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### Research Articles: Behavioral/Cognitive

### Grey Matter Volume Differences Associated with Extremely Low Levels of Cannabis Use in Adolescence

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### 145 Abstract

146 Rates of cannabis use among adolescents are high, and are increasing concurrent with 147 changes in the legal status of marijuana and societal attitudes regarding its use. Recreational 148 cannabis use is understudied, especially in the adolescent period when neural maturation may 149 make users particularly vulnerable to the effects of  $\Delta$ -9-tetrahydrocannabinol (THC) on brain 150 structure. In the current study, we used voxel-based morphometry to compare grey matter 151 volume (GMV) in 46 fourteen year old human adolescents (males and females) with just one or 152 two instances of cannabis use and carefully matched THC-naïve controls. We identified 153 extensive regions in the bilateral medial temporal lobes as well as the bilateral posterior 154 cingulate, lingual gyri, and cerebellum that showed greater GMV in the cannabis users. Analysis 155 of longitudinal data confirmed that GMV differences were unlikely to precede cannabis use. 156 GMV in the temporal regions was associated with contemporaneous performance on the 157 Perceptual Reasoning Index and with future generalized anxiety symptoms in the cannabis 158 users. The distribution of GMV effects mapped onto biomarkers of the endogenous cannabinoid 159 system providing insight into possible mechanisms for these effects.

### Significance Statement

Almost 35% of American 10<sup>th</sup> graders have reported using cannabis and existing research 162 163 suggests that initiation of cannabis use in adolescence is associated with long-term 164 neurocognitive effects. We understand very little about the earliest effects of cannabis use, 165 however, as most research is conducted in adults with a heavy pattern of lifetime use. This 166 study presents evidence suggesting structural brain and cognitive effects of just one or two 167 instances of cannabis use in adolescence. Converging evidence suggests a role for the 168 endocannabinoid system in these effects. This research is particularly timely as the legal status 169 of cannabis is changing in many jurisdictions and the perceived risk by youth associated with 170 smoking cannabis has declined in recent years.

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### 175 Introduction

177 Preclinical evidence has consistently demonstrated a causal relationship between cannabis 178 exposure and changes to brain morphology (refer(Panlilio and Justinova, 2018) for review). The 179 human evidence, however, has been variable reporting both increases and decreases in brain 180 volumes (Ashtari et al., 2011; Cousijn et al., 2012; Gilman et al., 2014), no volume differences 181 (Jager et al., 2007; Weiland et al., 2015; Gillespie et al., 2018), and modest effect sizes 182 (Weiland et al., 2015). Factors including the age of cannabis use initiation, comorbid substance 183 use, and levels of use are believed to contribute to variability in the human findings (Curran et 184 al., 2016). 185

186 Most neuroimaging research is conducted in adults with a heavy, chronic pattern of cannabis 187 use and does not reflect most people's experience, which is recreational (SAMHSA, 2014). 188 Dose-dependent associations with brain volumes have been reliably identified in preclinical 189 studies (refer (Lorenzetti et al., 2010) for review) with some evidence of the same in humans 190 (Battistella et al., 2014; French et al., 2015), suggesting consequences of lower levels of use. 191 One study has reported differences in grey matter density and shape of the amygdala and 192 nucleus accumbens in recreational cannabis users (Gilman et al., 2014), but subsequent 193 research has suggested that these findings may be associated with alcohol (Weiland et al., 194 2015) and nicotine (Gillespie et al., 2018) exposure in the cannabis users. 195

196 One mechanism by which cannabis may produce neurobiological changes is through the 197 endogenous cannabinoid system (eCB). The amygdala, hippocampus, striatum, and cerebellum 198 (Lorenzetti et al., 2016) are regions most frequently showing structural brain correlates of 199 cannabis use and are also components of the eCB system (Burns et al., 2007); the preclinical 200 literature suggests a causal role of this system in the effects of cannabis on brain morphology 201 (Downer et al., 2001). The eCB system mediates maturation-related neural reorganisation 202 (Fernández-Ruiz et al., 2000), which may place adolescents at heightened vulnerability to 203 structural brain effects of cannabis exposure as adolescence is a time of rapid neural maturation 204 (Rubino and Parolaro, 2008). Consistent with this suggestion, those who commenced cannabis 205 use in adolescence typically show greater structural brain differences than those who initiated 206 use in adulthood (Battistella et al., 2014; Lubman et al., 2015). These findings may also have 207 been influenced by the effects of other substances, however, as one study comparing 208 adolescent daily cannabis users with controls matched for alcohol and nicotine use found no 209 differences in subcortical grey matter density or morphology (Weiland et al., 2015).

210 In the present study we identified participants with just one or two instance of cannabis use from 211 a very large, population sample of adolescents (IMAGEN, n = 2400 (Schumann et al., 2010)) 212 and control participants matched on a range of variables, including alcohol and nicotine 213 consumption. We predicted that even extremely low levels of cannabis use would be associated 214 with structural brain differences in regions previously implicated in cannabis use studies and in 215 the eCB system: the amygdala, hippocampus, striatum, and cerebellum. We adopted a whole 216 brain, Voxel Based Morphometry (VBM) approach as it allows us to also test more extensive 217 regions of the eCB system including the frontal cortex and posterior cingulate (Burns et al., 218 2007). We explored whether grey matter volume (GMV) predicted behavioral features 219 previously associated with cannabis use and with the eCB system.

- 220 To test whether observed differences between cannabis users and controls may precede 221 cannabis use, we also identified participants who were cannabis-naïve at the time of imaging
- 222 but went on to use cannabis two years later and matched controls who remained abstinent.
- 223 Finally, in order to demonstrate association with the eCB system, we compared the spatial

224 distribution of GMV effects with two biomarkers of the eCB system using CB1 receptor

225 availability taken from a previously published, independent sample (D'Souza et al., 2016) and 226 the expression of the CNR1 gene, which encodes this receptor, taken from the Allen Human 227 Brain Atlas (Hawrylycz et al., 2012).

### Materials and Methods 228

229 Standard Operating Procedures: Standard operating procedures for the IMAGEN project are 230 available at https://imagen-europe.com/resources/standard-operating-procedures/ and contain 231 details on ethics, recruitment, and assessment.

### 232 233 Participants

234 Data were acquired from a large sample of adolescents recruited through high schools in four 235 European countries for the IMAGEN project (http://www.imagen-europe.com). Recruitment into 236 the IMAGEN study was managed through eight sites and targeted adolescents for whom all four 237 grandparents were the same nationality as the participant; as such, the sample is racially and 238 ethnically homogenous. Raw, T1-weighted images were visually inspected for the presence of 239 anatomical abnormalities or artifacts including head motion or reconstruction errors. After VBM 240 processing, images were again inspected for any errors in tissue segmentation or normalization 241 into MNI space. Images failing quality control for any reason were excluded. 242

243 Cohort 1: Forty-seven participants reported low levels of cannabis use at baseline (only one or 244 two lifetime instances of use) and complete demographics to facilitate matching; one participant 245 was excluded due to poor scan quality, leaving 46 adolescent cannabis using participants. The 246 groups were matched on age, sex, handedness, pubertal development, intelligence quotient 247 (IQ: verbal comprehension and perceptual reasoning index scores), socioeconomic status 248 (SES), total GMV, alcohol use, and nicotine use across group. All participants denied any other 249 illicit substance use, and none reported using the fictional control substance, relevin, supporting 250 the integrity of the self-report metrics. Table 1 summarizes the demographic information. We also ensured that similar numbers of cannabis users and controls were selected from each site 252 (Mann-Whitney U tests, Table 1) and confirmed that the proportion of cannabis users and 253 controls did not differ by site using a Kruskal-Wallis Test ( $\chi^2$  (6) = 5.919, p = 0.432). 254

255 For a subset of the 14-year-old cannabis-using participants, data were available at two-year 256 follow-up for substance use, cognitive ability, and psychopathology at age 16 to allow us to 257 assess the implications of cannabis-related GMV differences for future functioning in these 258 domains. Table 2 summarizes the demographic information for this subset of participants. 259

260 Cohort 2: In order to determine whether group differences between cannabis users and 261 matched controls may have preceded cannabis use, we also identified participants who were 262 cannabis-naïve at the age 14 baseline assessment but reported at least ten instances of 263 cannabis use by follow-up two years later. Sixty-nine participants who were cannabis-naïve at 264 baseline but with at least ten instances of cannabis use by follow-up provided complete 265 demographic data and all had GMV data that passed QC. Sixty-nine controls matched by group 266 on the same demographic measures as above and who reported no cannabis use at baseline or 267 follow-up were also identified. All participants denied any other illicit substance use at baseline 268 and follow-up. Table 3 summarizes the demographic information for this sample of participants. 269 We again ensured that similar numbers of cannabis users and controls were selected from each 270 site (Mann-Whitney U tests, Table 3) and confirmed that the proportion of cannabis users and 271 controls did not differ by site using a Kruskal-Wallis Test ( $\chi^2$  (7) = 4.633, p = 0.705).

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For both cohorts, the control subjects were selected from a larger pool of IMAGEN participants with T1 images that passed QC and who reported no illicit substance use. This selection was done using Python scripts written in our laboratory to randomly select subjects and compare them with the sample of cannabis users on nominated characteristics (in this case: age, sex, handedness, site (dummy coded as 8 binary variables), pubertal development, VCIQ, PRIQ, SES, total GMV, alcohol use, and nicotine use) without experimenter intervention.

### 280 Substance Use Measures

281 Substance use was assessed at baseline (age 14) and follow-up (age 16) via the European 282 School Survey Project on Alcohol and Drugs (ESPAD (Hibell et al., 2004)), a self-report 283 questionnaire that measures use of alcohol, nicotine, cannabis, inhalants, tranquilisers, 284 amphetamines, Lysergic acid diethylamide (LSD), magic mushrooms, crack, cocaine, heroin, 285 narcotics, Methylenedioxymethamphetamine (MDMA), ketamine, y-Hydroxybutyric acid (GHB), 286 anabolic steroids, and a fictional control measure (relevin). Participants indicated how frequently 287 they had used each of the substances in their lifetime, in the past 12 months, in the past 30 288 days, and in the past 7 days using a 7-point scale (0: never, 1: 1-2 times, 2: 3-5 times, 3: 6-9 289 times, 4: 10-19 times, 5: 20-39 times, and 6: 40 or more times); they also indicated the age at 290 which they had first tried each of the substances. 291

292 Cohort 1 comprised those participants with an ESPAD of 1 for cannabis (i.e. 1-2 instances of 293 cannabis use) and no reported use of any other illicit substances, and matched controls with no 294 cannabis use and no use of any other illicit substances. We also extracted lifetime alcohol and 295 nicotine use from the ESPAD in order to match the groups on these variables. In order to 296 explore possible relationships between GMV and cannabis use metrics, we also extracted from 297 the ESPAD age of first use, frequency of use in the past 30 days, and lifetime use by age 16 for 298 those who reported cannabis use at baseline.

Cohort 2 comprised those participants with an ESPAD of 0 for cannabis at baseline, an ESPAD of 4, 5, or 6 for cannabis at follow-up (i.e. cannabis-naive at age 14 and with 10+ instances of cannabis use by age 16) and no reported use of any other illicit substances at either baseline or follow-up, and matched controls with no cannabis use and no use of any other illicit substances at either time point. We also extracted lifetime alcohol and nicotine use from the ESPAD in order to match the groups on these variables.

### 307 Demographic Measures

Biological sex was determined by karyotype analysis (chromosome 23: XX=female, XY=male).
Participants provided blood samples, which were shipped to the Institute of Psychiatry, London
for genotyping with Illumina Human610-Quad Bead Chips (Illumina, San Diego, CA, USA). DNA
extraction was performed by a semiautomated process to ensure high quality and sufficient
quantity.(Schumann et al., 2010)

SES was indexed by a score that summed: Mother's Education Score, Father's Education Score, Family Stress Unemployment Score, Financial Difficulties Score, Home Inadequacy Score, Neighborhood Score, Financial Crisis Score, Mother Employed Score, and Father Employed Score from the parent report of the Development and Well-Being Assessment interview (DAWBA (Goodman et al., 2000), see also <u>http://www.dawba.info</u>).

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- 320 Participants completed the Perceptual Reasoning, Matrix Reasoning, Similarities and
- 321 Vocabulary subscales from the Wechsler intelligence scale for children WISC-IV(Wechsler,
- 322 1949) to generate Verbal Comprehension (VCIQ) and Perceptual Reasoning (PRIQ) indices.

323 Physical maturity was assessed using the Pubertal Development Scale (Petersen et al., 1988),

324 a self-report measure of physical signs associated with the onset, progression, and completion 325 of puberty.

### 326 Personality and Temperament Measures

327 Personality was assessed with the self-reported Substance Use Risk Profile Scale 328 (SURPS; Woicik, Stewart, Pihl, & Conrod, 2009), the NEO Five Factor Inventory (NEO-329 FFI; Costa Jr & McCrae, 1992), and the Temperament and Character Inventory (TCI; Cloninger, 330 Przybeck, Svrakic, & Wetzel, 1994). The SURPS produced summary measures for personality 331 traits of hopelessness, anxiety sensitivity, impulsivity, and sensation-seeking. The NEO-FFI 332 produced summary measures for five higher-order personality characteristics: neuroticism, 333 conscientiousness, extraversion, agreeableness, and openness to experience. The TCI 334 produced measures for exploratory excitability vs. stoic rigidity, impulsiveness vs. reflection, 335 extravagance vs. reserve, disorderliness vs. regimentation, and a novelty seeking summary 336 statistic. 337

### Cognitive Measures

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Delay discounting was assessed with the Monetary Choice Questionnaire (Kirby, 2009) that
required participants to complete 27 two-alternative forced choice items in which they indicated
whether they would prefer a "smaller sooner" or a "larger later" reward (e.g., "Would you prefer
€14 today or €25 in 19 days?"). The summary k statistic indexes the degree to which a
participant discounts more temporally remote rewards.

Psychomotor speed and manual dexterity were assessed using the Perdue Pegboard (Tiffin,
Participants were asked to place as many pins as possible in the small holes on the test
board in 30 seconds. Participants completed three trials in each of three conditions: using only
the dominant hand; only the non-dominant hand; and both hands.

Spatial working memory and decision-making were assessed using the Cambridge
Neuropsychological Test Automated Battery (CANTAB (Robbins et al., 1994)). We examined
the number of memory failures made during a visual search task and the risk-taking summary
statistic from a gambling task.

### 355 Psychopathology Measures

Psychiatric symptoms of conduct disorder, oppositional defiant disorder, attention deficit/
 hyperactivity disorder, generalized anxiety, depression, specific phobia, social
 phobia, agoraphobia, panic disorder, obsessive compulsive disorder and eating disorders were
 assessed via the DAWBA, which was administered to participants and their parents at baseline

assessed via the DAWBA, which was administered to participants and their parents at baseline and at follow-up. Computer generated band scores integrated reported symptoms and their impact with the approximate prevalence rates in an epidemiological sample for each disorder and reflect the likelihood that the participant would be diagnosed with the disorder in question (ranging from 0 to 5). Diagnostic criteria were based on the *Diagnostic Statistical Manual*, Version 4.

### 366 Neuroanatomical MRI acquisition

367 MRI scanning was conducted at the eight IMAGEN assessment sites using 3T whole body MRI 368 systems (Siemens: 4 sites, Philips: 2 sites, General Electric: 1 site, and Bruker: 1 site). A high-769 resolution, three-dimensional T1-weighted image was acquired using a magnetization prepared 370 gradient echo sequence based on the ADNI protocol (<u>http://adni.loni.usc.edu/methods/mri-</u>

<u>tool/mri-analysis/</u>), which specifies protocols designed to minimize differences in image contrast
 and signal-to-noise across scanner makes and models. Two additional quality control

procedures were regularly implemented: (1) the American College of Radiology phantom was
scanned every two months at each site and after every hardware and software upgrade to
provide information about geometric distortions and signal uniformity related to hardware
differences in radiofrequency coils and gradient systems, image contrast, and temporal stability;
and (2) twice per year at each site and after any hardware or software upgrade, human
volunteers were scanned to determine inter-site variability in raw MRI signal and tissue
relaxation properties. (Schumann et al., 2010).

### 381 Voxel-Based Morphometry:

382 T1-weighted images were processed using the Statistical Parametric Mapping version 8 (SPM8) 383 (http://www.fil.ion.ucl.ac.uk/spm/software/spm8/) VBM toolbox (http://dbm.neuro.uni-384 jena.de/vbm/) with default parameters incorporating the DARTEL toolbox implemented in 385 MATLAB 7.0 (MathWorks, Natick MA, USA). Image processing comprised iterative tissue 386 segmentation and spatial normalization using both linear (12-parameter affine) and non-linear 387 transformations (Ashburner and Friston, 2000; Ashburner, 2007) without skull stripping. SPM8 388 default settings were used to be consistent with other VBM publications from the IMAGEN 389 Consortium. To preserve information about absolute volume, the gray matter concentration 390 images were modulated by multiplying by the linear and non-linear components of the Jacobian 391 determinants generated during spatial normalization. Thus, the dependent measure in the 392 subsequent analysis was absolute gray matter volume. Voxel resolution after normalization was 393 1.5 x 1.5 x 1.5 mm. To make the residuals in later analyses conform more closely to a Gaussian 394 distribution and to account for individual differences in brain anatomy, the modulated GM 395 images were smoothed with an isotropic Gaussian kernel of 8 mm full-width at half maximum.

### 397 Experimental Design and Statistical Analyses

398 Whole brain voxel-wise analyses were conducted using the general linear model, implemented 399 in AFNI (Cox, 1996). We tested for GMV differences at baseline between: (1) Cohort 1, those 46 400 participants who reported low levels of cannabis use at baseline and their matched controls; and 401 (2) Cohort 2, those 69 participants who reported cannabis use by age 16 and their matched 402 controls. Age, sex, handedness, and total GMV were included in the models as covariates of no 403 interest. Imaging site was included as an additional covariate; given the cohort sizes and large 404 number of covariates already used, additional measures inter-site imaging variance were not 405 included in this analysis. Type 1 error was controlled using a combination of voxel-level 406 significance and cluster extent: following Eklund and colleagues (Eklund et al., 2016), the 407 updated AFNI program 3dTTest++ with the option –clustsim

416 We also conducted region-of-interest analyses in cohort 2 in which we extracted GMV from the 417 regions showing significant volume differences between baseline users and controls to confirm 418 that GMV differences in these specific regions did not precede cannabis use. Note that these 419 regions were defined by the analysis of cohort 1 (n = 46), and then tested on an independent 420 cohort (cohort 2, n = 69).

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422 A series of *post hoc* analyses were conducted to ensure that group differences in GMV between 423 baseline users and controls could not be accounted for by any differences in cognitive ability, 424 personality, or symptoms of psychopathology. Independent groups t-tests were used to test for
425 differences in the continuous variables and Mann-Whitney U tests were used to test for
426 differences in the ordinal DAWBA band scores. We did not correct for multiple comparisons for
427 these tests so as to have a liberal threshold for identifying any group differences. We then
428 repeated the voxel-wise GMV analyses with any behavioral variables that differed between the
429 groups included as additional covariates.

431 We explored whether individual differences in GMV in those regions that differed between 432 cannabis users and controls were associated with substance use factors (lifetime alcohol or 433 nicotine consumption, recent cannabis use, or age of onset of cannabis use) in those 434 participants reporting cannabis use at baseline. We also assessed whether GMV in regions that 435 differed between cannabis users and controls were associated with individual differences in 436 specific cognitive and psychopathological domains previously related to cannabis misuse in 437 those participants reporting cannabis use. Spatial working memory, risk-taking, delay 438 discounting, psychomotor speed, depression, generalized anxiety, and ADHD were assessed at 439 baseline. For a subset of those participants reporting cannabis use at baseline, 440 psychopathology (n = 33), delay discounting (n = 31), and substance misuse data (n = 31) were 441 also available at follow-up two years later. We assessed whether regional GMV at baseline 442 predicted symptoms of depression, generalized anxiety, or ADHD; delay discounting; or future 443 cannabis use. For all post hoc analyses, regional GMV was normalized by total GMV. 444

445 <u>Cannabinoid 1 Receptor Availability:</u> In order to test for associations between the spatial
 446 distribution of group differences in GMV and a receptor for the eCB system, we used a map of
 447 CB<sub>1</sub> receptor availability generated from the healthy control participants in a previously
 448 published study (D'Souza et al., 2016). Maps of CB<sub>1</sub> receptor availability were generated using
 449 positron emission tomography and the reversible ligand [11C]OMAR in 21 adult males aged 18 450 35 (D'Souza et al., 2016), the 21 individual participant maps were averaged to provide an
 451 estimate of CB<sub>1</sub> receptor availability at each voxel.

The map of the GMV comparison between cannabis users and controls was down-sampled to the resolution of the PET map (3x3x3mm<sup>3</sup> voxels) and Spearman correlations were conducted between the t-statistic at each voxel and the average CB<sub>1</sub> receptor availability at the same site using the AFNI program 1dCorrelate. First, we tested all voxels within a grey matter mask; we then tested only those voxels within regions showing significant GMV differences between cannabis users and controls.

460 Gene Expression: Associations between the spatial distribution of group differences in GMV and 461 expression of the gene that encodes the CB<sub>1</sub>R were tested with reference to the Allen Human 462 Brain Atlas (Hawrylycz et al., 2012). Using the alleninf toolbox, (Gorgolewski et al., 2014) we 463 extracted normalized gene expression values for CNR1 (averaged within spherical ROIs with 464 radii of 3mm) from within a grey matter mask and then used random-label permutation to test for 465 an association between CNR1 expression and the t-statistic of GMV effects. Distributions of 466 Spearman correlations between 50 randomly selected genes and the t-statistics of GMV effects 467 were obtained by 5,000 bootstrap resamples and then merged to build a null model. The 95% 468 confidence interval of this null distribution was calculated as the cut-off point against which the 469 strength of the association between GMV effects and CNR1 gene expression was assessed. 470 The list of randomly chosen genes, their expression at each sampling site, the expression of 471 CNR1, and the GMV t-statistic at each sampling site are available in the Extended Data.

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4 <u>Results</u>

476 Cohort 1: Group Differences in GMV associated with low rates of cannabis use 477 Figure 1 illustrates extensive regions of greater GMV in those participants who reported low 478 levels of cannabis use relative to matched controls. Bilateral medial temporal regions, including 479 the hippocampus, the amygdala, and the striatum, and bilateral parietal regions were implicated, 480 as were regions of the cerebellum and the left middle temporal gyrus (Table 4). Due to the 481 relevance of the striatal sub-regions, especially the ventral striatum, for addiction and substance 482 use, Table 5 details the number of voxels (and proportion of volume) implicated in each of the 483 putamen, caudate, and ventral striatum as defined by the Oxford-GSK-Imanova structural 484 striatal atlas (Tziortzi et al., 2011). Figure 2 illustrates the distribution of regional GMV, 485 normalized by total GMV, for those regions at which GMV differed between cannabis users and 486 controls.

488 Of all the variables describing cognitive ability, symptoms of psychopathology, and personality, 489 only agoraphobia (U = 868.00, puncorr = 0.038) and the sensation seeking measure from the 490 SURPS (t<sub>88</sub> = 2.824, p<sub>uncorr</sub> = 0.006) differed between the cannabis users and controls with the cannabis users reporting higher levels of both. When agoraphobia band score and sensation 491 492 seeking were included in the voxel-wise analysis as covariates, the three clusters reported in 493 Figure 1 and table 4 were still observed (albeit, with a small reduction in volume that may be 494 accounted for by the reduction in power due to the addition of extra covariates). One additional 495 cluster centered on the left inferior temporal gyrus (Table 6) was also revealed in this analysis 496 as showing significantly greater GMV in the cannabis users than the controls. 497

498 Cohort 1: Associations Between GMV and Contemporaneous Behavioral Measures

499 In light of the individual differences in normalized GMV effects in the cannabis using group, we 500 conducted post hoc analyses to explore whether any of the demographic variables on which the 501 groups were matched was associated with GMV in the regions of interest for those adolescents 502 reporting cannabis use. Age was not associated with normalized GMV in any of the identified 503 ROIs; the difference between males and females in GMV in the bilateral parietal cluster 504 approached but did not reach the corrected significance level ( $t_{44} = 2.226 p_{uncorr} = 0.031$ ) with 505 normalized GMV greater in males than females. When controlling for handedness, sex, and 506 age, normalized GMV in the left and right temporal clusters ( $r_{41} = -0.411$ ,  $p_{corr} = 0.037$  and  $r_{41} = -0.411$ . 507 0.457, p<sub>corr</sub> = 0.012, respectively) were negatively associated with PRIQ such that greater 508 relative volume in these regions was associated with reduced PRIQ (Figure 3). VCIQ, PDS, 509 SES, alcohol use, and nicotine use were not associated with GMV in any of the identified ROIs. 510 The cannabis use metrics (age of use or whether cannabis was used in the last month) were not 511 associated with GMV. 512

513 Of the specific cognitive and psychological domains assessed at baseline, only psychomotor 514 speed showed an association with GMV (Figure 4): normalized GMV in the left temporal cluster 515 showed a negative association with the number of pegs placed with the non-dominant hand ( $r_{39}$ 516 = -0.454,  $p_{corr}$  = 0.030). 517

518 Cohort 2: Associations Between GMV and Future Cannabis Use

519 There were no regions at which GMV differed between the future cannabis users and their 520 matched controls. Region of interest analyses focused on those regions from cohort 1 that 521 differed between baseline users and matched controls also revealed no significant differences 522 between future cannabis users and matched controls (Table 7).

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Cohort 1: Associations Between GMV and Future Behavioral Measures

A post hoc Mann-Whitney U test showed that baseline GMV in the right temporal cluster was
 significantly greater for those cannabis users who went on to have higher levels of generalized
 anxiety (DAWBA band scores of 1 or greater vs DAWBA band scores of 0: U = 43, p<sub>corr</sub> = 0.009,
 Figure 5). No other associations between regional GMV and cognition or psychopathology
 reached significance.

531 Cohort 1: Spatial Associations Between GMV Effects and CB<sub>1</sub> Receptor Availability 532 Comparison of the t-statistic map of GMV differences between cannabis users and controls with 533 the map of average CB1 receptor availability in an independent sample (D'Souza et al., 2016) 534 showed significant (p< 0.05) spatial association (r<sub>54041</sub> = 0.1131, 95% CI: 0.10468, 0.12152). 535 Comparison of only those voxels showing a significant GMV difference between cannabis users 536 and controls also showed a significant (p < 0.05) spatial association between the magnitude of 537 GMV effects and CB<sub>1</sub> receptor availability ( $r_{1229}$  = 0.0803, 95% CI: 0.02537, 0.13444). This more 538 conservative test illustrates that even within those regions showing a significant GMV difference 539 between cannabis users and controls, the magnitude of the difference was associated with CB1 540 receptor availability. 541

542Cohort 1: Spatial Associations Between GMV Effects and CNR1 Gene Expression543Comparison of the t-statistic map of GMV differences between cannabis users and controls with544the map of CNR1 gene expression showed significant (p< 0.05) spatial association ( $r_{3685}$  =5450.311, 95% CI: 0.279, 0.341), while the null model showed no association with GMV (95% CI: -54601930, 01977). .547

### 548 Discussion

549 We present evidence of GMV differences in adolescents associated with only one or two 550 instances of cannabis use. Although novel, this work is consistent with reports of a dose-551 response effect of cannabis on behavioral and brain measures following heavier use (Lorenzetti 552 et al., 2010; Silins et al., 2014). We identified GMV increases in large medial temporal clusters 553 incorporating the amygdala, hippocampus, and striatum, extending into the left prefrontal cortex. 554 GMV increases were also observed in the lingual gyri, posterior cingulate, and cerebellum. The 555 regions identified in this whole-brain, VBM approach replicated previous findings of differences 556 in volume (Yücel et al., 2008; Ashtari et al., 2011; Schacht et al., 2012) and shape (Smith et al., 557 2013; Gilman et al., 2014; Smith et al., 2015) associated with cannabis use in region of interest 558 studies and with the spatial distribution of the eCB system (Burns et al., 2007). Although 559 cannabis use has been associated with reduced brain volumes, studies typically report on 560 adults with heavy substance use histories (cf. (Ashtari et al., 2011)). Gilman and colleagues 561 (Gilman et al., 2014), however, have reported grey matter density increases in the amygdala 562 and nucleus accumbens of young adult recreational users and Medina and colleagues (Medina 563 et al., 2007) observed hippocampal enlargement in cannabis using adolescents. Our results are 564 also consistent with the Avon Longitudinal Study of Parents and Children (French et al., 2015), 565 which showed a trend for greater cortical thickness in male adolescents with fewer than 5 566 instances of cannabis use relative to THC-naïve controls.

567 Converging evidence suggests that these effects may be a consequence of cannabis exposure. 568 GMV differences could not be explained by group differences in demographic, personality, 569 psychopathology, or other substance use factors. Examination of THC-naïve 14 year olds who 570 later used cannabis showed no GMV differences, even using a more liberal regions of interest 571 test, suggesting that the differences do not precede cannabis use and are not due to 572 unidentified factors in those predisposed to use. Finally, the spatial distribution of GMV effects 573 was associated with the eCB system, suggesting cannabis exposure may cause these findings.

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574 The preclinical literature presents a number of possible mechanisms by which low levels of 575 cannabis exposure could result in greater GMV relative to THC-naïve controls. Adolescent rats 576 treated with cannabinoid agonist showed altered gliogenesis in regions including the striatum 577 and greater preservation of oligodendroglia relative to control animals (Bortolato et al., 2014). 578 Zebra finches treated with cannabinoid agonist showed greater dendritic spine densities (Gilbert 579 and Soderstrom, 2011); critically, these effects were observed in late-prenatal but not adult 580 animals. Of particular relevance to this study, a single dose of  $\Delta$ 9THC transiently abolished 581 eCB-mediated long-term depression (LTD) in the nucleus accumbens and hippocampus of 582 adolescent mice (Mato et al., 2004). Suspension of LTD may interrupt maturation-related neural 583 pruning and preserve grey matter. Future studies should assess whether these processes 584 operate in human adolescents and whether they produce persisting alterations in GMV.

585 These findings should be interpreted in light of the study's limitations. The IMAGEN sample is 586 racially and ethnically homogenous so it remains to be determined whether the findings 587 generalize to youth from more diverse backgrounds. Substance use was assessed using self-588 report and we do not have standard dose units of cannabis nor information on mode of use or a 589 measure of drug metabolites. Combining images from different sites and imaging platforms 590 remains controversial and is not completely controlled by including site as a covariate. Future 591 studies should replicate the present results using images acquired at the same site on the same 592 scanner or with equal numbers of cases and controls per scanner. We also note that the CNR1 593 gene expression (Hawrylycz et al., 2012) and CB1 receptor density (D'Souza et al., 2016) maps 594 were generated in independent samples of adults and may not accurately represent the eCB 595 system in our sample of adolescents. Although we report significant spatial associations 596 between GMV effects and both CNR1 gene expression and CB1 receptor density, the effect 597 sizes were small and any suggestion that these associations represent mechanisms for the 598 effects we observe is speculative and requires further investigation.

600 We adopted a whole brain, VBM approach to detect effects that were not limited by anatomical 601 boundaries and to allow exploration of spatial relationships between GMV effects and the eCB 602 system. There is evidence, however, that brain perfusion can influence VBM measures of local 603 volume ((Franklin et al., 2013; Franklin et al., 2015; Ge et al., 2017) but cf (Hawkins et al., 604 2018)) so future studies should combine VBM with other measures of brain structure to provide 605 confirmatory evidence. In particular, shape analysis has been shown to be sensitive to brain 606 structural differences associated with cannabis use (Smith et al., 2013; Gilman et al., 2014; 607 Smith et al., 2015; Weiland et al., 2015). Moreover, combining morphometry metrics allows for 608 testing of associations between them, which can identify different relationships between shape 609 deformations and local volume (e.g. (Gilman et al., 2014)) providing evidence of further 610 differences between cannabis users and controls.

611 One source of variability in the human findings on brain structural correlates of cannabis use 612 may be comorbid substance use (Weiland et al., 2015; Gillespie et al., 2018). Given recent 613 evidence of different patterns of functional connectivity in groups using alcohol, nicotine, and 614 cannabis alone and in combination (Vergara et al., 2018), it will be important to account for any 615 possible interaction effects of cannabis with other psychoactive substances. This issue is 616 particularly important considering the ways in which comorbid substance use has been 617 addressed in two recent, widely cited studies. Gilman and colleagues (Gilman et al., 2014) 618 covaried for alcohol and nicotine use and found grey matter density increases and shape 619 deformations associated with cannabis use. Weiland and colleagues (Weiland et al., 2015) 620 matched groups on alcohol and nicotine use and reported no morphometric differences 621 associated with cannabis use, concluding that previously reported differences associated with 622 cannabis may instead be attributable to alcohol use. The participants in Weiland and

623 colleagues' study (Weiland et al., 2015), however, were using alcohol and nicotine at higher 624 levels than those in Gillman and colleagues' study (Gilman et al., 2014). It is possible that 625 cannabis, alcohol and nicotine have differential effects on brain morphometry; specifically, 626 recreational cannabis use has been associated with volume increases, while alcohol has been 627 associated with volume reductions. In the current study, we matched the groups on alcohol and 628 nicotine use and, within the cannabis using group, neither alcohol nor nicotine use was 629 associated with individual differences in GMV, suggesting that the GMV increases we report are 630 associated with cannabis use.

631 We note individual differences in GMV effects: Although regional GMV was increased at the 632 group level for adolescents with low levels of cannabis exposure, the distributions showed a 633 high degree of overlap such that many cannabis users had GMV equivalent to that of controls. 634 None of the tested demographic, personality, or substance use factors stratified GMV in the 635 cannabis users. We note evidence that an association between cannabis use and cortical 636 thickness was stratified by genetic risk for schizophrenia (French et al., 2015) and that an 637 association between cannabis use and hippocampal shape was stratified by dopamine-relevant 638 genes (Batalla et al., 2018). Some adolescents may be vulnerable to GMV effects at extremely 639 low levels of cannabis use and it will be critical to identify those at risk as these structural brain 640 changes may be associated with individual risk for psychopathology and deleterious effects on 641 mood and cognition.

642 Of the behavioral variables tested, only sensation seeking and agoraphobia differed between 643 the cannabis users and controls and these factors were not related to GMV differences. In the 644 cannabis using participants, GMV in the medial temporal clusters was associated with PRIQ 645 and psychomotor speed such that greater GMV in these regions was associated with reduced 646 performance. The finding that right medial temporal GMV predicted generalized anxiety 647 symptoms at follow up for those participants who had used cannabis should be interpreted with 648 caution given the small sample size and that we were not able to identify factors that drove the 649 individual differences in cannabis effects on GMV at baseline. These findings are notable, 650 however, as panic and anxiety symptoms are frequently reported side effects by naïve and 651 occasional cannabis users (Hall and Solowij, 1998). We also note fMRI evidence of 652 hypersensitivity of the amygdala to signals of threat in a partly overlapping sample of cannabis 653 using adolescents (Spechler et al., 2015) and a relationship between adolescent cannabis use 654 and future mood complaints (Wittchen et al., 2007), even with comparatively low levels of use 655 (Cheung et al., 2010).

656 We have revealed GMV increases in adolescents with only one or two instances of cannabis 657 use in regions rich in CB<sub>1</sub> receptors and CNR1 gene expression. Critically, we were able to 658 control for a range of demographic and substance use effects, to confirm that these structural 659 brain effects were not associated with comorbid psychopathology, and to demonstrate these 660 effects were unlikely to precede cannabis use. The pattern of results is characterized by 661 individual differences in GMV effects in the cannabis users; these individual differences were 662 associated with PRIQ and with vulnerability to future symptoms of generalized anxiety. Given 663 the increasing levels of cannabis use amongst adolescents today, we suggest that studying the 664 effects of recreational use early in life is an area of particular importance that should be 665 addressed in the future by large scale, prospective studies.

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835 Figure 1 (a) Those regions showing significantly greater GMV in 14 year olds reporting one or two instances of cannabis use than in matched controls (p<sub>FWE</sub> < 0.05). From left to right, slices 836 are taken from anterior (y=-18) to posterior (y=72) in 15mm increments. The left hemisphere is 837 838 to the right of the image. (b) Outlines of anatomical regions (AAL atlas) superimposed on a 839 binarized mask of the voxels showing significantly greater GMV in 14 year olds reporting one or 840 two instances of cannabis use than in matched controls ( $p_{FWE} < 0.05$ ). For clarity, only those 841 regions for which at least 10% of their volume was included in the significant clusters are 842 represented. From left to right, slices are taken from anterior (y=-18) to posterior (y=72) in 843 15mm increments. The left hemisphere is to the right of the image. c) Outlines of striatal sub-844 regions (Oxford-GSK-Imanova structural striatal atlas (Tziortzi et al., 2011)) superimposed on a 845 binarized mask of the voxels showing significantly greater GMV in 14 year olds reporting one or 846 two instances of cannabis use than in matched controls (pFWE < 0.05). From left to right, slices 847 are taken from inferior (z=-10) to superior (z=8) in 6mm increments. The left hemisphere is to 848 the right of the image. 849

Figure 2: Distribution of Average GMV in the regions showing significantly different GMV
 between those 14 year olds reporting one or two instances of cannabis use and matched
 controls.

Figure 3: Inverse correlations were observed between Perceptual Reasoning Index (PRIQ) and normalized GMV in the left ( $r_{41} = -0.411$ ,  $p_{corr} = 0.037$ ) and right ( $r_{41} = -0.457$ ,  $p_{corr} = 0.012$ ) temporal clusters for those participants reporting one or two instances of cannabis use.

Figure 4. An inverse correlation was observed between normalized GMV in the left temporal cluster and contemporaneous pegboard performance in those participants with low levels of cannabis use ( $r_{39} = -0.454$ ,  $p_{corr} = 0.030$ ).

Figure 5: Associations between normalized GMV in the right temporal cluster at baseline and Generalized Anxiety Disorder DAWBA band scores at follow-up. For those participants with low levels of cannabis use at baseline, those with DAWBA band scores of 0 or 1 at follow up had significantly lower GMV than those who had DAWBA band scores of 3 or 4 (U = 113, p<sub>corr</sub> = 0.030).

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(a)

(b)



p = 5 x 10<sup>-15</sup>

Putamen
Caudate











Table 1: Demographic characteristics of Cohort 1, those 14 year olds reporting 1 or	2
instances of cannabis use (n= 46) and matched controls (n =46).	

Variable	Mean Cannabis	Mean Control	Statistic
Age	14.60	14.51	t <sub>90</sub> = 1.06
PDS	3.04	2.95	t <sub>90</sub> = 0.846
VCI	108.33	108.20	t <sub>90</sub> = 0.042
PRI	102.85	103.77	t <sub>90</sub> = 0.345
SES	18.80	18.72	t <sub>90</sub> = 0.091
Total GMV (mm <sup>3</sup> )	742955.12	728559.92	t <sub>90</sub> = 1.03
Lifetime Alcohol Consumption	3.46	3.52	t <sub>90</sub> = 0.214
Lifetime Nicotine Consumption	2.54	2.59	t <sub>90</sub> = 0.101
Average age of first cannabis	13.83 years		
use			
	Summary	Summary	
	Cannabis	Control	
Sex	65% male	48% male	U = 874
Handedness	87% right handed	87% right handed	<i>U</i> = 1058
Site 1	3	2	<i>U</i> = 1081
Site 2	12	7	<i>U</i> = 1173
Site 3	4	1	<i>U</i> = 1127
Site 4	6	8	<i>U</i> = 1012
Site 5	7	8	<i>U</i> = 1035
Site 6	3	8	<i>U</i> = 943
Site 7	11	12	<i>U</i> = 1035
Site 8	0	0	
# reporting cannabis use in the	10 (21.74%)		
past 30 days (%)			
# reporting cannabis use in the	6 (13.04%)		
past 7 days (%)			

Table 2: Demographic characteristics for those members of cohort 1 for whom specific substance use, psychopathology, and cognitive measures were available at 16 year old follow up.

	Substance Use (n = 31) Mean	Psychopathology (n=33) Mean	Delay Discounting
			(n = 31) Mean
Age	14.60	14.60	14.58
PDS	3.04	3.04	3.03
VCI	110.19	110.31	110.46
PRI	103.91	103.36	103.87
SES	19.01	19.47	19.40
Total GMV (mm <sup>3</sup> )	742793.69	741208.83	742428.43
Lifetime Alcohol	3.61	3.64	3.61
Consumption			
Lifetime Nicotine	2.48	2.45	2.39
Consumption			
	Substance Use	Psychopathology	Delay
	Summary	Summary	Discounting
		-	Summary
Sex	61% male	61% male	61% male
Handedness	90% right	88% right handed	87% right handed
	handed		

Table 3: Demographic characteristics of Cohort 2, those 16 year olds who were abstinent for cannabis use at baseline (age 14) but reported 10 or more instances of cannabis use by age 16 (n = 69) and matched controls (n = 69).

Variable	Mean Cannabis	Mean Control	Statistic
Age	14.43	14.50	t <sub>136</sub> = 0.944
PDS	2.80	2.79	t <sub>136</sub> = 0.290
VCI	112.48	110.29	t <sub>136</sub> = 0.859
PRI	109.16	108.26	t <sub>136</sub> = 0.367
SES	17.97	17.42	t <sub>136</sub> = 0.835
Total GMV (mm <sup>3</sup> )	755082.71	747752.65	t <sub>136</sub> = 0.647
Lifetime Alcohol	2.33	2.29	t <sub>136</sub> = 0.166
Consumption			
Lifetime Nicotine	1.33	1.16	t <sub>136</sub> = 0.577
Consumption			
Average age of first	14.97 years		
cannabis use			
	Summary	Summary	
	Cannabis	Control	
Sex	74% male	70% male	U = 2277
Handedness	93% right handed	91% right handed	<i>U</i> = 2346
Site 1	3	7	U = 2242.5
Site 2	11	9	<i>U</i> = 2449.5
Site 3	4	3	<i>U</i> = 2415
Site 4	8	6	<i>U</i> = 2449.5
Site 5	11	10	<i>U</i> = 2415
Site 6	8	13	<i>U</i> = 2208
Site 7	15	10	U = 2553
Site 8	9	11	<i>U</i> = 2311.5

Table 4: Those regions showing significantly greater GMV in 14 year olds reporting one or two instances of cannabis use than in matched controls.

Anatomical Region (AL)         # significant voxels         % anatomical region implicated           Offactory Cortex         136         20.57           Gyrus Rectus         43         20.57           Gyrus Rectus         43         20.57           Superior Frontal Gyrus (Pars Orbitalis)         23         1.34           Infenor Temporal Gyrus         420         7.87           Middle Temporal Gyrus         164         1.38           Superior Temporal Gyrus         3         0.45           Superior Temporal Gyrus         3         0.45           Superior Temporal Gyrus         5         0.07           Middle Temporal Gyrus         5         0.07           Suboritical         3         3.63           Anygdala         362         74.17           Hippocampus         777         35.63           Putamen         503         21.13           Palldum         126         18.13           Insula         640         14.79           ParaHippocampal Gyrus         202         4.08           Cuadate         92         4.08           Temporal Lobe         77         5.63           Temporal Cyrus         50         0.07	Cluster 1: Left Temporal (Vol. 4968 vox (1	6.767 μl): F <sub>1 80</sub> = 8.88. p <sub>corr</sub> =	0.008: peak voxel -55214)
Frontal Lobe         Image: Construct of the second se	Anatomical Region (AAL)	# significant voxels	% anatomical region implicated
Offactory Cortex         136         20.57           Gyrus Rectus         48         2.33           Superior Frontal Gyrus (Pars Orbitalis)         29         1.34           Inferior Frontal Gyrus (Pars Orbitalis)         39         0.95           Temporal Lobe         787           Middle Temporal Gyrus         164         1.38           Heschi's Cyrus         3         0.65           Superior Temporal Orle         13         0.43           Rolandic Operculum         5         0.07           Subcortical         747         35.63           Putamen         503         21.13           ParaHippocampus         777         35.63           Putamen         603         21.13           ParaHippocampal Gyrus         202         4.85           Caudate         92         4.08           Caudate         92         4.08           Caudate         92         4.08           Caudate         92         4.08           Superior Temporal Cyrus         66         11.62           Superior Temporal Cyrus         68         11.62           Superior Temporal Cyrus         68         11.62           Superior Temporal C	Frontal Lobe		
Gyrus Rectus         48         2.33           Superior Frontal Gyrus (Pars Orbitalis)         29         1.34           Inferior Frontal Gyrus (Pars Orbitalis)         39         0.95           Superior Temporal Gyrus         164         1.38           Hiedle Temporal Gyrus         164         1.38           Busch's Gyrus         3         0.65           Superior Temporal Pole         13         0.43           Subcortical	Olfactory Cortex	136	20.57
Superior Frontal Gyrus (Pars Orbitalis)         29         1.34           Inferior Frontal Gyrus (Pars Orbitalis)         39         0.955           Temporal Gyrus         420         7.87           Middle Temporal Gyrus         164         1.38           Heschi's Gyrus         164         1.33           Reischi's Gyrus         5         0.07           Rolandic Operculum         5         0.07           Inferior Temporal Gyrus         5         0.07           Subcortical         -         -           Arrygdala         382         7.41.7           Hippocampus         777         35.63           Putamen         503         21.13           Paidudum         126         18.13           Insula         640         14.79           ParaHippocampal Gyrus         202         8.56           Caudate         92         4.08           Superior Temporal Lobe         -         -           Temporal Lobe         -         -           Heschi's Gyrus         68         11.62           Superior Temporal Gyrus         610         0.67           Superior Temporal Cyrus         68         11.62           Su	Gyrus Rectus	48	2.33
Inferior Frontal Cyrus (Pars Orbitalis)         39         .0.95           Temporal Cyrus         420         .787           Niddle Temporal Cyrus         164         .138           Heschi's Cyrus         164         .138           Superior Temporal Cyrus         164         .138           Solardic Operation         5         .0.21           Inferior Temporal Cyrus         5         .0.21           Inferior Temporal Cyrus         5         .0.21           Arnygdala         382         .74.17           Hippocampus         .777         .35.53           Putamen         .503         .21.13           Insula         .640         .14.13           Paral-Hippocampal Gyrus         .202         .8.56           Caudate         .92         .4.08           Cluster 2: Right Temporal (Vol. 3710 vox (12,491 µl); Fias = 5.88, pcsr = 0.018; peak voxel 30, -11, -27)         % anatomical region implicated           Temporal Cyrus         .50         .0.67	Superior Frontal Gyrus (Pars Orbitalis)	29	1.34
Temporal Lobe         Hole           Superior Temporal Gyrus         420         7.87           Middle Temporal Gyrus         164         1.38           Resch's Gyrus         3         0.65           Superior Temporal Pole         13         0.43           Rolandic Operculum         5         0.21           Inferior Temporal Gyrus         5         0.07           Subcortical	Inferior Frontal Gyrus (Pars Orbitalis)	39	0.95
Superior Temporal Gyrus         164         .1.38           Hesch's Gyrus         3         .0.55           Superior Temporal Pole         13         .0.43           Rolandic Operculum         6         .0.21           Inferior Temporal Gyrus         .0.5         .0.07           Subcotrical         .0.17         .0.3563           Amygdala         .0.82         .77.7           Anggdala         .0.640         .14.17           Planem         .503         .21.13           Insula         .640         .14.17           ParaHippocampal Gyrus         .202         .8.66           Cluster 2: Right Temporal (Vol. 3710 vox (12.491 µl); F <sub>1.80</sub> = 5.88, Poor         = 0.018; peak voxel 30, -11, -27)           Anatomical Region (AL)         # significant voxels         % anatomical region implicated           Temporal Cyrus         .68         .11.62           Superior Temporal Cyrus         .68         .11.62           Superior Temporal Cyrus         .68         .164           Superior Temporal Cyrus         .746         .3.31           Hippocampus         .746         .3.31           Hippocampus         .746         .3.43           Hippocampla Gyrus (L)         .22.0	Temporal Lobe		
Middle Temporal Gyrus       164       1.38         Resch's Gyrus       3       0.43         Rolandic Operculum       5       0.21         Inferior Temporal Pole       3       0.43         Anygdala       382       0.47         Anygdala       382       74.17         Hippocampus       777       38.63         Putamen       603       21.13         Palidum       126       18.13         Palidum       126       18.13         ParaHippocampal Gyrus       202       8.66         Caudate       92       4.08         Cluster 2: Right Temporal (Vol. 3710 vox (12,491 µl); F <sub>1,80</sub> = 5.88, P <sub>corr</sub> = 0.018; peak voxel 30, -11, -27)       Anatomical region implicated         Matomical Region (AAL)       # significant voxels       % anatomical region implicated         Temporal Lobe       17       0.54         Superior Temporal Gyrus       50       0.67         Superior Temporal Gyrus       64       0.22.0         Palidum       172       264         Putamen       654       22.20         ParaHippocampus       746       3.313         Paldum       172       264         Putamen       50 <t< td=""><td>Superior Temporal Gyrus</td><td>420</td><td>7.87</td></t<>	Superior Temporal Gyrus	420	7.87
Hesch's Gyrus       3       0.55         Superior Temporal Pole       13       0.43         Rolandic Operculum       5       0.07         Subcortical       0       0         Amygdala       382       74.17         Hippocampus       777       3563         Putamen       503       21.13         Insula       640       14.79         ParaHippocampus       202       8.56         Caudate       92       4.08         Cluster 2: Right Temporal Lobe       % anatomical region implicated         Temporal Dobe       %       % anatomical region implicated         Hesch'is Gyrus       50       0.67         Superior Temporal Cycle       17       0.54         Superior Temporal Cycle       17       0.54         Armygdala       439       73.91         Hippocampus       746       33.13         Hippocampus       746       33.19         Hippocampus       746       33.19         Putamen       564       22.20         ParaHippocampal Gyrus       410       15.61         Insula       185       4.39         Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µ); F <sub>140</sub>	Middle Temporal Gyrus	164	1.38
Superior Temporal Pole         13         0.43           Inferior Temporal Cyrus         5         0.07           Subcortical         -         -           Amygdala         382         74.17           Hippocampus         777         35.63           Putamen         603         21.13           Pallidum         126         18.13           ParaHippocampus         202         8.56           Caudate         92         4.08           Cluster 2: Right Temporal (Vol. 3710 vox (12,491 µl); F <sub>1.80</sub> = 5.88, p <sub>cor</sub> = 0.018; peak voxel 30, -11, -27)         Anatomical Region (AL)           Anatomical Region (AL)         # significant voxels         % anatomical region implicated           Temporal Lobe         16         3.13           Hippocampus         50         0.67           Superior Temporal Gyrus         50         0.67           Superior Temporal Gyrus         746         3.313           Palldum         172         26.64           Putamen         564         22.20           Arrygdala         439         73.91           ParaHippocampal Gyrus         410         16.61           Insula         185         4.39           Anatomical Region (AL)	Heschl's Gyrus	3	0.55
Rolandic Operculum         6         0.21           Inferior Temporal Gyrus         5         0.07           Subcortical	Superior Temporal Pole	13	0.43
Inferior Temporal Gyrus         5         0.07           Subcortical	Rolandic Operculum	5	0.21
Subcortical         74.17           Hippocampus         777         35.63           Putamen         503         21.13           Palidum         126         18.13           Insula         640         14.79           ParaHippocampal Gyrus         202         8.56           Caudate         92         4.08           Cluster 2: Right Temporal (Vol. 3710 vox (12,491 µl); F <sub>1.80</sub> = 5.88, p <sub>cor</sub> = 0.018; peak voxel 30, -11, -27)         Anatomical Region (AAL)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         1         16.25           Hesch's Gyrus         50         0.67           Superior Temporal Pole         17         0.54           Mugdala         439         73.91           Hippocampus         746         33.13           Palidum         172         26.46           Putamen         564         22.20           ParaHippocampal Gyrus         410         16.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1.80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 31         Anatomical region implicated           Temporal Lobe         1	Inferior Temporal Gyrus	5	0.07
Amygdala         382         /4.1/           Hippocampus         777         35.63           Putamen         503         21.13           Palldum         126         18.13           Insula         640         14.79           ParaHippocampal Gyrus         202         8.56           Caudate         92         4.08           Cluster 2: Right Temporal (Vol. 3710 vox (12.491 µl); F <sub>180</sub> = 5.88, Pterr = 0.018; peak voxel 30, -11, -27)         Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         1         0.04         9         0.07           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated         1.62           Superior Temporal Cyrus         50         0.07         3.91           Hippocampus         7.46         33.13         9           Patamen         564         22.20         2.84           Putamen         564         22.20         2.84           Temporal Lobe         1         1.62         4.39           Testiform Gyrus (L)         410         1.561         1.91           Insula         185         4.39         5.16           Custer 3: Bilateral P	Subcortical		
Hippocampus         777         35.63           Putamen         503         21.13           Pallidum         126         18.13           Insula         640         14.79           ParaHippocampal Gyrus         202         8.56           Caudate         92         4.08           Cluster 2: Right Temporal (Vol. 3710 vox (12,491 μl); F <sub>180</sub> = 5.88, Ptor = 0.018; peak voxel 30, -11, -27)         Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         # significant voxels         % anatomical region implicated         162           Matomical Region (AAL)         # significant voxels         % anatomical region implicated         162           Superior Temporal Cyrus         50         0.07         0.97           Superior Temporal Pole         17         0.54           Subcortical	Amygdala	382	74.17
Putamen         503         21.13           Insula         640         14.79           ParaHippocampal Gyrus         202         8.56           Caudate         92         4.08           Cluster 2: Right Temporal (Vol. 3710 vox (12.491 µl); F <sub>1.80</sub> = 5.88, pcar         = 0.018; peak voxel 30, -11, -27)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         % anatomical region implicated         %           Superior Temporal Gyrus         50         0.67           Superior Temporal Pole         17         0.54           Superior Temporal Pole         17         0.54           Manygdala         439         73.91           ParaHippocampus         746         33.13           Palidum         172         26.46           Putamen         564         22.20           Paratippocampal Gyrus         410         16.51           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1.80</sub> = 14.32, pcar, = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)         %           Fusiform Gyrus (R)         114         1.91           Parietal Lobe         9         7.98           Parietal Lobe <td>Hippocampus</td> <td>777</td> <td>35.63</td>	Hippocampus	777	35.63
Pailidum         126         18.13           Insula         640         14.79           ParaHippocampal Gyrus         202         8.56           Caudate         92         4.08           Cluster 2: Right Temporal (Vol. 3710 vox (12,491 µl); F <sub>1.80</sub> = 5.88, pcorr         0.018; peak voxel 30, -11, -27)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         %         0.067           Superior Temporal Pole         17         0.54           Subcortical          0.54           Arnygdala         439         73.91           Hippocampus         746         33.13           Pallidum         172         26.46           Insula         185         4.39           Patarhippocampus         746         33.13           Pallidum         172         26.46           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F.1,80 = 14.32, pcorr         8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         -         -         -           Fusiform Gyrus (L)<	Putamen	503	21.13
Insula         640         14.79           ParaHippocampal Gyrus         202         8.56           Caudate         92         4.08           Cluster 2: Right Temporal (Vol. 3710 vox (12,491 µl); F <sub>1.80</sub> = 5.88, p <sub>corr</sub> = 0.018; peak voxel 30, -11, -27)         Matomical Region (AAL)         % anatomical region implicated           Temporal Lobe         % anatomical region implicated         % anatomical region implicated           Heschl's Gyrus         68         11.62           Superior Temporal Opte         17         0.54           Superior Temporal Pole         17         0.54           Migpdala         439         73.91           Hippocampus         746         33.13           Palidum         172         26.46           Putamen         564         22.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4559 vox (16,737 µl); F <sub>1.80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe	Pallidum	126	18.13
Paratrippocampal Gyrus         202         8.56           Caudate         92         4.08           Cluster 2: Right Temporal (Vol. 3710 vox (12,491 µl); F <sub>180</sub> = 5.88, p <sub>corr</sub> = 0.018; peak voxel 30, -11, -27)         Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         % anatomical region implicated         % anatomical region implicated         % anatomical region implicated           Heschl's Gyrus         68         11.62         %           Superior Temporal Pole         17         0.54           Supcorr Temporal Pole         17         0.54           Maygdala         439         73.91           Hippocampus         746         33.13           Palidum         17.2         264.66           Putamen         564         22.20           Paratippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         9         7.98         9         7.98           Posterior Cingulate (R)         59         7.98 <td>Insula</td> <td>640</td> <td>14.79</td>	Insula	640	14.79
Caudate         92         4.08           Cluster 2: Right Temporal (Vol. 3710 vox (12,491 µl); F <sub>180</sub> = 5.88, p <sub>corr</sub> = 0.018; peak voxel 30, -11, -27)         Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         % anatomical region implicated         % anatomical region implicated           Heschl's Gyrus         68         11.62           Superior Temporal Pole         17         0.54           Superior Temporal Pole         17         0.54           Amygdala         439         73.91           Hippocampus         746         33.13           Palidum         172         264.6           Putamen         564         22.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)         Anatomical region implicated           Temporal Lobe         7         0.84         518           Fusiform Gyrus (L)         283         5.18           Fusiform Gyrus (L)         283         5.18           Precureus (R)         20         1.97           Partilipocamplal Gyrus (L)         212         2.55 </td <td>ParaHippocampal Gyrus</td> <td>202</td> <td>8.56</td>	ParaHippocampal Gyrus	202	8.56
Cluster 2: Right Temporal (Vol. 3710 vox (12,491 µl); F <sub>180</sub> = 5.88, p <sub>corr</sub> = 0.018; peak voxel 30, -11, -27)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe           Meschi's Gyrus         68         11.62           Superior Temporal Oyrus         50         0.67           Superior Temporal Pole         17         0.54           Subcortical	Caudate	92	4.08
Cluster 2: Kight Temporal (vol. 3/10 vol. 17, 49 T μj; T <sub>160</sub> = 3.68, p <sub>corr</sub> = 0.016; peak vokef 30, -11, -27)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Heschl's Gyrus         68         11.62           Superior Temporal Pole         17         0.54           Mygdala         439         73.91           Hippocampus         746         33.13           Pallidum         172         266.46           Putamen         564         22.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>180</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)         Anatomical Region implicated           Temporal Lobe         59         7.98           Fusiform Gyrus (L)         283         5.18           Fusiform Gyrus (R)         114         1.91           Parietal Lobe         212         2.55           Occipital Lobe         212         2.55           Cerebellar Chy Carebellum         268         3.45<	Chuster 2: Dight Temporal (Mal. 2740 year)		- 0.049; mask wavel 20, 44, 27)
Antachinical Region (AAL)         # significant voxels         75 anatomical region implicated           Temporal Lobe         68         11.62           Superior Temporal Gyrus         50         0.67           Superior Temporal Pole         17         0.54           Amygdala         439         73.91           Hippocampus         746         33.13           Pallidum         172         26.46           Putamen         564         222.00           ParaHippocampal Gyrus         140         1561         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F1.80 = 14.32, pcorr = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)         Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         7         283         5.18         5.18           Fusiform Gyrus (L)         283         5.18         5.18           Posterior Cingulate (L)         22         1.97         7.98           Posterior Cingulate (L)         212         2.55         2.55           Occipital Lobe         79         1.137         1.62           Lingual Gyrus (R)         1158         21.14         1.61           Lingual Gyrus (R)         158         1.64	Cluster 2: Right Temporal (Vol. 3/10 Vox )	$(12,491 \mu I); F_{1,80} = 5.88, p_{corr} =$	= 0.018; peak voxel 30, -11, -27)
Interpolation         Interpolation           Busch's Gyrus         68         11.62           Superior Temporal Gyrus         50         0.67           Subcortical         0.54         0.67           Amygdala         439         73.91           Hippocampus         746         33.13           Pallidum         172         266.46           Putamen         564         22.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1,80</sub> = 14.32, porr = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)         Anatomical Region (AAL)           # significant voxels         % anatomical region implicated         165           Temporal Lobe         9         7.98           Fusiform Gyrus (L)         283         5.18           Fusiform Gyrus (R)         114         1.91           Precuneus (R)         21         2.55           Occipital Lobe         22         1.97           Precuneus (R)         268         3.45           Precuneus (R)         269         5.16           Calcarine (L)         269         5.16           Calcarine (L)         269		# significant voxels	% anatomical region implicated
Install         06         11.62           Superior Temporal Pole         17         0.54           Subcortical         0         0.67           Amygdala         439         73.91           Hippocampus         746         33.13           Pallidum         172         26.46           Putamen         564         22.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)         Anatomical Region (AAL)           Fusiform Gyrus (L)         283         5.18           Fusiform Gyrus (R)         114         1.91           Parietal Lobe			14.00
Subjenior Temporal Pole         30         0.00           Subcortical         0         0.54           Amygdala         439         73.91           Hippocampus         746         33.13           Pallidum         172         266.46           Putamen         564         22.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)         Anatomical Region (AAL)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe	Superior Temporal Cyrus	60	0.67
Subcortical         17         0.34           Amygdala         439         73.91           Hippocampus         746         33.13           Pallidum         172         266.46           Putamen         564         22.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>-4</sup> ; peak voxel -24, -59, 3)         Anatomical Region (AAL)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe	Superior Temporal Bolo	50	0.07
Amygdala         439         73.91           Hippocampus         746         33.13           Pallidum         172         26.46           Putamen         564         22.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)         Anatomical Region (AAL)           # significant voxels         % anatomical region implicated         % anatomical region implicated           Temporal Lobe         % anatomical region implicated         %           Fusiform Gyrus (L)         283         5.18           Fusiform Gyrus (R)         114         1.91           Parietal Lobe         1         1.97           Precuneus (R)         22         1.97           Precuneus (R)         212         2.55           Occipital Lobe         1158         21.14           Lingual Gyrus (L)         269         5.16           Calcarine (L)         269         5.16           Cerebellar Lobule 4 5         7.9         1.87           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         3		17	0.04
Arrygona         1000           Pallidum         172         26.46           Putamen         564         22.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 μ)); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>-4</sup> ; peak voxel -24, -59, 3)         Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         7         8         5.18         5.18           Fusiform Gyrus (L)         283         5.18         5.18           Fusiform Gyrus (R)         114         1.91           Parietal Lobe         7.98           Posterior Cingulate (R)         59         7.98           Posterior Cingulate (L)         22         1.97           Precuneus (R)         212         2.55           Occipital Lobe         1158         21.14           Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (L)         269         5.16           Calcarine (R)         7.98         1.60           Cerebellar Lobule 4 5 (R)         269         5.16           Calcarine (R)         1158         21.14 <t< td=""><td>Amyodala</td><td>439</td><td>73 01</td></t<>	Amyodala	439	73 01
Πηροσμημα         170         30.16           Pallidum         172         26.46           Putamen         564         22.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 μl); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>-4</sup> ; peak voxel -24, -59, 3)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         % anatomical region implicated         9           Fusiform Gyrus (R)         114         1.91           Parietal Lobe         9         7.98           Posterior Cingulate (L)         22         1.97           Precuneus (R)         212         2.55           Occipital Lobe         1158         21.14           Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (L)         269         5.16           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellum         102         269         5.16           Calcarine (L)         308         14.62 <td>Hippocampus</td> <td>746</td> <td>33.13</td>	Hippocampus	746	33.13
Tutinum         112         20-70           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1.80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)         Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         # significant voxels         % anatomical region implicated         114         1.91           Fusiform Gyrus (L)         283         5.18         5.18         5.18           Fusiform Gyrus (R)         114         1.91         9         1.97           Posterior Cingulate (R)         59         7.98         7.98           Posterior Cingulate (L)         222         1.97           Precuneus (R)         268         3.45           Precuneus (R)         268         3.45           Precuneus (L)         212         2.55           Occipital Lobe         0         0           Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (R)         1158         21.14           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellar Lobule 4 5 (	Pallidum	172	26.46
Totken         301         121.20           ParaHippocampal Gyrus         410         15.61           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>-4</sup> ; peak voxel -24, -59, 3)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe           Fusiform Gyrus (L)         283         5.18           Fusiform Gyrus (R)         114         1.91           Parietal Lobe          1.97           Posterior Cingulate (R)         59         7.98           Posterior Cingulate (L)         222         1.97           Precuneus (R)         268         3.45           Precuneus (L)         212         2.55           Occipital Lobe          2           Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (R)         158         21.14           Lingual Gyrus (R)         269         5.16           Calcarine (L)         269         5.16           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellar Lobule 4 5 (R)	Putamen	564	22.40
Insula         110         100           Insula         185         4.39           Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>4</sup> ; peak voxel -24, -59, 3)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         % anatomical region implicated           Fusiform Gyrus (L)         283         5.18           Fusiform Gyrus (R)         114         1.91           Parietal Lobe          7.98           Posterior Cingulate (R)         59         7.98           Posterior Cingulate (L)         22         1.97           Precuneus (R)         268         3.455           Precuneus (R)         212         2.55           Occipital Lobe             Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (L)         818         16.01           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20	ParaHippocampal Gyrus	410	15.61
Indust         Indus         Indus         Indus <td>Insula</td> <td>185</td> <td>4 39</td>	Insula	185	4 39
Cluster 3: Bilateral Posterior (Vol. 4959 vox (16,737 µl); F <sub>1,80</sub> = 14.32, p <sub>corr</sub> = 8.0 x 10 <sup>-4</sup> ; peak voxel -24, -59, 3)           Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         % anatomical region implicated           Fusiform Gyrus (L)         283         5.18           Fusiform Gyrus (R)         1114         1.91           Parietal Lobe         7.98           Posterior Cingulate (R)         59         7.98           Posterior Cingulate (L)         22         1.97           Precuneus (R)         268         3.45           Precuneus (R)         268         3.45           Precuneus (L)         212         2.55           Occipital Lobe         2         2.114           Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (L)         818         0.601           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellum         308         14.62           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20		100	1.00
Anatomical Region (AAL)         # significant voxels         % anatomical region implicated           Temporal Lobe         *         % anatomical region implicated           Fusiform Gyrus (L)         283         5.18           Fusiform Gyrus (R)         114         1.91           Parietal Lobe         *         *           Posterior Cingulate (R)         59         7.98           Posterior Cingulate (L)         22         1.97           Precuneus (R)         212         2.55           Occipital Lobe         *         *           Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (L)         818         16.01           Calcarine (L)         269         5.16           Cerebellum         79         1.87           Cerebellar Lobule 4 5 (R)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 6 (R)         7         0.88      <	Cluster 3: Bilateral Posterior (Vol. 4959 vo	ох (16.737 цl): F <sub>1 80</sub> = 14.32, р	orr = 8.0 x 10 <sup>-4</sup> ; peak voxel -24, -59, 3)
Temporal Lobe         10         11         10         11         10         11	Anatomical Region (AAL)	# significant voxels	% anatomical region implicated
Fusiform Gyrus (L)         283         5.18           Fusiform Gyrus (R)         114         1.91           Parietal Lobe         59         7.98           Posterior Cingulate (R)         22         1.97           Precuneus (R)         268         3.45           Precuneus (R)         212         2.55           Occipital Lobe         1158         21.14           Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (R)         1158         21.14           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellum         79         1.87           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Lobule 4 5 (L)         365         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8	Temporal Lobe		
Fusiform Gyrus (R)         114         1.91           Parietal Lobe         7.98           Posterior Cingulate (R)         59         7.98           Posterior Cingulate (L)         22         1.97           Precuneus (R)         268         3.45           Precuneus (L)         212         2.55           Occipital Lobe         1         1           Lingual Gyrus (L)         1158         21.14           Lingual Gyrus (L)         818         16.01           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Lobule 4 5 (L)         368         0.13	Fusiform Gyrus (L)	283	5.18
Parietal Lobe         Posterior Cingulate (R)         59         7.98           Posterior Cingulate (L)         22         1.97           Precuneus (R)         268         3.45           Precuneus (L)         212         2.55           Occipital Lobe         2         1.97           Lingual Gyrus (L)         1158         21.14           Lingual Gyrus (L)         818         16.01           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Lobule 4 5 (L)         368         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum (L)         8         0.13	Fusiform Gyrus (R)	114	1.91
Posterior Cingulate (R)         59         7.98           Posterior Cingulate (L)         22         1.97           Precuneus (R)         268         3.45           Precuneus (L)         212         2.55           Occipital Lobe           Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (L)         818         16.01           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Lobule 6 (R)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Parietal Lobe		
Posterior Cingulate (L)         22         1.97           Precuneus (R)         268         3.45           Precuneus (L)         212         2.55           Occipital Lobe         2         1           Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (L)         818         16.01           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellum         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (L)         265         6.18           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Posterior Cingulate (R)	59	7.98
Precuneus (R)         268         3.45           Precuneus (L)         212         2.55           Occipital Lobe	Posterior Cingulate (L)	22	1.97
Precuneus (L)         212         2.55           Occipital Lobe	Precuneus (R)	268	3.45
Occipital Lobe            Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (L)         818         16.01           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum (L)         8         0.13	Precuneus (L)	212	2.55
Occipital Lobe            Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (L)         818         16.01           Calcarine (L)         269         5.16           Calcarine (R)         79         1.37           Cerebellum           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13			
Lingual Gyrus (R)         1158         21.14           Lingual Gyrus (L)         818         16.01           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellum           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Occipital Lobe		
Lingual Gyrus (L)         818         16.01           Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellum         79         1.87           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Lingual Gyrus (R)	1158	21.14
Calcarine (L)         269         5.16           Calcarine (R)         79         1.87           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Vermis (4 5)         258         14.62           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Lingual Gyrus (L)	818	16.01
Calcarine (R)         79         1.87           Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Calcarine (L)	269	5.16
Cerebellum         Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Calcarine (R)	79	1.87
Cerebellar Vermis (4 5)         258         17.61           Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Cerebellum		
Cerebellar Lobule 4 5 (R)         308         14.62           Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Cerebellar Vermis (4 5)	258	17.61
Cerebellar Lobule 6 (L)         332         8.20           Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Cerebellar Lobule 4 5 (R)	308	14.62
Cerebellar Lobule 6 (R)         265         6.18           Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Cerebellar Lobule 6 (L)	332	8.20
Cerebellar Lobule 4 5 (L)         156         5.80           Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Cerebellar Lobule 6 (R)	265	6.18
Cerebellar Vermis (6)         7         0.88           Crus Cerebellum1 (L)         8         0.13	Cerebellar Lobule 4 5 (L)	156	5.80
Crus Cerebellum1 (L) 8 0.13	Cerebellar Vermis (6)	7	0.88
	Crus Cerebellum1 (L)	8	0.13

Table 5: The number of Ventral Striatum voxels (and % of total anatomical volume) showing significantly greater GMV in 14 year olds reporting one or two instances of cannabis use than in matched controls.

	# significant voxels	% anatomical region implicated
Ventral Striatum (Left)	131	30.32
Ventral Striatum (Right)	226	54.72

Table 6: Those regions showing significantly greater GMV in 14 year olds reporting one or two instances of cannabis use than in matched controls, when controlling for agoraphobia and sensation seeking.

Region	Vol.	Peak Voxel	
Left Temporal	4836 vox	-55, -2, -14	$F_{1,76} = 8.018, p_{corr} = 0.011$
	(16,321µl)		
Right Temporal	3425 vox	30, -11, -27	$F_{1,76} = 6.026, p_{corr} = 0.016$
	(11,559µl)		
Bilateral Posterior/ Inferior Parietal	4907 vox	-24, -59, 3	F <sub>1,76</sub> = 12.718, p <sub>corr</sub> = 0.002
	(16,561µl)		
Left Inferior Temporal Gyrus	603 vox	-50, -9, -42	F <sub>1,76</sub> = 12.755, p <sub>corr</sub> = 0.002
	(2,038µl)		

Table 7: No significant GMV differences were observed at baseline between those participants who were abstinent for cannabis use at age 14 but reported at least 10 instances of use by age 16 and matched controls (i.e. Cohort 2) in those regions defined in Cohort 1.

Region	Vol.	Peak Voxel	
Left Temporal	4968 vox	-55, -2, -14	F <sub>1,125</sub> = 3.026, p <sub>corr</sub> = 0.252
	(16,767 μl)		
Right Temporal	3710 vox	30, -11, -27	$F_{1,125} = 5.626, p_{corr} = 0.057$
	(12,491 μl)		
Bilateral Posterior/ Inferior Parietal	4959 vox	-24, -59, 3	F <sub>1,125</sub> = 0.021, p <sub>corr</sub> ≈ 1
	(16,737 μl)		