

1 **Assessment of the relationship between synaptic density and metabotropic glutamate**
2 **receptors in early Alzheimer’s disease: a multi-tracer PET study**

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23 Running Title: mGluR5 and SV2A PET in Alzheimer’s disease

24 **Abstract**

25 **Background:** The pathological effects of amyloid β oligomers ($A\beta$) may be mediated through
26 the metabotropic glutamate receptor subtype 5 (mGluR5), leading to synaptic loss in
27 Alzheimer's disease (AD). Positron emission tomography (PET) studies of mGluR5 using
28 [^{18}F]FPEB indicate a reduction of receptor binding that is focused in the medial temporal lobe in
29 AD. Synaptic loss due to AD measured through synaptic vesicle glycoprotein 2A (SV2A)
30 quantification with [^{11}C]UCB-J PET is also focused in the medial temporal lobe, but with clear
31 widespread reductions is commonly AD-affected neocortical regions. In this study, we used
32 [^{18}F]FPEB and [^{11}C]UCB-J PET to investigate the relationship between mGluR5 and synaptic
33 density in early AD.

34 **Methods:** Fifteen amyloid positive participants with early AD and 12 amyloid negative,
35 cognitively normal (CN) participants underwent PET scans with both [^{18}F]FPEB to measure
36 mGluR5 and [^{11}C]UCB-J to measure synaptic density. Parametric *DVR* images using equilibrium
37 methods were generated from dynamic. For [^{18}F]FPEB PET, *DVR* was calculated using
38 equilibrium methods and a cerebellum reference region. For [^{11}C]UCB-J PET, *DVR* was
39 calculated with a simplified reference tissue model – 2 and a whole cerebellum reference region..

40 **Result:** A strong positive correlation between mGluR5 and synaptic density was present in the
41 hippocampus for participants with AD ($r = 0.81, p < 0.001$) and in the CN group ($r = 0.74, p =$
42 0.005). In the entorhinal cortex, there was a strong positive correlation between mGluR5 and
43 synaptic in the AD group ($r = 0.85, p < 0.001$), but a weaker non-significant correlation in the CN
44 group ($r = 0.36, p = 0.245$). Exploratory analyses within and between other brain regions
45 suggested significant positive correlations between mGluR5 in the medial temporal lobe and
46 synaptic density in a broader set of commonly AD-affected regions.

47 **Conclusion:** Medial temporal loss of mGluR5 in AD is associated with synaptic loss in both
48 medial temporal regions and more broadly in association cortical regions, indicating that
49 mGluR5 mediated A β toxicity may lead to early synaptic loss more broadly in AD-affected
50 networks. In CN individuals, an isolated strong association between lower mGluR5 and lower
51 synaptic density may indicate non-AD related synaptic loss.

52

53 **Keywords:** Alzheimer's disease, mGluR5 availability, synaptic density, SV2A, PET, [¹⁸F]FPEB,
54 [¹¹C]UCB-J.

55

56

57 **Background**

58 Alzheimer's disease (AD) results in early and pronounced synaptic loss as a prominent
59 pathological feature (1-4). Evidence supports a robust correlation between synaptic loss and level
60 of cognitive impairment (5, 6), as determined by postmortem and brain biopsy studies, as well as
61 synaptic positron emission tomography (PET) imaging (7-10). [¹¹C]UCB-J was developed as a
62 PET tracer for synaptic vesicle glycoprotein 2A (SV2A) in the past decade and has shown
63 promising results in investigations of synaptic density in human studies, including studies of AD
64 (11-13). [¹¹C]UCB-J has a high in vivo affinity for SV2A, which resides within synaptic vesicles
65 located at presynaptic terminals (14, 15). We have reported widespread reductions in synaptic
66 density in the medial temporal lobe and in common AD-affected neocortical brain regions using
67 [¹¹C]UCB-J PET (7, 13, 16). This has been corroborated by multiple other groups (17-22).
68 Glutamate is the primary excitatory neurotransmitter in the nervous system with ionotropic
69 glutamate receptors being the main conduit for information transfer within the central nervous

70 system (23). However pre- and postsynaptic metabotropic glutamate receptors (mGluRs) are
71 commonly present and help with fine-tuning synaptic communication between neurons by
72 regulating strength and timing of network activity (24). Metabotropic glutamate receptor subtype
73 5 (mGluR5) is a seven-transmembrane G protein-coupled receptor expressed in neurons and glial
74 cells throughout the cortex and hippocampus that has a non-homogeneous distribution pattern
75 (24-28). Based on mouse hippocampal neuron studies, mGluR5 have been considered primarily
76 post-synaptic and involved in inducing long-term depression (LTD) at NMDAR synapses (26,
77 29, 30). However, more recent evidence indicates a heterogenous localization and function for
78 mGluR5 with presynaptic, postsynaptic, and intracellular expression. Non-human primate studies
79 indicate that mGluR5 is expressed in both presynaptic and postsynaptic terminals in the
80 dorsolateral prefrontal cortex (31). Additionally, studies in rats demonstrate the existence of
81 functional intracellular mGluR5 in hippocampus. Based on animal models of AD, it has been
82 hypothesized that mGluR5 contributes to amyloid- β oligomer ($A\beta$) toxicity through various
83 mechanisms. This includes facilitating the clustering of $A\beta$ as an extracellular scaffold for
84 mGluR5 – leading to $A\beta$ -induced abnormal mGluR5 accumulation and subsequent increase in
85 intracellular calcium levels and synaptic deterioration (32), as well as mGluR5 acting as a co-
86 receptor with cellular prior protein (PRPc) and subsequent postsynaptic activation of the tyrosine
87 kinase Fyn (33, 34). The latter finding asserts mGluR5 as a link between $A\beta$ and tau pathology
88 where the activation of Fyn leads to downstream tau phosphorylation (35). Recognition of
89 mGluR5 as a mediator of AD pathology has spurred research into its role as a therapeutic target
90 in AD mouse models as well as in human clinical trials (36-42).

91 Several recent human PET imaging studies with mGluR5 specific radiotracers have made
92 it possible to assess mGluR5 changes in individuals affected by clinical AD. Our previous work

93 quantifying mGluR5 availability in AD with [¹⁸F]FPEB PET showed a significant reduction of
94 hippocampal mGluR5 due to AD with non-significant, but numerically lower mGluR5 binding
95 in association cortical regions (25). This finding was corroborated in studies by Wang *et al.* and
96 Treyer *et al.* using [¹⁸F]PSS232 and [¹¹C]-ABP699 PET respectively (43, 44).

97 As an extension of our previous work showing synaptic density and mGluR5 reductions
98 in AD we performed analyses to investigate the spatial relationships between both biomarkers in
99 a cohort of individuals who underwent both [¹⁸F]FPEB and [¹¹C]UCB-J PET. Because the largest
100 reductions of mGluR5 and synaptic density are found in the medial temporal lobe, in our primary
101 analyses we focused on the hippocampus and entorhinal cortex. We then examined brain wide
102 regional correlations between mGluR5 and synaptic density. We hypothesized that mGluR5 and
103 synaptic density would be strongly correlated in participants with AD, not in CN participants.

104

105

106 **Methods**

107 The study protocol was approved by the Yale University Human Investigation Committee
108 and Radiation Safety Committee. All participants provided written informed consent prior to
109 participating in the study.

110

111 **Study Participants**

112 Participants between 55 and 85 years of age were evaluated with a screening diagnostic
113 evaluation, as previously described (45). Participants with AD were required to either i) meet the
114 diagnostic criteria for probable dementia based on National Institute on Aging-Alzheimer's
115 Association (NIA-AA) guidelines, have a Clinical Dementia Rating (CDR) score of 0.5 -1, and

116 Mini-Mental Status Examination (MMSE) score of ≥ 16 or ii) meet the NIA-AA diagnostic
117 criteria of amnesic mild cognitive impairment (aMCI), have a CDR score of 0.5, and an MMSE
118 score of 24-30. Moreover, participants in the AD group were required to demonstrate impaired
119 episodic memory, as evidenced by a Logical Memory (LM) II score of 1.5 standard deviations
120 (SD) below an education-adjusted norm. CN participants were required to have a CDR score of
121 0, an MMSE score > 26 , and a normal education adjusted LMII.

122 All participants underwent PET with [^{11}C]Pittsburg Compound B ([^{11}C]PiB) to assess for
123 the presence of brain A β . [^{11}C]PiB PET scans were required to be negative for A β in CN
124 participants and positive in AD participants. Participants were considered A β + if the [^{11}C]PiB
125 PET scan was positive based on visual interpretation of 2 expert readers and confirmed with
126 quantitative read criteria of cerebral-to-cerebellar distribution volume ratio (*DVR*) of at least 1.40
127 in at least 1 AD-affected region of interest (ROI) (7, 46).

128

129 **Magnetic resonance imaging**

130 Magnetic resonance imaging (MRI) was conducted using a 3T Trio (Siemens Medical
131 Systems, Erlangen, Germany) equipped with a circularly polarized head coil. MRI acquisition
132 consisted of a Sag 3D magnetization-prepared rapid gradient-echo (MPRAGE) sequence with
133 the following parameters: 3.34-msec echo time, 2500-msec repetition time, 1100-msec inversion
134 time, 7-degree flip angle, and 180 Hz/pixel bandwidth. The resulting images have dimensions of
135 $256 \times 256 \times 176$ with a pixel size of $0.98 \times 0.98 \times 1.0$ mm. The MRI procedure was used to make
136 sure that patients did not show evidence of infection, infarction, or other brain lesions. Moreover,
137 it served to delineate brain anatomy, assess atrophy, and perform partial volume correction
138 (PVC) of PET images. Version 6.0 of FreeSurfer (<http://surfer.nmr.mhg.harvard.edu/>) was used

139 to reconstruct cortical regions and perform volumetric segmentation used to define ROIs in
140 participant native space (47).

141

142 **Positron emission tomography methods**

143 PET images were acquired on the High-Resolution Research Tomograph (Siemens
144 Medical Solution, Knoxville, TN, USA, 207 slices, resolution < 3 mm full width half maximum)
145 (48). Dynamic [^{11}C]PiB scans were obtained over a period of 90 minutes after the bolus
146 administration of a tracer dose of up to 555MBq (49). [^{18}F]FPEB was used to quantify regional
147 brain availability of mGluR5. Using the previously evaluated bolus plus constant infusion
148 paradigm (Kbol = 190 min), dynamic [^{18}F]FPEB scans were taken for 60 minutes, beginning at
149 60 minutes after the initial injection of up to 185 MBq of tracer. Lastly, [^{11}C]UCB-J PET was
150 used for evaluating synaptic density by acquiring dynamic scans up to 90 minutes after
151 administration of a tracer bolus of up to 740 MBq (50).

152 Using the Motion-compensation OSEM List-mode Algorithm for Resolution-recovery
153 (MOLAR), list-mode data was reconstructed with event-by-event motion correction based on
154 Vicra optical detector (NDI Systems, Waterloo, Canada) (51, 52). Software motion correction
155 was applied to the dynamic PET images using a mutual-information algorithm (FSL-FLIRT, FSL
156 3.2; Analysis Group, FMRIB, Oxford, UK) to perform frame-by-frame registration to a summed
157 image (0 - 10 min for [^{11}C]UCB-J and 60 - 70 min for [^{18}F]FPEB. A summed motion corrected
158 PET image was used to create a registration between the MRI and PET scans for each
159 participant. This PET to MRI registration was used to apply participant specific ROIs to
160 parametric PET images.

161 For [¹¹C] PiB, parametric images of binding potential (BP_{ND}), the ratio at equilibrium of
162 specifically bound radioligand to that of nondisplaceable radioligand in tissue, were generated
163 using simplified reference tissue model-2 (SRTM2) using whole cerebellum as reference region.
164 In order to account for potential partial volume effects, we performed partial volume correction
165 of dynamic series for [¹⁸F]FPEB and [¹¹C]UCB-J using the iterative Yang method (53, 54).
166 Kinetic modeling was performed both with and without PVC of dynamic PET series. For
167 [¹⁸F]FPEB image analysis, parametric images of DVR were generated with equilibrium methods
168 using data collected from 90 to 120 minutes post bolus injection and a whole cerebellum
169 reference region, as previously described (25). Lastly, for [¹¹C]UCB-J, SRTM2 was applied to
170 generate parametric BP_{ND} images PET frames from 0 to 60 minutes post injection and a whole
171 cerebellum reference region.(55) For [¹¹C]UCB-J, BP_{ND} was converted to DVR using the formula
172 $DVR = BP_{ND} + 1$.(16, 49)

173 Reported values for each ROI are bilateral regions except where specified as left or right
174 hemisphere. ROIs used for the composite of AD-affected brain regions are defined in
175 Supplementary Table 1. ROIs used for the medial temporal composite included bilateral
176 hippocampus, entorhinal cortex, parahippocampal cortex, and amygdala.

177

178 **Statistical analysis**

179 Statistical analyses were performed using MATLAB R2018b (Mathworks, Natick, MA,
180 USA) and SPSS 28 (IBM Corp, Armonk, NY). Between group comparisons were performed
181 using χ^2 tests for categorical variables, independent two-tailed t tests for continuous variables, as
182 well as Mann-Whitney U tests for CDR global and CDR sum of boxes scores. Separate
183 univariate regression analyses were used to evaluate the relationship between mGluR5 and

184 synaptic density with the primary analysis focused on hippocampus and entorhinal cortex.
185 Pearson's correlation coefficients (r) and associated two-tailed p values were calculated to assess
186 the strength of linear correlation between mGluR5 and synaptic density in each ROI, as well as a
187 medial temporal composite region. Fisher r -to- z transformation was used to compare the strength
188 of correlation of mGluR5 and synaptic density between AD and CN groups. Significant p value
189 was defined as < 0.05 . Analyses were performed both without and with PVC of PET data.
190 Analyses including all brain regions did not include correction for multiple comparisons due to
191 the exploratory nature of these investigations.

192

193

194 **Results**

195 **Participants characteristics**

196 The study sample consisted of 15 amyloid positive participants with AD and 12 amyloid
197 negative participants with normal cognition. Demographic characteristics, cognitive assessment
198 results, *APOE* genotype, and PET *DVR* measures for each group are shown in Table 1.
199 Diagnostic groups were well balanced for age, sex, and education. Additionally, all participants
200 with AD demonstrated typical clinical characteristics of aMCI or mild dementia, with significant
201 deficits indicated by MMSE (24.1 ± 3.9), CDR global score (0.7 ± 0.2), and CDR sum-of-boxes
202 score (4.0 ± 2.2) in comparison to participants with normal cognition (Table 1). *APOE* genotypes
203 reflected expected patterns with higher copy numbers of the $\epsilon 4$ in the AD participant group. As
204 expected from our previous studies, synaptic density ($[^{11}\text{C}]\text{UCB-J}$ PET *DVR*) was lower in both
205 the hippocampus and a composite of common AD-affected brain regions in participants with AD
206 compared to the CN group. In this slightly smaller sample than our previous study with

207 [¹⁸F]FPEB PET (25), hippocampal mGluR5 density was lower, but not significantly different, in
208 participants with AD compared to the CN group. In line with our previous observation, mGluR5
209 density in the composite of AD-affected regions was not lower in participants with AD compared
210 to the CN group.

211

212 **Correlations between mGluR5 and synaptic density in hippocampus and entorhinal cortex**

213 Our primary analyses used univariate linear regression to assess the relationship between
214 mGluR5 and synaptic density in the hippocampus and entorhinal cortex, regions known to be
215 involved in early AD pathogenesis and with significant AD related reductions of synaptic density
216 and mGluR5 density based on our previous studies. A strong, significant positive correlation was
217 demonstrated between hippocampal mGluR5 and synaptic density in participants with AD ($r =$
218 $0.81, p < 0.001$) and a slightly weaker, significant positive correlation in the CN group ($r = 0.74,$
219 $p = 0.005$, Figure 1A). Significant correlations of similar strength were also present in the
220 hippocampus with PVC of the PET data ($r = 0.82, p < 0.001$ for AD and $r = 0.73, p = 0.007$ for
221 CN). A Fisher r -to- z transformation indicated no statistically significant difference in the strength
222 of correlations in the hippocampus between the two groups without PVC ($z = 0.35, p = 0.704$)
223 and with PVC ($z = 0.50, p = 0.617$). In the entorhinal cortex, a strong, significant positive
224 correlation was demonstrated between mGluR5 and synaptic density in participants with AD ($r =$
225 $0.85, p < 0.001$), but no significant correlation was found in the CN group ($r = 0.36, p = 0.245,$
226 Figure 1B). Correlations of similar strength were also present in the entorhinal cortex with PVC
227 of the PET data ($r = 0.83$ with $p < 0.001$ for AD, $r = 0.40, p = 0.196$ for CN). Although group
228 differences in correlation strength were similar in magnitude, the correlation between mGluR5
229 and synaptic density in the entorhinal cortex was significantly stronger in participants with AD

230 compared to the CN group without PVC ($z = 1.99, p = 0.046$), but not with PVC ($z = 1.76, p =$
231 0.078).

232

233 **Correlations between mGluR5 and synaptic density in other medial temporal regions**

234 To better understand the pattern of correlations in brain areas affected in early AD, we
235 next focused on the medial temporal lobe in analyses of a composite medial temporal region, as
236 well as the individual regions used to construct the composite (Supplementary Figure 1). There
237 was a strong, positive correlation between mGluR5 and synaptic density in the medial temporal
238 lobe of the AD group ($r = 0.84, p < 0.001$), and no significant correlation in the CN group ($r =$
239 $0.56, p = 0.055$). In addition to the relationships described in the primary analyses for
240 hippocampus and entorhinal cortex, mGluR5 and synaptic density had a strong, positive
241 correlation in the amygdala for the AD group ($r = 0.84, p < 0.001$), and a weaker non-significant
242 correlation in the CN group ($r = 0.56, p = 0.057$). In the parahippocampal cortex, there was a
243 strong, positive correlation in the AD group ($r = 0.85, p < 0.001$) and a weaker, non-significant
244 correlation in the CN group ($r = 0.42, p = 0.170$). A very similar pattern of correlation and
245 significance existed with application of PVC to the PET data, as well as after adjustment for
246 multiple comparisons using false discovery rate (Supplementary Figure 1). All statistically
247 significant correlations in the figure remained statistically significant after correction for multiple
248 comparisons using false discovery rate (FDR) method except for amygdala in data with PVC in
249 participants with normal cognition.

250

251 **Correlations between mGluR5 and synaptic density in all brain regions**

252 We performed exploratory analyses in all brain regions to have a better understanding of
253 the whole brain pattern of correlations between mGluR5 and synaptic density. Stronger
254 significant correlations between mGluR5 and synaptic density were observed more broadly in
255 participants with AD compared to the CN group both without and with PVC (Figure 2, Table 2,
256 and Supplementary Table 2). Without PVC, regions with significant correlations in the AD group
257 included bilateral temporal poles, entorhinal cortices, hippocampi, parahippocampal cortices,
258 amygdalae, fusiform gyri, inferior/middle/superior temporal gyri, banks of the superior temporal
259 sulci, insular cortices, medial orbitofrontal cortices, rostral anterior cingulate gyri, as well as left
260 caudate, right pars opercularis, right transverse temporal gyrus, right supramarginal gyrus, right
261 isthmus of the cingulate, right inferior parietal cortex, and right lingual gyrus. In the CN group,
262 significant correlations existed only in the bilateral hippocampi, bilateral caudate, left pallidum,
263 right transverse temporal cortex, left insular cortex, and left thalamus. When using Fisher r -to- z
264 transformation to assess the difference in correlation strength between AD and CN groups, the
265 bilateral temporal poles, left banks of superior temporal sulcus, and right entorhinal cortex had
266 significantly stronger positive correlations in participants with AD as compared to the CN group.
267 Similar relationships were seen with and without PVC of PET data (Table 2 and Supplementary
268 Table 2).

269 To explore the relationships of mGluR5 and synaptic density between different brain
270 regions, we constructed a matrix of inter-tracer correlations for all region pairs in each diagnostic
271 group. A review of these matrices with an overlaid heatmap of the correlation strength indicates
272 strong correlations between synaptic density in the medial orbitofrontal and temporal lobes with
273 mGluR5 in widespread brain regions in participants with AD (Figure 3). In the CN group,
274 significant moderate to strong correlations were more isolated between hippocampal synaptic

275 density and mGluR5 in widespread brain regions (Figure 4). Similar relationships existed with
276 PVC of the PET images (Supplementary Figures 2 and 3).

277

278

279 **Discussion**

280 In this study we investigated the relationship between mGluR5 availability measured
281 with [¹⁸F]FPEB PET and synaptic density measured with [¹¹C]UCB-J binding to SV2A in early
282 AD compared to individuals with normal cognition with an initial focus on medial temporal
283 brain regions, followed by region-based whole brain analyses. We found strong correlations
284 between mGluR5 binding and synaptic density in the hippocampus and entorhinal cortex in
285 individuals with AD. This differed from the CN group where a strong positive correlation
286 between mGluR5 availability and synaptic density was present in the hippocampus, but not the
287 entorhinal cortex. In the whole brain region-based analyses, widespread significant positive
288 correlations between mGluR5 binding and synaptic density were found in the group with AD.

289 Our previous work using [¹⁸F]FPEB and [¹¹C]UCB-J PET showed that both mGluR5
290 binding and synaptic density are significantly lower in the medial temporal lobe of individuals
291 with AD with largest effect sizes in the hippocampus (13). While the AD-related reduction in
292 mGluR5 was significant only in the hippocampus the magnitude of mGluR5 was significantly
293 lower in many commonly AD-affected association cortical regions (25). The synaptic density
294 reduction due to AD was observed was of larger magnitude and more widespread in neocortical
295 brain regions (7). Our results indicate that mGluR5 and synaptic density are highly correlated
296 within a group of participants with early AD. Considering that synaptic density is highly
297 correlated with cognitive performance in a larger sample of participants with AD (45), it is

298 possible that loss of mGluR5 and SV2A are markers of disease progression that are highly
299 related due to their locations at the synapse. Interestingly, mGluR5 binding and synaptic density
300 were strongly correlated in the hippocampus, but not the entorhinal cortex in the CN group. This
301 hippocampal correlation in the CN group was similar in magnitude and not significantly different
302 in comparison to the group with AD. The meaning of this correlation in CN participants is not
303 clear, but there may be non-AD-related reductions in mGluR5 and synaptic density in this group
304 of older adults that are correlated, but not causing clinical symptoms.

305 We also investigated the within-region relationships between mGluR5 and synaptic
306 density in all individual brain ROIs. We found that mGluR5 binding and synaptic density were
307 significantly correlated with a widespread spatial extent in the AD group, but that intraregional
308 correlations were more isolated in the CN group. In addition to the possibility that some of
309 these intraregional relationships may be driven by non-AD disease processes – such as those in
310 the hippocampus – it is also possible that age-related neurodegeneration could contribute in some
311 regions. Of particular interest, we found the strongest correlation between mGluR5 binding and
312 synaptic density in the CN group exists in the caudate. In our work and the work of others, the
313 caudate has the strongest correlation between age and synaptic density, suggesting that this may
314 be a site of age-related synaptic loss (56-58). We speculated that this association may be present
315 because the caudate is the site of nerve terminals for multiple major tracts that undergo
316 substantial age-related neurodegeneration (56). Similarly, mGluR5 binding and age are most
317 strongly correlated in the caudate, although this age-related reduction in mGluR5 binding may be
318 largely mediated by brain volume loss (59).

319 There is one other study investigating the relationship between mGluR5 binding
320 measured with [¹⁸F]PSS232 PET and synaptic density measured with [¹⁸F]SynVesT-1 PET in a

321 cohort of 20 participants (10 CN and 10 AD). In this study by Wang *et al.*, they reported
322 significant correlations between mGluR5 binding and synaptic density within and between many
323 typically AD-affected regions and also performed a more speculative analysis that suggested
324 mGluR5 binding in the medial temporal lobe may mediate the association between global
325 amyloid and synaptic density in that region (60). The results of Wang *et al.* are not that surprising
326 since their analysis was performed in the entire cohort of CN and AD participants and therefore,
327 likely driven by the large differences in these groups due to the presence or absence of AD
328 pathogenesis. A strength of our paper that builds on previous findings is separate analyses in CN
329 and AD groups that may help distinguish AD from non-AD related associations between
330 mGluR5 binding and synaptic density.

331

332 **Limitations**

333 Our study has a few limitations. The diagnosis and stage of AD was determined with
334 clinical criteria and amyloid PET positivity with no assessment of brain tau accumulation that
335 may have provided a better understanding of AD pathological stage. Moreover, the relatively
336 small sample size limits our ability to detect subtle relationships when signal-to-noise ratios may
337 be low. Future studies with larger sample sizes could confirm the absence of correlations, and
338 also allow investigations into the relationship between mGluR5 and synaptic density with
339 cognition. Additionally, our study is cross-sectional which limits the ability to determine causal
340 relationships. Longitudinal assessments with both radiotracers starting at preclinical AD stages
341 would allow validation of findings and a more thorough investigation of the temporal and spatial
342 changes of mgluR5 and synaptic density due to AD progression.

343

344 **Conclusion**

345 We observed significant, strong positive correlations between mGluR5 binding and
346 synaptic density in the hippocampus and entorhinal cortex of participants with AD. Cognitively
347 normal participants showed slightly weaker but still strong positive correlations between
348 mGluR5 and synaptic density in the hippocampus only. Whole brain region-based analyses
349 suggested a more widespread pattern of positive correlations between mGluR5 binding and
350 synaptic density due to AD that was not present in older adults with normal cognition. Our
351 findings suggest that loss of mGluR5 in AD may be closely linked to AD related synaptic loss.
352 Further studies may provide insight into the role of mGluR5 at various stages of AD pathologic
353 change, expand our understanding of AD pathogenesis, and aid in the development of novel
354 biomarkers and treatments.

355

356

357 **List of abbreviations**

358

359 Metabotropic glutamate receptor subtype 5 = mGluR5; Alzheimer's disease = AD; Positron
360 emission tomography = PET, Cognitively normal = CN; Distribution Volume Ratio = *DVR*;
361 Binding Potential Non-displaceable = BP_{ND} ; Synaptic vesicle glycoprotein 2A = SV2A; Long-
362 term depression = LTD; long-term potentiation = LTP; Positive allosteric modulator = PAM;
363 Amyloid- β oligomer = A β ; Cellular prion protein = PrPc; National Institute on Aging-
364 Alzheimer's Association = NIA-AA; Clinical Dementia Rating = CDR; Mini-Mental Status
365 Examination = MMSE; Amnesic mild cognitive impairment = aMCI; Logical Memory = LM;
366 Standard deviations = SD; [^{11}C]Pittsburg Compound B = [^{11}C]PiB; Region of interest = ROI;

367 Magnetic resonance imaging = MRI; Magnetization-prepared rapid gradient-echo = MPRAGE;
368 Partial volume correction = PVC; Motion-compensation OSEM List-mode Algorithm for
369 Resolution-recovery = MOLAR; Simplified reference tissue model-2 = SRTM2; Pearson's
370 correlation coefficient = r ; False discovery rate = FDR

371

372

373 **Declarations**

374 **Ethics approval and consent to participate**

375 The study protocol was approved by the Yale University Human Investigation Committee and
376 Radiation Safety Committee. All participants provided written informed consent prior to
377 participating in the study.

378

379 **Consent for publication**

380 Not applicable

381

382 **Availability of data and materials**

383 The data used for these analyses are available from the corresponding author on reasonable
384 request.

385

386 **Competing interests**

387 Adam P. Mecca, Richard E. Carson, and Christopher H. van Dyck report grants from the
388 National Institutes of Health for the conduct of the study. Adam P. Mecca reports grants for
389 clinical trials from Eli Lilly and Janssen Pharmaceuticals outside the submitted work. Yiyun

390 Huang reports research grants from UCB and Eli Lilly outside the submitted work. Yiyun
391 Huang, Nabeel B. Nabulsi, and Richard E. Carson have a patent for a newer version of the
392 tracer. Richard E. Carson is a consultant for Rodin Therapeutics and has received research
393 funding from UCB. Richard E. Carson reports having received grants from AstraZeneca,
394 Astellas, Eli Lilly, Pfizer, Taisho, and UCB outside the submitted work. Ryan S. O'Dell reports
395 grants for clinical trials from Cognition Therapeutics and Bristol-Myers Squibb outside of the
396 submitted work. Christopher H. van Dyck reports consulting fees from Kyowa Kirin, Roche,
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407

408 **Author contributions**

409 Dr. Salardini and Dr. Mecca had full access to the data and take responsibility for the integrity of
410 the data and the accuracy of the data analysis. Study concept and design:
411 Carson, Huang, Mecca, Nabulsi, O'Dell, Salardini, van Dyck. Acquisition, analysis, or
412 interpretation of data: Carson, Mecca, O'Dell, Salardini, Tchorz, van Dyck. Drafting of the

413 manuscript: Mecca, O'Dell, Salardini, van Dyck. Critical revision of the manuscript for
414 important intellectual content: Carson, Huang, Mecca, Nabulsi, O'Dell, Salardini, van Dyck.
415 Statistical analysis: Mecca, O'Dell, Salardini. Study supervision: Mecca, Salardini, van Dyck.

416

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420

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422 **References**

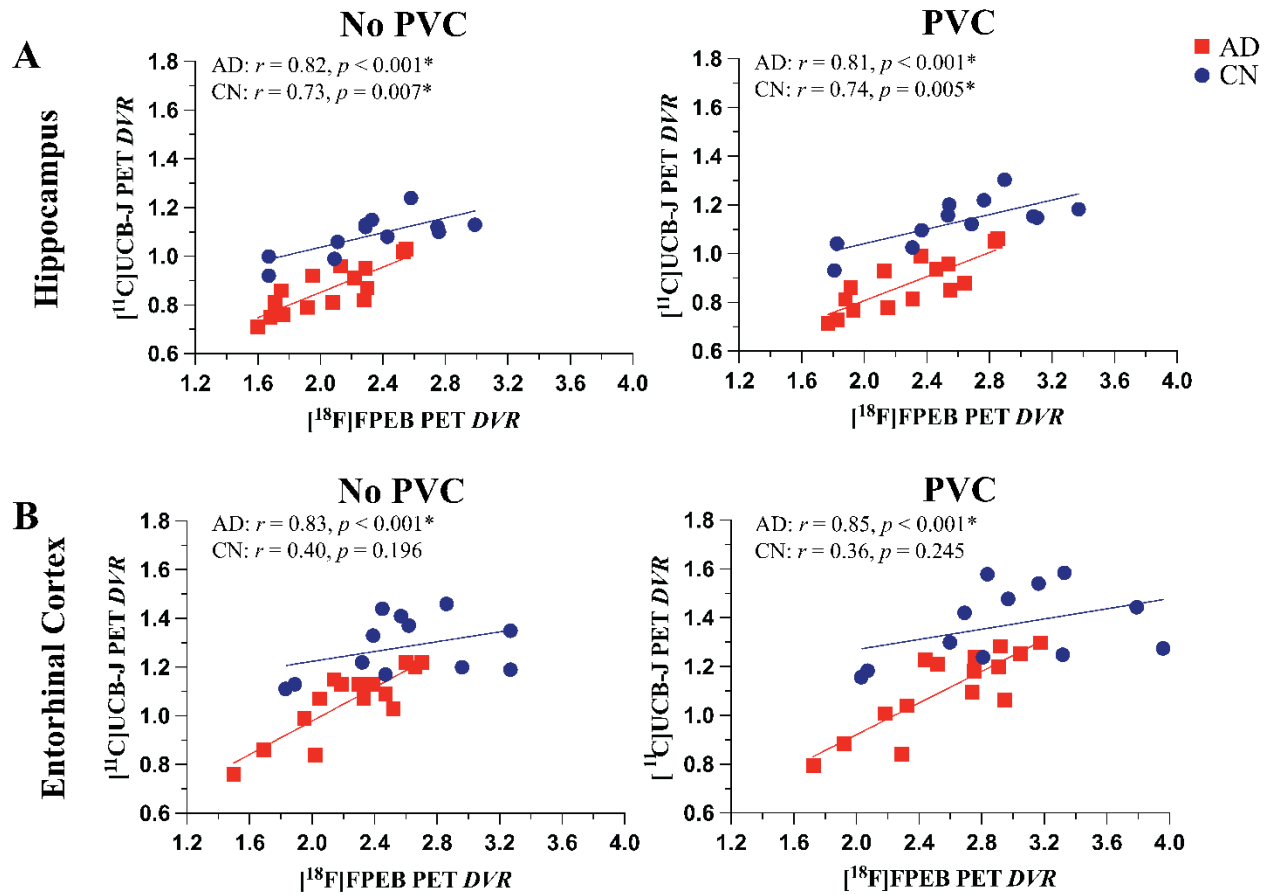
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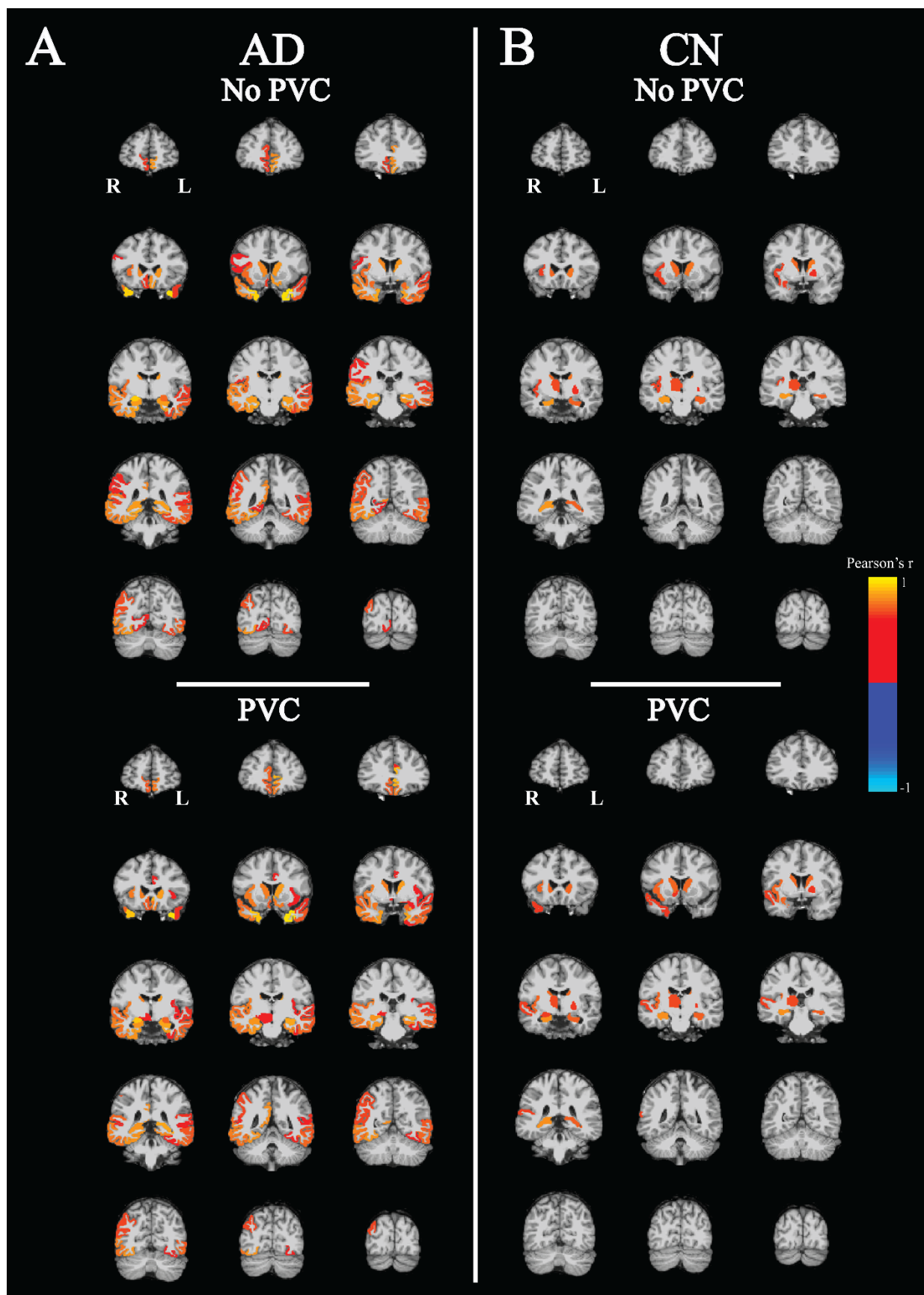
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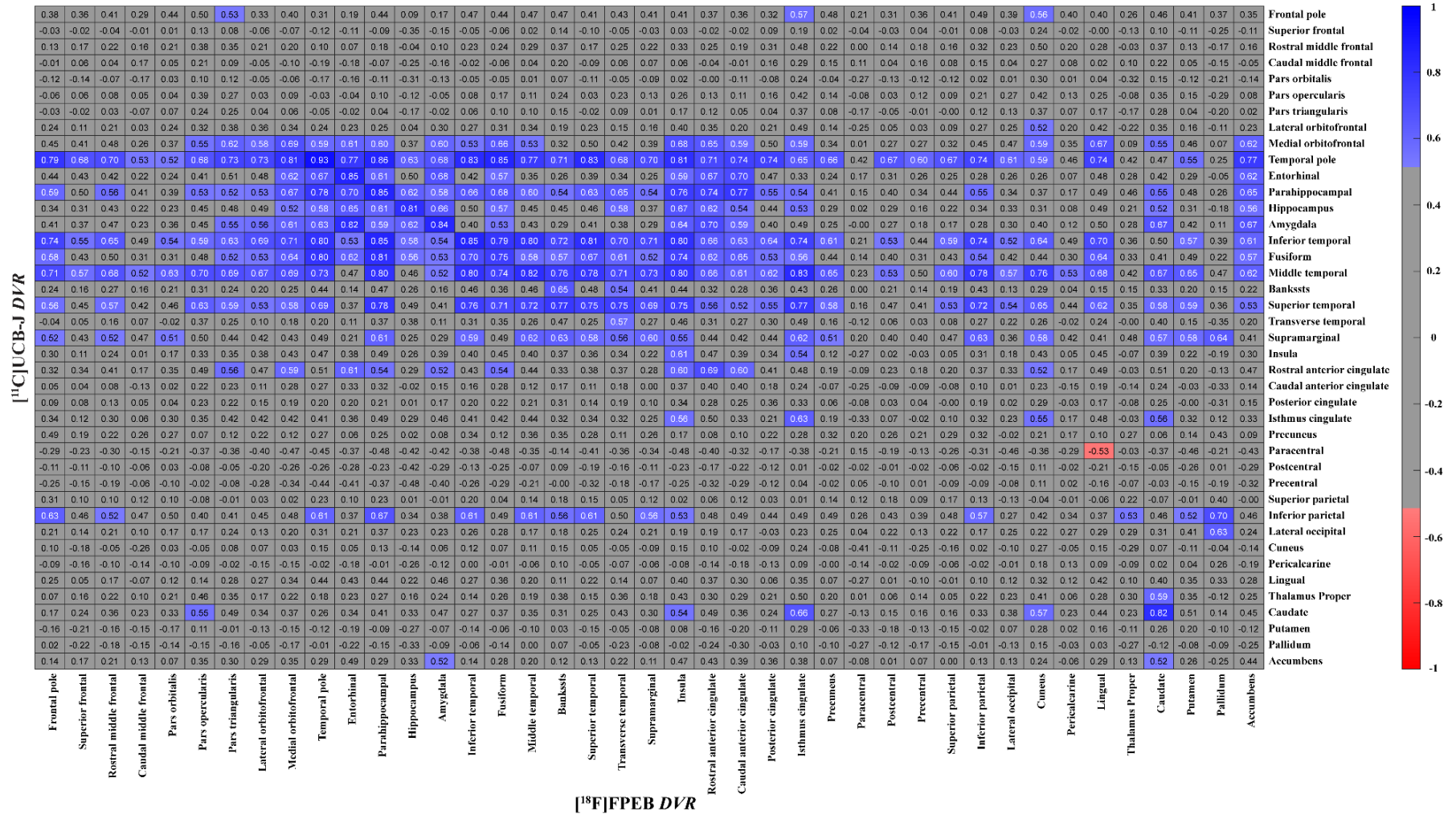
590 **Figure 1. Correlations between mGluR5 and synaptic density in the hippocampus and**
591 **entorhinal cortex.** $[^{18}\text{F}]\text{FPEB}$ (mGluR5) and $[^{11}\text{C}]\text{UCB-J}$ (synaptic density) *DVRs* are plotted
592 for participants with CN (blue, $n = 12$) and AD (red, $n = 15$). Univariate linear regression line of
593 best fit, Pearson's correlation coefficients (r) and the associated p values are shown for each
594 group for **(A)** hippocampus and **(B)** entorhinal cortex. $^*p < 0.05$. Abbreviations: AD =
595 Alzheimer's disease; CN = cognitively normal; *DVR* = Distribution volume ratio; mGluR5 =
596 metabotropic glutamate receptor subtype 5; PVC = partial volume correction



598 **Figure 2. Correlation maps of mGluR5 and synaptic density in all regions.** Pearson's
599 correlation coefficients (r) and associated p values were calculated between [^{18}F]FPEB
600 (mGluR5) and [^{11}C]UCB-J (synaptic density) PET DVR s in all regions in participants with **(A)**
601 Alzheimer's Disease ($n = 15$) and **(B)** normal cognition ($n = 12$). All voxels in each region were
602 colored uniformly for regions that had an uncorrected $p < 0.05$ and displayed as an overlay on
603 the MNI template T1 MRI. Abbreviations: AD = Alzheimer's disease; CN = cognitively normal;
604 DVR = Distribution volume ratio; mGluR5 = metabotropic glutamate receptor subtype 5; PVC =
605 partial volume correction

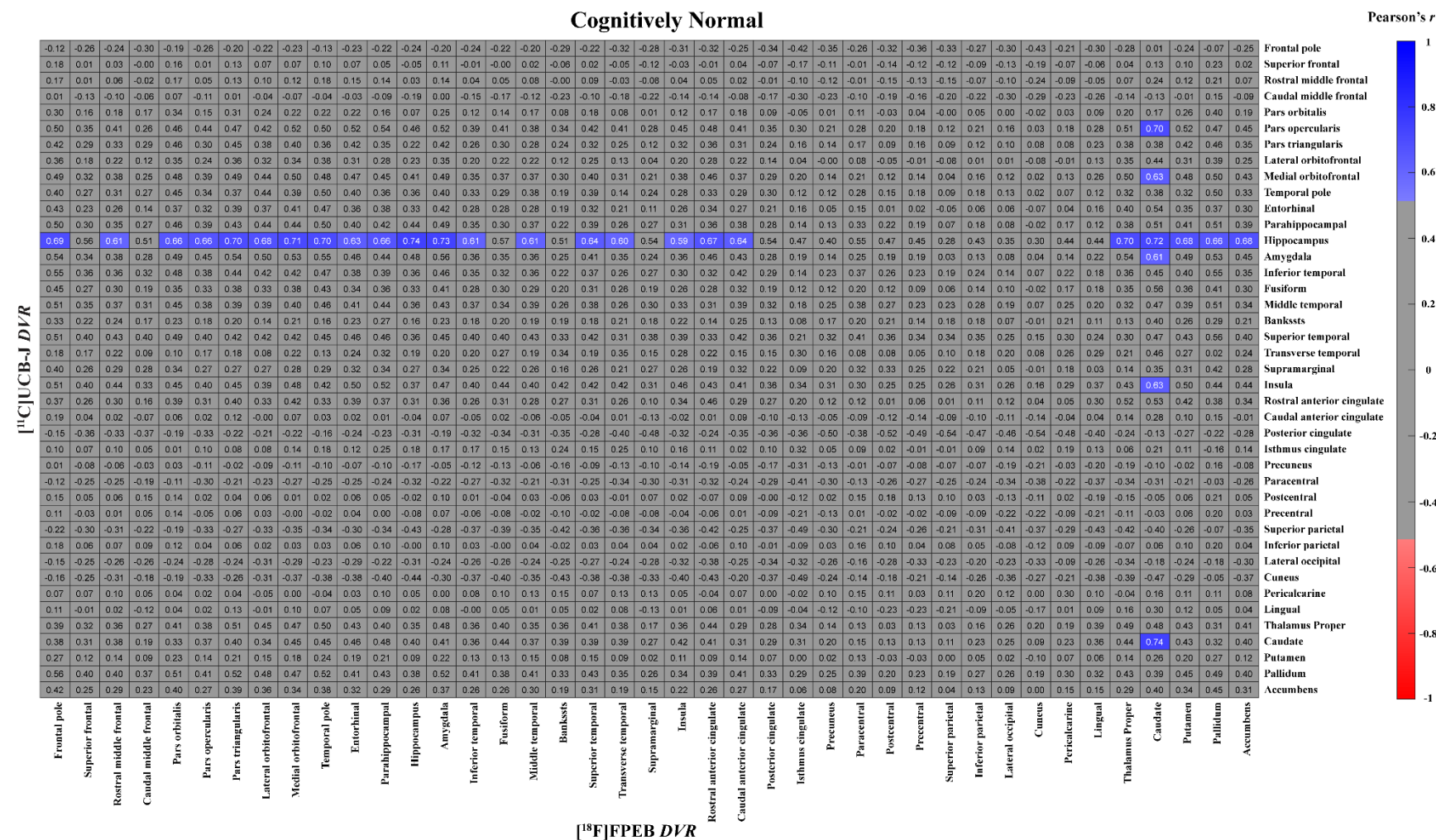
Alzheimer's Disease

Pearson's r



combinations that had an uncorrected $p < 0.05$. Abbreviations: AD = Alzheimer's disease; CN = cognitively normal; $DVR =$

Distribution volume ratios; mGluR5 = metabotropic glutamate receptor subtype 5; PVC = partial volume corrected



combinations of regions. Data are from 12 cognitively normal participants. The heat map shows the r for all combinations that had an uncorrected $p < 0.05$. Abbreviations: AD = Alzheimer's disease; CN = cognitively normal; DVR = Distribution volume ratios; mGluR5 = metabotropic glutamate receptor subtype 5; PVC = partial volume corrected

Table 1. Participant characteristics

	CN (Aβ-)	AD (Aβ+) 	CN vs. AD
Sex (Male/Female)	6/6	7/8	$\chi^2(1) = 0.03, p = 0.863$
Age (years)	70.1 \pm 8.2	73.5 \pm 5.9	$t(25) = 1.27, p = 0.213$
Education (years)	17.5 \pm 2.1	16.8 \pm 2.4	$t(25) = 0.78, p = 0.441$
CDR-global	0.0 \pm 0.0	0.7 \pm 0.2	$U = 180, p < 0.001^*$
CDR-SB	0.0 \pm 0.0	4.0 \pm 2.2	$U = 180, p < 0.001^*$
MMSE	29.1 \pm 1.3	24.1 \pm 3.9	$t(17.7) = 4.59, p < 0.001^*$
LMII	14.0 \pm 4.2	1.9 \pm 2.6	$t(25) = 9.16, p < 0.001^*$
APOE Genotype ($\epsilon 2 \epsilon 3, \epsilon 3 \epsilon 3, \epsilon 3 \epsilon 4, \epsilon 4 \epsilon 4$)	3, 7, 2, 0	0, 4, 8, 3	$\chi^2(3) = 0.61, p = 0.012^*$
Composite [^{18}F]FPEB PET DVR	2.92 \pm 0.44	2.81 \pm 0.30	$t(25) = 0.81, p = 0.424$
Composite [^{11}C]UCB-J PET DVR	1.70 \pm 0.09	1.57 \pm 0.07	$t(25) = 3.95, p < 0.001^*$
Hippocampal [^{18}F]FPEB PET DVR	2.33 \pm 0.41	2.05 \pm 0.31	$t(25) = 2.02, p = 0.054$
Hippocampal [^{11}C]UCB-J PET DVR	1.09 \pm 0.08	0.86 \pm 0.10	$t(25) = 6.18, p < 0.001^*$

Mean \pm standard deviation (continuous variables) or frequency (categorical variables) are shown for the group with normal cognition (n = 12) and Alzheimer's disease (n = 15). Test statistics, degrees of freedom, and associated *p* values are reported for independent two-tailed t tests, Mann-Whitney U tests, or a χ^2 test. **p* < 0.05. Abbreviations: A β = amyloid β ; AD = Alzheimer's disease; CDR-global = clinical dementia rating global score; CDR-SB = clinical dementia rating sum of boxes; CN = cognitively normal; DVR = Distribution volume ratio; LMII = Logical Memory delayed recall; MMSE = Mini-Mental Status Exam

Table 2. Regional correlations between mGluR5 and synaptic density (no PVC)

Region	Left hemisphere						Right hemisphere					
	AD		CN		Fisher's r to z		AD		CN		Fisher's r to z	
	r	p	r	p	z	P	r	p	r	p	z	P
Frontal pole	0.30	0.283	0.09	0.771	0.48	0.631	0.40	0.143	-0.37	0.229	1.85	0.064
Superior frontal	-0.03	0.905	0.05	0.881	0.19	0.852	0.02	0.927	-0.02	0.939	0.11	0.909
Rostral middle frontal	0.13	0.650	-0.08	0.804	0.47	0.636	0.31	0.255	0.15	0.635	0.39	0.699
Caudal middle frontal	0.10	0.715	-0.17	0.590	0.63	0.528	0.39	0.152	0.08	0.813	0.76	0.449
Pars orbitalis	0.23	0.406	0.49	0.108	0.67	0.503	-0.10	0.720	0.19	0.554	0.66	0.506
Pars opercularis	0.21	0.453	0.36	0.244	0.38	0.702	0.52	0.048*	0.44	0.149	0.22	0.827
Pars triangularis	0.24	0.397	0.33	0.287	0.24	0.807	0.37	0.171	0.47	0.126	0.26	0.796
Lateral orbitofrontal	0.40	0.136	0.12	0.714	0.70	0.483	0.29	0.290	0.46	0.136	0.43	0.663
Medial orbitofrontal	0.79	<0.001*	0.44	0.147	1.35	0.178	0.59	0.020*	0.49	0.101	0.31	0.759
Temporal pole	0.93	<0.001*	0.36	0.247	2.99	0.003*	0.91	<0.001*	0.53	0.076	2.12	0.034*
Entorhinal	0.81	<0.001*	0.46	0.127	1.42	0.155	0.83	<0.001**	0.18	0.575	2.27	0.023*
Parahippocampal	0.78	<0.001*	0.46	0.135	1.27	0.205	0.83	<0.001*	0.36	0.245	1.82	0.068
Hippocampus	0.81	<0.001*	0.68	0.014	0.64	0.523	0.80	<0.001*	0.80	0.002*	0.05	0.960
Amygdala	0.70	0.003*	0.49	0.108	0.77	0.440	0.90	<0.001*	0.56	0.056	1.87	0.061
Inferior temporal	0.71	0.003*	0.38	0.225	1.13	0.259	0.74	0.002*	0.34	0.278	1.35	0.177
Fusiform	0.62	0.014*	0.26	0.420	1.04	0.298	0.79	<0.001*	0.29	0.355	1.75	0.081
Middle temporal	0.65	0.008*	0.23	0.464	1.23	0.220	0.76	<0.001*	0.49	0.107	1.07	0.286
Bankssts	0.57	0.026*	-0.43	0.162	2.51	0.012*	0.60	0.017*	0.46	0.131	0.45	0.652
Superior temporal	0.59	0.019*	0.25	0.435	0.97	0.329	0.74	0.002*	0.57	0.050	0.67	0.504
Transverse temporal	0.37	0.175	0.18	0.572	0.46	0.643	0.75	0.001*	0.61	0.033*	0.60	0.545
Supramarginal	0.50	0.058	0.16	0.626	0.88	0.377	0.55	0.032*	0.31	0.318	0.68	0.497
Insula	0.46	0.087	0.21	0.511	0.63	0.526	0.70	0.003*	0.61	0.035*	0.38	0.704
Rostral anterior cingulate	0.77	<0.001*	0.47	0.118	1.13	0.258	0.61	0.016*	0.27	0.387	0.96	0.336
Caudal anterior cingulate	0.47	0.079	0.19	0.557	0.72	0.473	0.37	0.176	0.03	0.915	0.80	0.424
Posterior cingulate	0.34	0.219	-0.40	0.198	1.75	0.079	0.38	0.158	-0.24	0.460	1.46	0.144
Isthmus cingulate	0.45	0.093	0.42	0.175	0.08	0.934	0.75	0.001*	0.24	0.453	1.68	0.092
Precuneus	0.27	0.335	-0.24	0.458	1.17	0.242	0.44	0.098	-0.02	0.949	1.12	0.260
Paracentral	0.07	0.811	0.002	0.993	0.15	0.883	0.28	0.315	-0.14	0.656	0.98	0.329
Postcentral	0.009	0.973	0.24	0.450	0.54	0.592	0.25	0.360	0.13	0.683	0.30	0.772
Precentral	0.06	0.837	-0.02	0.952	0.18	0.860	0.13	0.648	0.03	0.924	0.22	0.824
Superior parietal	0.20	0.468	-0.27	0.397	1.09	0.275	0.14	0.605	-0.10	0.763	0.55	0.580
Inferior parietal	0.27	0.325	-0.08	0.807	0.81	0.416	0.64	0.010*	0.15	0.640	1.38	0.167
Lateral occipital	0.15	0.600	-0.29	0.351	1.03	0.304	0.35	0.204	-0.16	0.628	1.18	0.238
Cuneus	0.21	0.440	-0.36	0.246	1.36	0.174	0.34	0.208	-0.03	0.926	0.88	0.376
Pericalcarine	0.20	0.477	0.38	0.225	0.44	0.656	0.44	0.102	0.21	0.512	0.58	0.559
Lingual	0.28	0.302	-0.02	0.944	0.72	0.472	0.54	0.037*	0.23	0.477	0.85	0.395
Thalamus	0.26	0.344	0.25	0.431	0.03	0.978	0.35	0.203	0.63	0.026*	0.87	0.381
Caudate	0.79	<0.001*	0.72	0.008	0.35	0.722	0.76	<0.001*	0.70	0.012*	0.33	0.738
Putamen	0.24	0.397	0.21	0.504	0.05	0.958	0.12	0.656	0.18	0.563	0.14	0.888
Pallidum	0.18	0.526	0.58	0.049	1.08	0.279	0.08	0.765	0.37	0.232	0.70	0.485
Accumbens area	0.33	0.230	0.07	0.820	0.61	0.542	0.41	0.126	0.57	0.052	0.48	0.632
Ventral Diencephalon	0.24	0.395	0.009	0.976	0.52	0.599	0.50	0.054	0.09	0.768	1.05	0.295

Pearson's r and associated p value is reported for the correlation between [^{18}F]FPEB (mGluR5) and [^{11}C]UCB-J PET (synaptic density) DVR in each brain region. Fisher r -to- z transformation was used to compare correlation coefficients of CN and AD groups. The data were from 12 CN participants 15 participants with AD. * $p < 0.05$.

Abbreviations: AD = Alzheimer's disease; CN = cognitively normal; *DVR* = Distribution volume ratio; PVC = Partial volume correction