hypothesis is 30 31 the temporo-spatial divide between plaques and NFTs. This has been addressed by the inclusion of 32 soluble species of A^β and tau in the revised amyloid cascade hypothesis, however, the demonstration 33 of a correlative relationship between AB and tau burden in post-mortem human tissue has remained 34 elusive. Employing frozen and fixed frontal cortex grey and associated white matter tissue from non-AD controls (Con; n=39) and Alzheimer's diseases (AD) cases (n=21), biochemical and 35 36 immunohistochemical measures of A β and AT-8 phosphorylated tau were assessed. Native-state dot-37 blot from crude tissue lysates demonstrated robust correlations between intraregional AB and AT-8 38 tau, such increases in A β immunoreactivity conferred increases in AT-8 immunoreactivity, both when 39 considered across the entire cohort as well as separately in Con and AD cases. In contrast, no such 40 association between AB plaques and AT-8 were reported when using immunohistochemical 41 measurements. However, when using the non-amyloid precursor protein cross reactive MOAB-2, 42 antibody to measure intracellular A β within a subset of cases, a similar correlative relationship with 43 AT-8 tau as that observed in biochemical analysis was observed. Collectively our data suggests that accumulating intracellular A β may influence AT-8 pathology. Despite the markedly lower levels of 44 45 phospho-tau in non-AD controls correlative relationships between AT-8 phospho-tau and AB as 46 measured in both biochemical and immunohistochemical assays were more robust in non-AD controls, suggesting a physiological association of A^β production and tau phosphorylation, at least 47 within the frontal cortex. Such interactions between regional A β load and phospho-tau load may 48 49 become modified with disease potentially, as a consequence of interregional tau seed propagation, 50 and thus may diminish the linear relationship observed between AB and phospho-tau in non-AD 51 controls. This study provides evidence supportive of the revised amyloid cascade hypothesis, and demonstrates an associative relationship between AT-8 tau pathology and intracellular Aß but not 52 extracellular Aβ plaques. 53

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60 Introduction

61 The original amyloid cascade hypothesis stated that the extracellular deposition of insoluble beta-62 amyloid (A β) plaques drives intracellular tau phosphorylation, the formation of neurofibrillary tangles (NFTs), and the subsequent neurodegeneration which underlies the pathology of Alzheimer's disease 63 (AD) [1]. Owing to the lack of correlation between plaque burden and cognitive impairment, as well 64 as a growing understanding of the toxicity of fibrillar and pre-fibrillar intermediate species, the 65 66 hypothesis has been revised to include roles for A β oligomers and tau oligomers[2]. Whilst the recent 67 outcomes of plaque clearing and A β oligomer targeted immunotherapies[3-5] support this revised 68 amyloid cascade hypothesis, several inconsistencies relating to the interaction of A β and tau remain.

Foremost, is the spatio-temporal disconnect between the emergence and progression of Aβ plaque and tau NFT pathology. Based on the post-mortem neuropathological Thal phases of Aβ deposition and positron emission tomography (PET) imaging studies Aβ plaques originate within the neocortex, specifically within the orbito-frontal and medial parietal cortices, before spreading to the hippocampus, the brain stem and cerebellum[6, 7]. In contrast, as reflected by Braak NFT staging, tau pathology initially occurs in the entorhinal cortex and hippocampus and subsequently spreads to the lateral temporal and parietal cortices and finally to the frontal and occipital cortices[8, 9].

76 Cross sectional population studies further highlight the independent nature of the two hallmark 77 pathologies, reporting that tau pathology consistent with Braak Stages I-II, occurs more readily with 78 age than that of plaque depositions[10]. Consequently, $A\beta$ deposition is not a prerequisite for NFT 79 formation in aging, nor in the cases of primary tauopathies[11]. The independence of tau pathology 80 from Aβ plaques is particularly evident in cases of primary age-related tauopathies (PART), which present with AD related NFTs in a spatial pattern consistent with Braak NFT stages up to stage IV 81 82 without A β plaque deposition[12]. Moreover, the demonstration of prion-like spreading via tau seed 83 templating and pathology propagation, provides a mechanistic process by which the presence of tau 84 pathology may occur independent from the influence of A β plaques[13]. Such tau seed propagation 85 of pathology likely contributes to the progression of tau pathology in many tauopathies, including AD.

Taken together, the direct causation of NFTs, purely as a consequence of Aβ plaque burden is difficult
to ratify, with the differential emergence in time and space of the neuropathological hallmarks, as well
as the independent occurrence of tau aggregations in other neurodegenerative conditions.

However, it remains likely that plaque deposition or rather the process of amyloid plaque formation,
influences the generation of tau pathology. This is perhaps most strongly supported by numerous
biochemical studies of human brain tissue, which report robust correlations with pathological Aβ and

tau species[14-17]. Despite the close relationship between tau and Aβ levels in various biochemical
assays, histochemical approaches frequently fail to detect such correlations. The disconnect between
biochemical and histochemical analysis clearly highlights differences in the pathological species
measured within the different methodological approaches.

In line with the revised amyloid cascade hypothesis[2], a range of experimental models demonstrate 96 97 that soluble pathological species of both tau and A β exert toxic influence within the brain, evident in 98 injection models [18-21], familial AD (FAD) [22-24], tauopathy[25-27] mouse models and cell culture approaches[28, 29]. Whilst APP centric FAD mouse models, do not develop NFTs, there is clear 99 100 evidence of increased tau phosphorylation within these models. Moreover in multi-genic mice, in 101 which a human mutant tau gene is included, APP centric mutations can accelerate and enhance NFT 102 pathology[30]. Nevertheless, evidence for the induction of tau pathology following Aβ intracerebral 103 delivery is sparse in non-transgenic animals[31, 32] and even in tau transgenic mice[33].

Whilst there are several possible explanations for the failure of exogenous Aβ to drive tau pathology
in-vivo, one possible contributing factor is that the sole delivery of Aβ to the extracellular space may
not be sufficient to drive tau pathology. Indeed, a growing body of evidence suggests that intracellular
Aβ accumulation may also play a role in the pathobiology of AD[34], influencing cellular
dysfunction[35-37] and tau phosphorylation[22].

109 Despite the potential for non-plaque A β to contribute to the production of tau pathology, few studies 110 have sought to examine both biochemical and immunohistochemical quantification of AB and tau 111 pathology within the same cases, thus, allowing for a direct comparison between the correlative 112 strength of total A β , plaque A β and intracellular A β with tau pathology. This current study aimed to quantify such parameters across cases to establish potential correlative relationships between 113 114 pathological species in order to further understand the influence that $A\beta$, in various forms, may have 115 upon the regional generation of tau pathology, both in neuropathologically confirmed cases of AD as 116 well as non-AD control cases.

117

118 Methods

119 Human post-mortem brain tissue

A study cohort of post-mortem human brains from clinico-pathologically classified AD (n=21) and non neurodegenerative control cases (Con, n=39) was obtained from the Newcastle Brain Tissue Resource
 (NBTR). AD Subjects had been clinically assessed during life prior to brain tissue donation and
 diagnosed with dementia due AD. Control cases similarly had been assessed during life and at the time

124 of death did not have dementia. The final clinico-pathological diagnoses were established by 125 combining clinical neuropathological data reviewed at regular meetings involving JA and AT. Neuropathological diagnoses were based on assessment of brain tissue according to the National 126 127 Institute of Ageing – Alzheimer's Association (NIA-AA) criteria[38], including Braak NFT staging[39], 128 Thal phases [6], and Consortium to Establish a Registry for Alzheimer's Disease (CERAD) scoring[40], 129 as well as Braak LB stages[41], and Newcastle / McKeith Criteria [42, 43] (Table 1 and S1 for full 130 details). 131 For histology and tissue micro-array (TMA; see below), tissue sections were prepared from the right

hemisphere of the brain and fixed for 4-6 weeks in 4% paraformaldehyde. Corresponding frozen frontal grey (GM) and white matter (WM) tissue (Brodmann's area (BA) 9) was obtained from the left hemisphere, dissected in a coronal plane and snap frozen between copper plates at -120°C prior to being stored at -80°C. Due to limitations in tissue availability, it was not possible to obtain both fixed and frozen tissue for all cases (see Table 1 and S1 for full details). Comparative analysis of age and post-mortem interval (PMI) between disease groups determined no significant difference in either measure (p>0.05).

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140 Tissue lysis

~250mg of frozen frontal tissue was electronically homogenised 1:10 (W/V) in 0.2M tetraethyl
ammonium bicarbonate (TEAB, pH 7.2, Sigma) with 1% SDS, containing protease (1 per 10ml,
Complete , Roche) and phosphatase inhibitors (1 per 10mls, PhosSTOP, sigma) using an Ultra-turrax
T10 homogeniser (5 mm diameter probe; 30,000rpm) for 15 sec. Lysate were aliquoted and stored at
-80°C, prior to use.

146

147 Immunoblot quantification of AD markers:

148 Dot blots were conducted for total A β and AT-8 phospho-tau in both grey and white matter samples. 149 The protein concentration of GM and WM crude lysate were adjusted to 0.5µg/µl as per Bradford assay 150 and dotted directly to a nitrocellulose membrane at 10μ l (5μ g/dot) and left to dry for 20 minutes 151 before further processing. The membranes, briefly washed in Tris-buffered saline (TBS; in mM; 50 152 Trizma base, 150 NaCl, pH= 7.6) prior to being blocked in 5% milk powder containing Tris-Buffered 153 Saline with 0.1% Tween 20 (TBST) at room temperature for 1 hour. After blocking, blots were rinsed 154 in TBS washing buffer 3 times for 5 minutes each. Membranes where subsequently placed in primary 155 antibody solution (TBST, 5% bovine serum albumin and 0.05% sodium-Azide) containing either MOAB-

156 2 (1:1000, Cat# M-1586-100, Biosensis) for the detection of Aβ or AT-8 (1:1000, Cat# AB_223647, Thermofisher) for phospho-tau and incubated overnight at 4°C. The membranes were then washed 157 158 in TBST before being incubated for 1 hour at room temperature in horse radish peroxidase conjugated 159 goat anti-mouse secondary IgG antibody (TBST+5% milk powder+1:5000 dilution) prior to repeated 160 washing before being development. Immunoreactivity was visualized via enhanced 161 chemiluminescence (1.25 mM Luminol, 25µl of 3%H202 and 50µl coumaric acid was incubated for 1 162 minute). The signal was captured by using digital western blot camera, with high sensitivity and auto 163 exposure being selected. The images were saved as 8- bit for illustration and 16-bit for quantification. 164 Total protein loading was determined via Ponceau S general protein stain (0.1% Ponceau S (w/v) and 165 5.0% acetic acid (w/v) in ddH₂O water) and resulting loading staining captured.

166 Immuno-blot quantification

167 Immunoreactivity of enhanced chemiluminescent (ECL) luminol blots and Ponceau S-stained blots was 168 quantified from 16-bit digitized images based on area under the curve measurements as computed by 169 ImageJ (Ver 1.53e, NIH, USA). Normalization of immunoblot intensity values were then performed 170 using total protein adjusted values. The 52 samples of human frontal cortex GM and WM were 171 processed in 4 separate batches and each batch normalized to the mean value of control cases (each 172 blot containing >3 Br 0-IV control cases) prior to pooling values between blots.

173

174 Immunohistochemical quantification of Aβ plaques and phospho-tau (AT-8)

175 Regional quantification of the A β plaque and AT-8 phospho-tau load within the frontal cortex (BA9) 176 via TMA slides, as described previously[44]. Sections (6 µm, thick) were cut from paraffin embedded 177 TMA blocks tissue blocks comprising of cylindrical tissue cores taken from multiple brain region 178 specific blocks and mounted on glass slides. Slides containing a 3 mm diameter samples of BA9 frontal 179 cortex were baked at 60°C for 1hr prior to being dewaxed in xylene, rehydrated in descending 180 concentration of ethanol (5 mins immersion) and washed in TBS. Slides intended for phospho-tau staining were treated with microwave assisted antigen retrieval (800 W, 10 mins) in citrate buffer (10 181 182 mM Citric acid, 0.05 % Tween 20, pH 6) and those intended for Aβ plaque staining were submerged in 183 90% Formic acid for 1hr at RT, before endogenous peroxidases were quenched in H_2O_2 (3%, 20 mins 184 submersion). Following consecutive washes in TBS and TBST, slides were incubated with either mouse 4G8 (1:16000, Cat# SIG-39200, Covance) or anti-AT-8 (1:4000) in TBS for 1 hr and immunoreactivity 185 186 visualised via the MENAPATH HRP polymer detection kit (Menarini diagnostics, Wokingham, UK) and 187 3,3'- Diaminobenzidine (DAB) chromogen with appropriate TBS and TBST washes performed between steps. Slides were co-stained with haematoxylin prior to being dehydrated in ethanol, cleared inxylene and mounted in dibutylphthalate polystyrene xylene (DPX).

Stained BA9 frontal cortex samples were imaged at x100 magnification with a semi-automated microscope (Nikon Eclipse 90i microscope, DsFi1 camera and NIS elements software V 3.0, Nikon). For each case, multiple images were captured to form a 3x3 image grid with 15% overlap in adjacent images, such that an area of 1.7mm was sampled from each case.

Following visual quality control inspection and the application of regions of interest (ROI) to exclude areas of tissue folds and tears, a consistent restriction threshold for 4G8 (R50-180, G20-168, and B8-139) and AT-8 (R25-170, G27-156, B11-126) was applied producing a binary signal image from which the percentage area of immunoreactivity could be acquired. For the quantification of A β plaques, 4G8 images were further processed by means of size exclusion, restricting object detection to >100 μ m², thus avoiding inclusion of intracellular APP and A β .

200

201 Immuno-fluorescent histochemical analysis of intracellular Aβ and phospho-Tau (AT-8)

Paraffin embedded tissue blocks of the frontal cortex BA9 were used to prepare sections (6 µm thick) 202 203 for the purpose of multiplex intracellular Aβ and phospho-tau fluorescent staining. Slide mounted 204 frontal cortex sections were baked at 60°C for 1hr, dewaxed and rehydrated and subjected to antigen 205 retrieval in citrate buffer and formic acid treatment (as above). Slides were then blocked in TBST 206 containing 10% normal goat serum for 1hr at RT and incubated in mouse IgG2b anti-MOAB-2 and 207 mouse IgG1 anti-AT-8 (1:500, for both) overnight at 4°C, prior to incubation in secondary antibodies 208 (goat-anti mouse IgG1 – Alexa 488 and goat-anti mouse IgG2b-Alexa 594, 1:1000 for both, Invitrogen). 209 Endogenous tissue fluorescent was quenched via post-staining treatment with Sudan Black (0.01%, 210 70% ethanol, 5mins submersion) before slides were coverslipped with DAPI-containing Prolong 211 Diamond Mounting media (Fisher Scientific). In a subset of slides, the limited co-localization of MOAB-212 2 labelled Aβ and APP was established, staining sections with mouse- IgG2b anti-MOAB-2 and Rabbitanti-APP (1:500, Cat# ab15272, abcam) and appropriate secondary antibodies. Fluorescence antibody 213 214 labelled sections were imaged via a wide-field fluorescence microscope system (Nikon Eclipse 90i microscope, DsQi1Mc camera and NIS elements software V 3.0, Nikon). 215

One section per case was examined at 400x magnification with 3 images per grey matter and white matter region selected at random. As these images were used for quantification of intracellular Aβ, excluding Aβ-plaques, any region selected which contained multiple plaques was excluded and another region selected. ROI were manually applied to each image and folds, tears and plaques were

excluded, before images were converted to grey scale and a consistent threshold applied to generate
 a binary image from which percentage area of immunoreactivity was determined. The mean
 percentage area of immunoreactivity was calculated per grey and white matter area per case.

224 Data analysis

225 Data were subjected to Shapiro-Wilk normality test for normal distribution, prior to statistical 226 comparison between control and AD cases using a non-parametric Mann-Whitney U test (GraphPad Prism Ver. 5). In SPSS, two-tailed Spearman's correlation was used for correlation analysis. Given the 227 228 association of increasing Braak stage with age, all correlations with Braak staging were performed with 229 partial correlations controlling from age. A series of one- tailed t-test were performed to identify the 230 initial stage at which measures were significantly elevated from Braak 0 pathological controls. For all 231 analysis, p<0.05 was considered as statistically significant, with increasing statistical reliability for p<0.01, p<0.001 and p<0.0001. 232

233

234 Results

235 **Biochemical analysis of Aβ and phospho-tau pathology.**

In order to limit the potential confounding influence of age-related A β independent tau pathology as seen in PART[12], Brodmann's area 9 of the superior frontal cortex, a region which does not develop NFTs until late stage of AD related disease progression (Braak NFT V-VI) was selected for investigation. Using a native state dot-blot quantification, crude tissue lysates of grey matter of this region as well as the associated white matter of AD (n=17) and non-AD controls (n=35), were probed for A β via the non-APP cross-reactive MOAB-2 antibody and for tau pathology using the phospho-tau specific antibody AT-8.

243 When considered purely based on the neuropathological diagnosis of either non-AD and AD, levels of AT-8 phospho-tau were elevated in AD cases, both in GM (13.24±3.3-fold cf. non-AD, p<0.001, fig1 a.i 244 245 + b.i) and WM (9±2.6 fold cf. non-AD, p<0.001, fig1 a.i + b.i). Despite a numerically higher mean within the GM compared to the WM, there was no statistically significant difference between the magnitude 246 of increase between GM and WM in AD cases (p>0.05). Similarly, when A_β levels were examined based 247 on the neuropathological cohort stratification of Non-AD and AD, elevations were apparent within the 248 249 GM (3.35±0.59 fold c.f non-AD, p<0.01, fig1 a.ii + c.ii) and WM (2.56±0.56 fold c.f non-AD, p<0.05, fig1 250 a.ii + b.ii) of AD cases. Again, no difference between the magnitude of increase within AD cases relative 251 to control cases was observed between GM and WM (p>0.05).

Such an outcome from the analysis of phospho-tau and Aβ between AD and non-AD cases is not
surprising but serves to validate the use of dot-blots to measure biochemical changes in phospho-tau
and Aβ.

To further place the observed changes of tau and Aβ within the context of disease progression, the 255 256 cohort was subdivided into their respective Braak stages (0-VI) and the association of crude lysate 257 measures of AT-8 phospho-tau and A β with disease progression was determined (Fig 1 c.i + ii). 258 Phospho-tau AT-8 immunoreactivity correlated with increasing Braak stages when considered across 259 the entire cohort within the GM (r=0.7, p<0.001) as well as in the WM (r=0.67, p<0.001). Additionally, 260 a modest but significant correlation between Braak NFT stage and age was reported (r=0.34, p<0.05) although age did not correlate with biochemical measures of AT-8 (p>0.05, data not shown). 261 262 Correlations between AT-8 immunoreactivity and Braak NFT remained when adjusting for age (r=0.62 263 and r=0.57 for GM and WM respectively, p<0.001 for both, fig 1 c). Interestingly when probed for the stage at which phospho-tau levels were significantly elevated from that of "pathologically-free" Braak 264 265 stage 0 cases, Braak stage IV was indicated (p<0.05, for both GM and WM). Braak stage IV being the 266 stage prior to gross affection of the frontal cortex with NFTs. Furthermore, when split according to 267 neuropathological diagnosis, and controlled for age, a significant correlation was observed in the non-268 AD control group in GM (Fig 1 c.i, r=0.47, p<0.01) and WM (Fig 1 c.i, r=0.43, p<0.05), but not in AD 269 cases (Fig 1 c.i, p>0.05). Together the data suggest that in non-AD control cases, AT-8 phospho-tau 270 increases within the frontal cortex in line with the intra-regional spatial AT-8 positive NFT progression, 271 somewhat independent of age. Thus, initial levels of frontal cortex AT-8 positive phospho-tau may be 272 regionally generated.

273 Following a similar line of investigation for the accumulation of A^β in relation to disease progression, 274 Aβ levels were correlated with individual Braak NFT stages. Across the entire cohort, robust 275 correlations were reported for both the GM (r=0.67, p<0.001, fig 1 d) and WM (r=0.6, p<0.001, fig 1 276 d). Again, correlations remained when controlling for age (r=0.54 and r=0.53 for GM and WM 277 respectively, p<0.001 for both). In line with observations of AT-8 phospho-tau, individual comparisons 278 with Braak 0 cases reported an initial significant elevation from the "pathologically-free" baseline at 279 Braak 4, in GM and WM samples (p<0.05). Consistent with similar measures of AT-8 phospho-tau, 280 when divided into non-AD and AD categories, increasing total A β correlated with progressive Braak 281 stages in the non-AD group in both the GM and WM (r= 0.36, p<0.05 and r=0.46, p<0.01 in GM and WM respectively, fig 1 d) and not in the AD group (p>0.05). Similar to correlation with Braak staging, 282 283 all other neuropathological classifications also reported robust correlations when considered as a single cohort (See S.table 2). When considering Non-AD controls only, significant correlation with AT8 284 285 phospho-tau was only apparent with CERAD scores for GM and WM (r=0.4, p<0.05 and r=0.47,

p<0.001, respectively) whilst GM and WM scores for Aβ correlated with Thal, CERAD and NIA-AA
(r=0.38-0.52, p<0.05, see S.table 2 for full details). In terms of AD cases, correlations were apparent
for all biochemical measures with all neuropathological scheme (see S.table 2).

Collectively, the data suggest a robust relationship of AT-8 and Aβ with disease progression within the GM and WM of the frontal cortex, occurring not only in confirmed cases of AD but also non-AD controls. In the frontal cortex, intra-regional pathology progresses in accordance with global ADrelated brain tau pathology as quantified by Braak NFT stages and this was most evident in non-AD controls.

294 Independent of global brain AD related pathological changes, a critical element to regional pathology 295 is to determine whether biochemical measures of tau pathology and A_β correlate on a case-by-case 296 basis, as the disconnect of histochemical tau and AB pathological hallmarks has long been a major 297 caveat to the amyloid cascade hypothesis. Indeed, a robust correlation between biochemical 298 measures of AT-8 phospho-tau and total A β measures was observed when considered as a single 299 cohort (Non-AD + AD cases) in the GM (r=0.75, p<0.001, fig 1 d.i) and in the WM (r=0.67, p<0.001, fig 1 300 e.i). Remarkably, when examined separately within Non-AD controls and AD cases, correlations 301 between AT-8 phospho-tau and A β were apparent in both control cases (GM; r=0.52, p<0.001 and 302 WM; r=0.67, p<0.001, fig 1 d ii + e ii) as well as AD cases (r=0.5, p<0.05 and r=0.73, p<0.01 in GM and 303 WM, fig 1 d iii + e iii).

304

305 Histochemical quantifications of AT-8 phospho-tau and Aβ plaques.

306 In order to establish if the biochemical derived relationship of increased in AB immunoreactivity 307 conferring increased in AT-8 immunoreactivity was primarily driven by an association of A β plaques 308 with AT-8 phospho-tau, semi-quantitative immunohistochemistry analysis was performed (Fig 2 a, and 309 see S.table 1 for details). Based on % area coverage, AT-8 immunoreactivity, a composite of NFTs and 310 NTs reported a marked ~ 100 fold increase in % coverage in the frontal grey matter of AD cases 311 compared to controls (7.6±2.5% vs. 0.07±0.02%, in AD and non-AD cases respectively, p<0.001, fig 2 312 b.i). Equally, guantification of the % area coverage of A β plagues between AD cases and non-AD controls, also, unsurprisingly reported a significant increase with the AD cases (14.26±1.7% cf. 313 314 3.55±1%, p<0.001, fig 2 b.ii).

When measures were considered in relation to progressive Braak NFT staging, AT-8 phospho-tau
(r=0.55, p<0.001, fig 2 c.i) and Aβ plaques (r=0.67, p<0.001, fig 2 c.ii) strongly correlated with Braak
stage, following a correction for age. Again, a significant elevation in AT-8 phospho-tau and Aβ plaque

318 coverage from "pathologically-free" Braak stage 0 cases was reported at Braak Stage IV (p<0.05), in 319 line with observations from biochemical measurements. When controlling for age, no significant 320 correlations were observed for AT-8 phospho-tau or A β plaques for either non-AD or AD groups 321 (p>0.05). Further analysis with additional neuropathological classification reported strong correlations with Thal and CERAD and NIA-AA when considered as a single cohort and in control only (S.table 3). 322 323 Collectively, the data largely suggests that correlative relationships reported within the overall cohort 324 likely stems from group effects driven by the general increase of pathological hallmarks between non-325 AD and AD groups and not as incremental regional increase in line with progressive Braak stage. Such 326 observations contrast with the findings of the biochemical investigation.

327 Equally correlations between IHC quantified Aβ and AT-8 phospho-tau also reported a correlation only 328 when the data were analysed as a single cohort combining non-AD controls and AD cases (r=0.65, 329 p<0.001, fig 2 d) and not when examined as a separate data set of non-AD controls or AD cases 330 (p>0.05, for both, fig 2 dii+iii). Nevertheless, biochemical measures correlated with respective 331 histochemical measures for AT-8 and A β (see Stable 4). Thus, despite the contradiction between biochemically measured association of AT-8 phospho-tau and total AB and histologically measured AT-332 333 8 phospho-tau and Aβ plaque load, the measures by differing methodological means are somewhat 334 inter-related.

335

336 Quantification of intracellular Aβ

337 The absence of a correlation between IHC measures AT-8 and plaques (fig 2), despite a robust 338 correlation with of biochemical AT-8 and total A β measures from crude tissue lysates (fig 1), suggests the possible inclusion of additional A β sources within the biochemical quantification. Accordingly, the 339 340 application of MOAB-2 AB antibody to fixed post-mortem human brain tissue sections, labelled both 341 extracellular plaques and intracellular pools of A β (fig 3). Comparisons of APP and MOAB-2 A β 342 labelling, demonstrated a clear distinction in subcellular and plaque labelling in both GM (fig 3 a) and 343 WM (fig 3 b). Whilst there is a degree of overlap as would be expected given the spatial limitation and the production of A β from APP, co-localisation is absence in the majority of puncta (fig 3; white 344 345 arrows), confirming the specificity of MOAB-2 for the labelling of A β , as previously reported[45, 46]. 346 Across a subset of the cohort, both in the GM and WM, a progressive increase in intracellular AB 347 staining in line with increase AT-8 immunoreactivity was observed (fig 4.a). When intracellular Aβ was quantified according to % area coverage, a significant increase in the levels was observed in AD cases 348 349 compared to controls (fig 4b.i, p<0.001 in GM and p<0.05 in WM). Similarly, AT-8, as measured by 350 immunofluorescence was again elevated in AD cases (fig 4 b ii, p<0.001 in GM and WM). Both AT-8

351 (r=0.62, p<0.01 in GM and r=0.5, p<0.05 in WM) as well as intracellular A β (r=0.61, p<0.01 in GM) correlated with Braak stage when controlling for age. Equally, strong correlations were observed with 352 Thal, CERAD and NIA-AA also (see S.table 5). No correlations with any neuropathological classifications 353 354 were reported when the cohort was spilt according to Non-AD and AD groups. Nevertheless, 355 correlative measures between intracellular AB and AT-8 phospho-tau reported a significant 356 relationship across all cases (in GM, r=0.76 and in WM, r=0.71, p<0.01 for both, fig 4 c i + d i,) and in 357 control cases only (in GM, r=0.82 and in WM, r=0.76, p<0.01 for both, fig 4 c ii + d ii). Interestingly 358 when considering only the AD cases, a significant inverse correlation of intracellular A β with AT-8 phospho-tau in the GM, was apparent (r=-0.74, p<0.05, fig 4 c iii), although no correlation was 359 360 reported in the WM (fig 4 d iii). As with IHC quantifications of plagues and AT-8 as per TMA slides, 361 measures of intracellular A β and AT-8 here, also correlated with biochemical measures when 362 considered as a single group, although no significant correlations were observed when split into 363 Control and AD categories (S. table4).

364 Collectively quantification of the intracellular A β and its correlation with AT-8 phospho-tau appears 365 to support the biochemical findings of a close relationship between AT-8 and A β , particularly in non-366 AD control cases. However is should be noted that spatial colocalization of AT-8 and intracellular A β 367 was not common and appeared to be the exception rather than the rule (fig 5), instead here we find 368 a correlation of regional AT-8 burden with regional intracellular A β .

369

370 Discussion

371 Collectively this study reports an apparent correlation between biochemical Aβ and AT-8 phospho-tau 372 measures. Such a relationship was not reproduced when comparing IHC based quantification of Aβ 373 plaques and AT-8 phospho-tau but was observed when considering IHC measures of phospho-tau and 374 intracellular Aβ. These correlative relationships are perhaps surprisingly strongest within the non-AD 375 control group and are evident in both grey and white matter. Together the data suggests a close 376 relationship between non-plaque Aβ and tau, which is at least partially due to the accumulation of 377 intracellular Aβ and its potential influence on tau phosphorylation.

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381 Regional correlation of tau and Aβ

382 A long-standing critique of the amyloid cascade hypothesis has been the disconnect between NFT and 383 Aβ-plaque burden within a given brain region of either non-AD controls or indeed AD cases [47, 48]. 384 However many biochemical approaches have previously found correlations between $A\beta$, either total 385 Aβ or specifically Aβ1-42, with a range of phosphorylated and oligomeric tau markers within the GM 386 of a given cortical region [14-17]. Furthermore, disease dependent changes in white matter A β levels 387 have also been previously observed using $A\beta_{40}$ and $A\beta_{42}$ ELISAs[49] and like-wise hyperphosphorylated tau has also been observed biochemically within the white matter of AD cases[13, 50]. Here we 388 389 observed that levels of AT-8 reactive phospho-tau and AB increased in non-AD controls in line with 390 Braak stage progression, in both the frontal grey and white matter and exhibited a positive correlation 391 between A β levels and AT-8 when measured biochemically. Whilst our observations cannot 392 determine causality, our findings are consistent with many *in-vitro* experiments in which the 393 application of A β to various cellular preparations results in downstream tau phosphorylation[28, 51, 394 52]. Moreover, support for the interaction of A β with tau pathology, can be gained from studies 395 reporting that interventions targeting A^β levels consequently reduce tau pathology both *in-vitro* and 396 in-vivo models[53, 54] as well as in biofluids obtained from human clinical trials[3, 4, 55].

397 When considering the association of A β and tau phosphorylation within non-AD controls, our findings 398 are similar to the observation of a previously seen correlation between $A\beta_{1-40}$ level and p-181 tau in 399 the CSF of control cases[56]. However here we can extend this finding to show that within the frontal 400 cortex there is a regional dependence between A β levels and tau pathology. As work by others have 401 reported no or minimal seed competent tau within the frontal cortex in non-AD controls [50], the 402 source of pathology within the frontal cortex is likely to mainly be intraregional, with minimal 403 influence of extra-regional spread. Such intraregional dependence would be consistent with the 404 strong correlation between A β and AT-8 signals within the non-AD controls, which is somewhat 405 weakened in AD cases, when seed competent tau species have presumably invaded the frontal 406 cortex[13], the self-propagation of tau pathology thus diminishing the relative contribution of A β to 407 the production of phospho-tau species. Interestingly, a linear relationship between regional soluble 408 tau phospho-species and $A\beta_{1-42}$ has previously been reported alongside a bimodal relationship within 409 the insoluble fraction, implying a weakening of the relationship between $A\beta_{1-42}$ and tau 410 phosphorylation once aggregated [17]. Thus, given that this present study did not seek to distinguish 411 between soluble and insoluble pathology, the increased representation of insoluble tau pathology 412 within lysates from AD cases, likely further explains the weakening of the linear relationship between 413 A β and tau phosphorylation in AD cases.

Accordingly, no linear correlation was observed between plaque load and AT-8 load when measured
 histologically. Such differences may relate to the presumed loss of extracellular soluble Aβ and

416 exclusion of intracellular Aβ pools, as is common practice when assessing Aβ burden as part of a417 neuropathological assessment[44].

418

419 Intracellular Aβ

Historically, intracellular Aβ has been problematic in its quantification, largely due to the crossreactivity of Aβ antibodies with APP and other intermediate APP metabolites. Howecer, several
commercial Aβ antibodies are available, including MOAB-2 which shows no cross-reactivity with APP
under many conditions [45, 46].

Although often overlooked, the production of AB occurs intracellularly following endosomal APP 424 425 cleavage via β -secretase [57] and sequential γ -secretase processing within either Golgi [58] or 426 lysosomal [59] compartments. Whilst the majority of A β is trafficked to the extracellular space, age-427 related changes in the relative production of AB peptide length [60] and disease alterations to 428 trafficking mechanisms, such as Rab GTPases [61, 62], may act synergically to enhance the retention of intracellularly produced A β and or indeed its reuptake, leading to its intracellular accumulation [63]. 429 430 Accordingly, post-mortem examination of the entorhinal cortex and hippocampus of non-diseased 431 non-AD cases, suggests an increase in intracellular A β in line with increasing age [64, 65] and 432 furthermore AD animal models also show an age-related accumulation of intracellular A β [46, 66].

433

434 Intracellular Aβ and tau

435 Here, when selectively measuring intracellular A β , a positive correlation between A β and AT-8 in the frontal cortex of non-AD controls was observed. This is consistent with biochemical measures of 436 437 frontal cortex lysates in the same cases as present here or indeed previously reported biochemical 438 measures in the lateral temporal cortex of a different study cohort[15]. However, in work by others, 439 no such relationship has been observed in the entorhinal cortex [64][67]. Such a contrast in 440 relationships may relate to the regions selected for investigation. The entorhinal cortex is one of the 441 earliest affected cortical regions with tau pathology in AD, and thus presumably represents a cortical 442 area of increased vulnerability to tau pathology. Such vulnerability may mean that tau pathology may 443 be generated in an A β independent manner within the region as is the case of primary age-related 444 tauopathy [12]. Nevertheless, within the prefrontal cortex, a region which does not demonstrate 445 robust age related NFT tau pathology and is not burdened with NFTs until late into the Braak NFT staging criteria (Braak V-VI), such modest pre-tangle tau-pathology generated in this region may be 446 447 largely dependent on AB mediated mechanism. Such a mechanism may become modified under pathological conditions, either upon reaching a critical threshold of intracellular Aβ accumulation
and/or via seed component invasion.

450 Interestingly within the grey matter, although a positive relationship between intracellular A β and 451 phospho-tau was observed in non-AD control cases, an inverse relationship was observed in AD cases. 452 Whilst intracellular AB levels remained elevated compared to controls, there has been reports of a 453 reduction of intracellular A β levels in line with the deposition of A β plaques in mice models [68] as 454 well as in serial observations in Down's syndrome brains [69] and cases of late stage NFT mediated neurodegeneration [67]. Such contradictions may be due to differences in the specific regions of 455 456 investigations. However, it is equally plausible that elevation of intracellular AB precedes and indeed 457 acts as a source for extracellular plaque deposition, with the excessive deposition of plaques at later 458 stages subsequently reducing intracellular A β levels as observed in animal model [68]. In turn tau 459 pathology may continue to grow due to the influence of seed component species.

In any case, excessive intracellular Aβ accumulation, is unlikely to be benign. The familial Osaka E639Δ 460 461 APP mutant which produces non-fibril E22 Δ A β gives rise to an accumulation of intracellular A β 462 oligomers in the absence of plaques. In AD patients or mouse models carrying the Osaka mutation, 463 pronounced cognitive impairments, cellular stress, synaptic spine loss and critically pathological tau 464 phosphorylation and conformational changes are observed [22, 45]. Whilst several of the downstream 465 cellular dysfunction may be independent of tau [70], these studies nevertheless highlight the induction 466 of tau pathology via AB independent of plaque formation, in support of the observed correlation of intracellular Aβ and phospho-tau observed here. 467

Given emerging evidence from clinical Aβ antibody trials[71], which supporting the targeting of soluble fibrillar Aβ species to consequently reduce tau pathology, further understanding the degree of interaction between Aβ and tau will provide greater insight into the mechanisms of AD related pathogenesis. Equally, in light of the facilitation of fibril seeding by the existence of pre-existing tau phosphorylation/pathology in mice [72-74], the targeting of pre-tangle soluble tau elevations in late stage affected brain regions, may protect against tau seed infiltration as part of AD disease progression, and may provide an effective stalling of the condition.

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479 Conclusions

480 Collectively this study demonstrates the robust correlation of AT-8 reactive tau and A^β in the frontal 481 cortex of both non-AD controls and AD cases when measured biochemically. Given that such linear 482 increases in AB plaques and AT-8 pathology is not observed when quantified via IHC, the study 483 demonstrates the potential influence of non-plaque A β in the intra-regional generation of tau 484 pathology. Specifically, the occurrence and accumulation of intracellular $A\beta$, in line with AD 485 pathological progression, at least in part correlated with AT-8 pathology and thus may contribute to production of tau pathology. This finding is supportive of the amyloid cascade hypothesis, yet in late-486 487 stage AD cases such a relationship may be diminished, with additional factors contributing to tau 488 pathology, at least within the frontal cortex. Critically, the observation of a localised relationship 489 between AB and phospho-tau in cases with low Braak NFTs stages implies that there is a degree of 490 regionally generated AD related pathology, which may be tolerated within a physiological range. 491 Following the age-related accumulation of pathology, this regionally produced burden may prime the 492 region for the invasion of seed competent forms of tau originating from connected regions, raising the burden beyond a critical threshold, and thus removing the necessity of $A\beta$ for the propagation of tau 493 494 pathology.

495

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509 Legends

510 Table legends

511 Table 1. Post-mortem human tissue cases and use. Human cases use for immuno-blots and 512 immunohistochemistry for plagues and AT-8 as well as intracellular A β and AT-8 are listed. Case are separated by disease classification according to non-diseased controls (Con) and Alzheimer's disease 513 514 (AD). Case numbers (n), sex, age, post-mortem interval (PMI), neurofibrillary tangle (NFT) Braak stage, 515 Thal phase, Consortium to Establish a Registry for Alzheimer's Disease (CERAD), the National Institute 516 of Ageing – Alzheimer's Association (NIA-AA) criteria, Lewy body (LB) Braak stage and McKeith criteria 517 are provided. For age and PMI both range and mean ±SEM are provided. For numerical scores of 518 pathology, range and percentage composition are given. For CERAD scores, negative (neg), A and B 519 reported. For NIA-AA, not, low and intermediate (inter) risk for Alzheimer's disease. For McKeith 520 criteria, only percentage composition is given, where cases free of LBs (No LB), brainstem, Limbic and 521 neocortical (Neo) predominate are indicated.

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523 Figure legends

524 Figure 1. Biochemical quantification of AT-8 phospho-tau and Aβ in the frontal cortex of non-AD and AD cases. A). Example dot blots of AT-8 (i) and MOAB-2 (A β ; ii) immunoreactivity and associated 525 526 ponceau total protein stain, produced from crude tissue lysates of frontal grey (GM) and white matter 527 (WM) in control (Con; black lettering) and Alzheimer's disease (AD; red lettering) cases. Braak NFT 528 stage of each sample is shown. B). Comparison of mean AT-8 (i) and A β (ii) immunoreactivity between 529 control (Con) and Alzheimer's disease (AD) cases in the GM and WM C). Association of AT-8 (i) and AB 530 (ii) immunoreactivity with Braak NFT stages across the cohort in GM and WM. Correlative analysis 531 (Spearman's r) is shown for when analysis as a single group or when separated into Con and AD groups. Combined (i), Con(ii) and AD (iii), linear correlations between AT-8 and A β in the GM (D) and WM(E). 532 Immunoreactive shown as relative to control (Rel. Control). *=p<0.05, **=p<0.01, ***=p<0.001 and 533 534 ****=p<0.0001. \$ denotes initial Braak NFT stage at which immunoreactivity is significantly elevated 535 from Braak 0 controls.

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540 Figure 2. Immunohistochemical quantification of AT-8 tau and Aß plaque burden in the frontal cortex of non-AD and AD cases. A). Example micrographs of DAB based AT-8 and 4G8 (AB) 541 immunoreactivity, area of quantification following threshold application shown in red. Note the size 542 543 exclusion in this parameter of intracellular 4G8 labelling to negate potential APP cross-reactivity. B). 544 Quantification of % area coverage of AT-8 (i) and plaque (ii) immunoreactivity in control (Con) and 545 Alzheimer's disease (AD) cases. C). Association of % area coverage of AT-8 (i) and plaques (ii) with 546 Braak NFT stage with correlative analysis (Spearman's r) shown. Combined(i), Con(ii) and AD(iii), linear correlations between AT-8 and plaques in the GM (D). N.S = not significant, ***=p<0.001 and 547 ****=p<0.0001. \$ denotes initial Braak NFT stage at which immunoreactivity is significantly elevated 548 549 from Braak 0 controls. Scale in a = $20 \,\mu m$.

550 Figure 3. Immunohistochemical distinction between MOAB-2 labelled Aβ and APP immunoreactivity

in the frontal cortex of an AD case. Example micrographs of APP (N-terminal-APP antibody) and Aβ
 (MOAB-2) from an AD case, in the grey (GM; a) and white matter (WM; b). Note the distinctive labelling
 of subcellular pools within insert (white arrows) and differential labelling of plaques (in a) Scale =20
 µm.

- Figure 4. Immunohistochemical quantification of intracellular A β and AT-8 tau in the frontal cortex of non-AD and AD cases. A). Example micrographs of AT-8 phosphorylated tau and MOAB-2 labelled intracellular A β in GM and WM of low (i), intermediate (ii), and high (iii) Braak stage cases. B). Quantification of intracellular A β (i) and AT-8 phospho-tau (ii) expressed as percentage area coverage in control (Con) and Alzheimer's disease (AD) cases. Combined(i), Con(ii) and AD(iii), spearman's correlations (r) between AT-8 and plaques in the GM (C) and WM (D). N.S = not significant, *=p<0.05, **=p<0.01, ***=p<0.001 and ****=p<0.0001. Scale =20 µm.
- Figure 5. Rare instances of intracellular Aβ and tau colocalization. Example micrographs
 demonstrating an overlap of AT-8 and Aβ immunoreactivity within cells from a non-AD control Braak
 stage IV. Scale=10 µm.
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572 Supplemental

Supplemental table 1 – Individual case list for tissue used. Each case was assigned an arbitrary case
 number (Case no) and sex, age (years), post-mortem delay (PMD, hrs), Braak stage, Thal phase,
 Consortium to Establish a Registry for Alzheimer's Disease (CERAD), the National Institute of Ageing –
 Alzheimer's Association (NIA-AA) criteria, Lewy body (LB) Braak stage and McKeith criteria and tissue

- 577 use of either biochemical (Biochem) analysis, tissue microarray for plaque and AT-8 pathology (TMA)
- and immunohistochemistry for intracellular A β and AT-8 (IHC) is provided.

579 Supplemental table 2. Correlation matrix of biochemical measures of Aβ and AT-8 reactive tau

- 580 with neuropathological assessment. Spearman correlations (r) between AT-8 and Aβ and Braak NFT
- staging, Thal phase, CERAD, NIA-AA, Age and postmortem delay (PMD) are shown for both the grey
- 582 matter (GM) and white matter (WM) samples. Note for Braak stage correlations reported are 583 controlled for the influence of age. N.S = not significant, *=p<0.05, and **=p<0.01.
- _____
- 584 Supplemental table 3. Correlation matrix of immunohistochemical measures of Aβ plaques and
- 585 **AT-8 reactive tau with neuropathological assessment.** Spearman correlations (r) between AT-8 and
- Aβ and Braak NFT staging, Thal phase, CERAD, NIA-AA, Age and postmortem delay (PMD) are shown.
 Note for Braak stage correlations reported are controlled for the influence of age. N.S = not
- 588 significant, *=p<0.05, and **=p<0.01.
- 589 Supplemental table 4. Correlation matrix of all measures of Aβ and AT-8 reactive tau across the
- 590 **study** Spearman correlations (r) between AT-8 and Aβ and Braak NFT staging, Thal phase, CERAD,
- 591 NIA-AA, Age and post-mortem delay (PMD) are shown. Note for Braak stage correlations reported
- are controlled for the influence of age. N.S = not significant, *=p<0.05, and **=p<0.01.
- 593 Supplemental table 5. Correlation matrix of immunohistochemical measures of intracellular Aβ
- and AT-8 reactive tau.. Spearman correlations (r) between AT-8 and Aβ and Braak NFT staging, Thal
 phase, CERAD, NIA-AA, Age and postmortem delay (PMD) are shown for both the grey matter (GM)
 and white matter (WM) samples. Note for Braak stage correlations reported are controlled for the
 influence of age. N.S = not significant, *=p<0.05, and **=p<0.01.
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Disease	N	Sex (% Male)	Age (years)	PMI (hrs)	NFT Braak stage	Thal Phase	CERAD	NIA-AA	LB Braak stage	McKeith Criteria
Immuno-blots	Immuno-blots									
Con	35	57.1%	47-97 80.5±2.2	16-95 45.8±3.7	0-IV 17.1% -0 17.1%- I 25.7% - II 25.7%-III 14.3%-VI	0-4 20%-0 22.9%-1 25.7%-2 22.9%-3 8.6%-4	0-2 80%-0 5.7%-1 14.3%-2	0-2 20 %- 0 65.7 % - 1 14.3 % - 2	0-3 91.4%-0 2.9%-1 5.7%-3	91.4%-No LB 8.6%-Brainstem
AD	17	35.3%	74-96 85.5±1.5	5-90 57.2±5.7	V-VI 29.4% -V 70.6% -VI	4-5 15.4%-4 84.6%-5	3 100% -3	3 100% - 3	0-3 82.3%-0 11.8%-2 5.9%-3	76.5% -No LB 17.6% -Brainstem 5.9%-Limbic
Immunohistoc	chemistry (AT-8	and 48G plaque	s)			1				
Con	36	56.2%	55-97 81.5±2.2	16-95 47.8 ±3.9	0-IV 16.7% -0 19.4%- I 25% - II 25%-III 13.9%-VI	0-4 16.7%-0 25%-1 30.6%-2 19.4%-3 8.3%-4	0-2 77.8%-0 8.3%-1 13.9%-2	0-2 16.7%-0 69.4%-1 13.9%-2	0-3 88.1%-0 5.8%-1 6.1%-3	88.7%-No LB 8.3%-Brainstem 2.8%-Limbic
AD	20	30%	70-93 85±1.3	5-90 54.4±4.9	V-VI 25%-V 75%-VI	4-5 13.9%-4 86.1%-5	3 100%-3	3 100%-3	0-3 85%-0 10%-2 5%-3	80% -No LB 15%-Brainstem 5%-Limbic
Immunohistoc	hemistry (AT-8	and MOAB-2 int	racellular Aβ)	I					1	
Con	14	28.6%	70-97 85.6±2.4	16-95 45.1 ±7.3	0-IV 7.1% -0 14.3%- I 42.9% - II 35.7%-III	0-2 28.6%-0 35.7%-1 35.7%-2	0-1 85.7%-0 14.3%-1	0-1 35.7%-0 64.3%-1	0-3 85.7%-0 14.3%-3	78.6%-No LB 14.3%-Brainstem 7.1%-Limbic
AD	8	12.5%	78-93 85.4±1.8	29-90 57±7.5	V-VI 12.5%-V 87.5%-VI	4-5 12.5%-4 87.5%-5	3 100%-3	3 100%-3	0-3 87.5%-0 12.5%-2	75% -No LB 25%-Brainstem

Table 1. Human tissue cohort. Human cases using in tissue microarray and western blot separated by disease classification according to non-diseased controls (Con), Alzheimer's disease (AD) and dementia with Lewy bodies (DLB). Case numbers (n), sex, age, post-mortem interval (PMI), neurofibrillary tangle (NFT) Braak stage, Thal phase, Consortium to Establish a Registry for Alzheimer's Disease (CERAD), the National Institute of Ageing – Alzheimer's Association (NIA-AA) criteria, Lewy body (LB) Braak stage and McKeith criteria are provided. For age and PMI both range and mean ±SEM are provided. For numerical scores of pathology, range and percentage composition are given. For McKeith criteria, only percentage composition is given, where cases free of LBs (No LB), amygdala predominate (Amyg), limbic predominate (Limb) and neocortical predominate (Neo) are indicated. * indicates composition based on available data.









Figure 3





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