

Volumetric assessment and longitudinal changes of brain structures in formalinized Beagle brains

Beagle formalinized brains: volumetric and longitudinal changes

Francesca Del Signore¹, Germain Arribarat², Leonardo Della Salda^{1*}, Giovanni Mogenicato², Alexandra Deviers², Benjamin Cartiaux², Massimo Vignoli¹, Patrice Peran^{2}, Francesco de Pasquale^{1**}**

¹ **Veterinary Faculty, University of Teramo, Teramo, Italy**

²**ToNIC Toulouse Neuroimaging Center UMR1214 - Inserm/UPS, Toulouse, France**

*** Corresponding author**

Email: ldellasalda@unite.it (LDS)

**** The authors contributed equally**

25 **Abstract**

26 High field MRI represents an advanced technique both for diagnostic and research purposes on animal
27 models such as the Beagle dog. The increasing interest in non-invasive neuroscience, aging, and
28 neuropathological research led to a need of reference values (in terms of volumetric assessment) for
29 the typical brain structures involved and, nowadays, several canine brain MRI atlases have been
30 provided. Since no reports are available regarding the measurements reproducibility and few
31 information are available about formalin fixation effect on brain structures when applied to MRI
32 segmentation, we assessed the segmentation variability of selected structures as a function of the
33 operator (two operators segmented the same data) and their intrinsic variability within a sample of 11
34 Beagle dogs (9 females and 2 males, 1.6 ± 0.2 years). Then, we analyzed for one further Beagle dog
35 (2 years old) the longitudinal changes in the brain segmentations of these structures corresponding
36 four conditions: *in vivo*, *post mortem* (after euthanasia), *ex vivo* (brain extracted and studied after 1
37 month in formalin and after 11 months); all the MRI images were collected with a 3 T MRI scanner.
38 Our findings suggest that the segmentation procedure can be considered overall reproducible since
39 only slight statistical differences were detected, apart from the ventricles.

40 Furthermore, in the *post mortem/ ex vivo* comparison, the majority of the structures showed a higher
41 contrast leading to more reproducible segmentations across operators and a net increase of volume of
42 the studied structures; this could be justified by the intrinsic relaxation time changes observed as a
43 consequence of formalin fixation, that led to an improvement of brain structures visualization and
44 then segmentation.

45 To conclude, MRI based segmentation seems to be a useful and accurate tool that allows longitudinal
46 studies, especially when applied to formalin fixed brains.

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50 **Introduction**

51

52 High field MRI, thanks to its high SNR and short acquisition time, represents an advanced technique
53 both for diagnostic and research purposes on animal models such as the dog [1,2]. This model offers
54 several advantages over more standard rodent and primate ones as testified by the growing literature
55 on neurocognitive, aging, and clinical applications [3]. Neurocognitively, the canine shares similar
56 behavioral/emotional responses with humans, e.g. in linking learning, memory, and other cognitive
57 functions. These convergent sociocognitive skills places the dog in a unique position to increase our
58 understanding of sociocognitive in humans [4]. Being gyrencephalic, the dog brain, as compared to
59 rodent and avian, it represents a better experimental model for several disorders, e.g. gliomas and
60 aging [5–7]. Among domestic canines, the Beagle is the breed most commonly used in laboratories,
61 thanks to its moderate size, docile nature, and strong immunity [8–10]. Such increasing interest in
62 non-invasive neuroscience, aging, and neuropathological research led to a need of reference values
63 (in terms of volumetric assessment) for the typical brain structures involved. To this aim, recent
64 studies developed a standard canine brain atlas to provide a common spatial referencing and
65 architectonic-based cortical segmentation of coregistered data [4]. Similarly, a stereotactic cortical
66 atlas for the mesaticephalic canine brain has been recently developed for functional and structural
67 MRI analyses [3]. In general, the currently available atlases, see for example [11–13], are affected by
68 some limitations such as the small number of subjects used [11], the acquisition of non-isovolumetric
69 data [13], the use of dogs non neurologically/clinically healthy [12] and the mixing of different breeds
70 included in the sample [3]. To overcome the uncertainty related to the breed variability, very recently
71 Liu et al. realized a specific atlas for the Beagle breed, which is the most used breed in this kind of
72 studies [14]. However, all these studies were performed on alive subjects, apart from the work of
73 Datta and colleagues, 2012, where formalin-fixed brains (*ex vivo*) were segmented to obtain a
74 diffeomorphic brain atlas of mesaticephalic dogs coregistered onto an *in vivo* template [11].

75 Nevertheless, in this study, some important aspects such as the reproducibility of the measurements
76 across different operators and the volumetric variation from *in vivo* to *ex vivo* phases were not
77 assessed [11]. Such a relationship is very important since MRI findings can be linked and often
78 validated through histopathology. This can be very time-consuming and challenging for several
79 reasons, e.g. the inaccurate correspondence MRI-anatomical sections (due to different slice thickness
80 and orientations) [15]. For this reason, when whole-brain histopathology is not feasible, *in vivo* and
81 *post mortem* MRI can be used as a guide for limited pathological sampling [16–18]. Similarly, in
82 forensic radiology, *post mortem* MRI has been recognized as a supplementary tool to address specific
83 forensic questions [19- 21]. However, *ex-vivo* MRI can be very challenging. First, after death, the
84 brain undergoes several changes, e.g. microbial degradation, autolysis, breakdown of cell
85 membranes, and stochastic diffusion of molecules. Second, also the chemical fixation, needed to
86 ensure the longitudinal stability of the macromolecular structure, might affect the tissue properties.
87 Therefore, due to death and fixation, a series of artifacts and changes of tissue properties are expected.
88 This will affect MR imaging and the conclusions based on MRI measurements in fixed tissue may
89 not reflect directly the *in-vivo* environment [21]. For example, it has previously been reported that
90 formalin fixation causes a tissue shrinkage that not be homogeneous among the various brain
91 structures, and it might vary over time [22]. However, these aspects are still under debate and the
92 literature is scares. For this reason, in this study, we analyzed the effect of long-term fixation (12
93 months) on brain structures in a sample of 11 Beagle dogs. We manually segmented a set of brain
94 structures, e.g. Globus Pallidus, Caudate, and Substantia Nigra, whose volumetric changes have been
95 reported to correlate with many neurodegenerative disorders such as Parkinson's [23]. First, we
96 assessed the variability of the extracted volumes as a function of the operator (two operators
97 segmented the same data) and their intrinsic variability within the sample. Then, we analyzed for one
98 further dog the longitudinal changes in the brain segmentations of these structures corresponding four
99 conditions: *in vivo*, *post mortem* (after euthanasia), *ex vivo* (brain extracted and studied after 1 month

100 in formalin and after 11 months). In this way, the last condition overlaps with the previous sample of
101 11 dogs.

102 The estimated volumetric data can represent an important reference for future studies and the
103 longitudinal effects observed might shed light on which structures are segmented more accurately as
104 a function of time spent in formalin. As far as we know, this is the first study reporting brain structures
105 in formalinized dogs and their longitudinal changes.

106

107 **Materials and Methods**

108

109 **Animals**

110 A sample of 12 healthy Beagle dogs were evaluated in two studies. In the first study, a group of 11
111 dogs (9 females and 2 males, 1.6 ± 0.2 years) was used for the evaluation of the effect of long-term
112 fixation on MRI properties of the brain. In this context, a single MRI scan was performed on isolated
113 heads that remained fixed for 11 months. Dogs originated from a laboratory in which they completed
114 their research time and were euthanized for teaching purposes (i.e. preparation of veterinary
115 anatomical teaching materials: MRI brain atlas and embalmed cadavers for dissection).

116 In the second study, one dog (male, 2 years) was used for the longitudinal evaluation of the effect of
117 death and fixation on MRI properties of the brain. This dog underwent 4 MRI exams: 1 *in vivo*, 1
118 *post-mortem* performed just after euthanasia, and 2 *ex vivo* performed on the brain removed from the
119 skull after 1 (*ex_vivo_1*) and 12 months of fixation (*ex_vivo_12*). While the terms *post-mortem* and
120 *ex vivo* are normally interchangeable, in this study, they refer to two different conditions which are
121 evaluation of the brain confined by the skull just after death (*post-mortem*) and evaluation of formalin-
122 fixed brains (*ex vivo*, either isolated or confined by the skull).

123 The experimental procedures related to the preparation of veterinary anatomical teaching materials
124 were approved by the Animal Ethics Committee of the National Veterinary School of Toulouse with
125 authorization n° 21559-2019071917392588. Dogs were euthanized by an IV injection of ≥ 100 mg/kg
126 sodium pentobarbital while they were deeply anesthetized (anesthetic protocol: IV injection of
127 butorphanol (0,4 mg/kg), medetomidine (20 μ g/kg), and diazepam (0,2 mg/kg)). Heparin sodium
128 (1000 IU) was injected by IV route 5 minutes before euthanasia to optimize post-mortem perfusion
129 of fixative solution.

130

131

132 **Fixation protocol**

133 Dogs were anesthetized to acquire in vivo MR images (not used in the present study) after which they
134 were euthanized, and their heads were fixed according to the procedure described above. Heads were
135 then stored in containers filled with 10% formalin solution and were scanned after 11 months of
136 fixation.

137 For the dog in the second study, in vivo MRI scans and euthanasia were carried out under anesthesia
138 with the protocol described above. The post-mortem MRI examination was performed just after
139 euthanasia and once this acquisition was completed, the cadaver was transferred to a special room for
140 fixation. The head was then separated from the body to be perfused via the common carotid arteries
141 with a rinsing solution (NaCl, flow rate: 15 mL/minute, perfusion time: 5 minutes) and a fixative
142 solution (10% formalin solution, 15 mL/minute, perfusion time: 5 minutes). The head was stored in
143 a container filled with 10% formalin solution. After one month of fixation, the brain was removed
144 from the skull for an ex vivo MRI acquisition. An additional ex vivo MRI examination of the brain
145 was performed after 11 months of fixation.

146

147 **MRI acquisition**

148 MRI examinations were performed at the Institute for Brain Sciences of Toulouse using a high field
149 3.0 Tesla magnet (Philips ACHIEVA dStream) at the Inserm/UPS UMR1214 ToNIC
150 Technical Platform, and a, 8-channel, human elbow coil (serving as dog head coil) for signal
151 reception. The Ex-vivo examinations were performed with a 1-channel solenoid antenna. To
152 guarantee a homogeneous and accurate signal, no acceleration or preparation factors were used. For
153 the dog with longitudinal follow-up of the brain, the imaging protocol comprised of T1 and T2
154 weighted images. The 3D whole-brain T1 and T2 weighted images were obtained in the sagittal plane.
155 For T1 imaging (Fast Field Echo), the sequence parameters were the following: echo time/repetition
156 time = 4.0/9.0 ms, flip angle = 8°. For T2 imaging (Spin Echo) the sequence parameters were the

157 following: echo time/repetition time = 266/2500 ms, flip angle = 90°. The spatial resolution
158 parameters were the same for both acquisitions: pixel spacing $0.5 \times 0.5 \text{ mm}^2$, slice thickness
159 = 0.5 mm, matrix size = 288x288, numbers of slices = 300, voxel size = $0.5 \times 0.5 \times 0.5 \text{ mm}^3$. The
160 total duration of the imaging protocol was 60 min.

161 These sequences were used for the 4 MRI examinations of the follow-up. Twenty-four hours before
162 the ex vivo MRI scans, the brain was rinsed with water and then submerged in a 0,9% saline solution
163 (NaCl). Just before acquisition, the brain was put in an MRI-compatible container (a plastic container
164 with a leakproof screw cap) totally filled with saline solution, the container was then placed in the
165 elbow coil.

166 Twenty-four hours before their acquisition, the 11 fixed heads were rinsed with water and then
167 submerged in NaCl. For the acquisition, they were dried, wrapped up in hermetic packages, held
168 horizontally on the MRI table, and placed in the human elbow coil. The imaging protocol comprised
169 T1-weighted images (repetition time TR = 8.5 ms; echo time TE = 3.8 ms; voxel size $0.5 \times 0.5 \times 0.5$
170 mm, matrix 288x288x300) and T2-weighted images (repetition time TR = 265.71 ms; echo time TE
171 = 2500 ms; voxel size $0.5 \times 0.5 \times 0.5 \text{ mm}$, matrix 288x288x300).

172

173 **Assessment of inter-operator reproducibility**

174 Two veterinarians experienced in canine brain structure segmentation operators (O1- and O2
175 respectively) blindly segmented the following structures: Ventricles (from T1 scans) and Caudate
176 Nucleus, Hippocampus, Substantia Nigra, Putamen, Globus Pallidus, Lateral and Medial Geniculate
177 Nucleus (from T2 scans), dividing the measurements in left and right parts on the group of 11 dogs
178 and the longitudinal study. The choice to perform the segmentations either in T1 or T2 was driven by
179 the contrast observed in the various structures.

180 As far as it regards the analysis on the measurements collected on the group of 11 dogs, on each
181 structure mean (M) and standard deviation (SD) was evaluated. Based on them, to assess the intra-

182 operator reproducibility, the percentage volumes were computed as $CV = SD/M$. Finally, the
183 statistical comparison between the two operators was provided by a t-test.

184

185 **Longitudinal study**

186 For the second part of the study, the two operators performed the same segmentations of the previous
187 phase, and the percentage (%) variations of the volumes between the different scanning phases were
188 estimated and compared between the two operators. We considered the following conditions: *in vivo*
189 *vs post mortem* (defined as $(post\ mortem - in\ vivo)/in\ vivo * 100$), the *post mortem vs ex vivo* – 1-
190 month volumes (expressed as $(ex_vivo_1 - post\ mortem)/post\ mortem * 100$), the *in vivo vs ex vivo*
191 1 month volumes (expressed as $(ex_vivo_1 - in\ vivo)/in\ vivo * 100$) and, finally, *ex vivo* 1 month vs
192 *ex vivo* 12 months (expressed as $((ex_vivo_12 - ex_vivo_1)/ex_vivo_1 * 100)$).

193 ITK SNAP software (version 3.8.0, 2019) was used as a tool to collect the segmentation and statistical
194 analysis was performed with MATLAB; the statistical significance was defined as an alpha level of
195 alpha = 0.05.

196

197 **Results**

198

199 **Brain segmentation after 12 months in formalin**

200 The segmentations obtained from two operators after 12 months in formalin are reported in

201 Table 1.

202

203 **Table 1. Summary table for the 11 dogs.**

		O1			O2			
		<i>Mean</i>	<i>ST</i>	<i>CV%</i>	<i>Mean</i>	<i>ST</i>		<i>p value</i>
		<i>(mm³)</i>	<i>DEV</i>		<i>(mm³)</i>	<i>DEV</i>	<i>CV%</i>	
Ventricles	T1	1059.27	240.87	0.22	674.9	197.52	0.25	<0.05
Caudate	T2 L	576.09	56.82	0.09	600.36	71.69	0.11	0.38
Caudate	T2 R	576.77	58.46	0.10	596.72	64.26	0.10	0.31
Hippocampus	T2 L	588.06	54.16	0.09	635.3	62.15	0.09	0.07
Hippocampus	T2 R	588.61	35.57	0.06	620.04	48.87	0.07	0.10
Sub Nigra	T2 L	48	11.04	0.22	50	9.73	0.19	0.64
Sub Nigra	T2 R	54.19	9.73	0.17	57.9	9.90	0.17	0.37
Lat Geniculate	T2 L	63.55	4.78	0.07	71.2	14.83	0.20	0.11
Lat Geniculate	T2 R	67.26	6.92	0.10	74.68	11.98	0.16	0.09
Med Geniculate	T2 L	78.19	12.85	0.16	88.64	18.22	0.20	0.13
Med Geniculate	T2 R	85.50	12.45	0.14	93.24	14.27	0.15	0.19
Putamen	T2 L	76.14	11.88	0.15	77.25	13.46	0.17	0.84
Putamen	T2 R	69.96	9.50	0.13	71.65	13.93	0.19	0.74
Globus Pallidus	T2 L	55.06	9.10	0.16	60.18	11.36	0.18	0.25
Globus Pallidus	T2 R	58.02	8.96	0.15	61.79	11.32	0.18	0.39

204 Estimated volumes of the considered structures. The obtained measurements reported for operator 1
205 and 2 (O1/O2), expressed the volume averaged across the considered subjects, their standard
206 deviation and CV%. Within this sample the only statistically significant difference between the
207 operators is observed for the ventricles (red).

208

209 For each operator (O1/O2), we report the mean and standard deviations of the volumes and
210 their percentage variations expressed as CV%. The ventricles expressed the highest variability within
211 the same operator (22% for O1 and 25% for O2), followed by the Substantia Nigra and Geniculate
212 (around 20%) while the remaining structures showed a variability around 15% or less (see Fig 1).

213

214 **Fig 1. Segmentation stability across operators.**

215 The percentage variability of each structure operator O1 (solid line) and O2 (dotted line). It can be
216 noticed that the lowest variability, except for the ventricles, was observed for both operators for the
217 largest and more defined brain structures, thus suggesting that the MRI volume of the segmented
218 structure is influenced by the actual size and its intrinsic contrast with the surrounding parenchyma.

219

220 The spatial topography of these variations is shown in Fig 2, where the % CV, averaged across
221 operators, is overlaid on a T1w image of a representative dog.

222

223 **Fig 2. The spatial topography of the % CV of the considered structures.** The % CV averaged
224 across operators overlaid on T1w images of a representative dog.

225

226 As it can be seen in Table 1 and Fig 3, where the whisker plots of the analyzed structures are
227 reported, apart from the ventricles ($p < 0.01$ – dotted box in Fig 3), a t-test evidenced no statistically
228 significant differences among the two users.

229

230 **Fig 3. Data distribution of the segmented structures.** Whisker plot for the distribution of the
231 segmented volumes from the two operators O1 (black) and O2 (white).

232 A) The set Ventricles, Caudate, and Hippocampus. Only the ventricles resulted statistically different
233 between the two operators (dotted box).

234 B) The set Substantia Nigra, Lateral and Medial Geniculate, Putamen and Globus Pallidus. In this set
235 no statistical difference is observed.

236

237 This seems to suggest that the segmentation of the considered structures is quite stable and
238 reproducible across operators. The observed difference in the ventricles might be ascribed to the
239 difficulty to identify their exact border with cerebrospinal fluid (CSF). To summarize, these findings
240 suggest that manual segmentation provides globally reproducible measurements. Further, we
241 observed some structure variability across dogs, that cannot be ascribed to operators' skills. Such
242 intrinsic structure variability shows that some structures are more stable than others. Of note, larger
243 structures such as Caudate and Hippocampus showed the lowest values of dispersion, while smaller
244 structures such as Medial and Lateral Geniculate were characterized by higher values of dispersion.
245 This might be justified since both operators experienced higher difficulty to segment the smaller
246 structures due to their relatively poor signal contrast.

247 **A longitudinal study on brain structures**

248 In this part of the study, for a representative dog, we assessed the longitudinal changes of
249 segmentations performed *in vivo*, *post mortem*, and *ex vivo_1* (after one month in formalin) and *ex*
250 *vivo_12* (after 12 months in formalin).

251 First, we observed that both T1w and T2w images showed a change of signal contrast between the
252 *post mortem* and *ex-vivo* data (Fig 4).

253

254 **Fig 4. T1 and T2 progressive change of contrast.** Panels A-D: T_w transverse section of the studied
255 dog in the four different phases of the experiment, respectively *in vivo* (A), *post mortem* (B), *ex vivo*
256 1 month (C), and *ex vivo* 12 months (D). *In vivo* and *post mortem* grey and the white matter appeared
257 respectively hypointense and hyperintense, on *ex vivo* images the contrast appeared exactly the
258 opposite, with grey and white matter respectively hyperintense and hypointense. The solid arrows
259 point at the Substantia Nigra and it can be observed the progressive increase of contrast. Panels E-H
260 represent the same four phases from T_{2w} images. In this case, rather than an inversion of the normal
261 contrast, it has been observed a strong decrease of intensity of the white matter in the *ex vivo* phase,
262 thus leading to an increased definition of the contours of the various structures.

263

264 Specifically, on T_{1w} images, while *in vivo* (A) and *post mortem* (B) grey and white matter
265 (GM and WM) appeared respectively hypointense and hyperintense, on *ex vivo* images (C-D) the
266 contrast seems the opposite, with GM and WM matter respectively hyperintense and hypointense.
267 This is evident for the Substantia Nigra, (white arrow in the figure). It can be observed that while in
268 panels A and B this structure is barely visible, in panels C (*ex vivo_1*) and D (*ex vivo_12*) the contrast
269 increased with borders of the structure more evident. On T_{2w} images, it is observed a strong decrease
270 in the white matter in both *ex vivo* phases, leading to an increased definition of the contours of the
271 various structures. This is most evident for the smallest structures (e.g. Putamen or Globus Pallidus),
272 whose borders were difficult to define by both operators *in vivo* and *post mortem* phases (Fig 4 E-H).

273 The volumetric percentage variations (see Materials and Methods) across time of the considered
274 structures are reported in Fig 5.

275

276 **Fig 5. Longitudinal variations.** A) Volume variation between the *in vivo* and *post mortem* phases.
277 Volume variations were obtained for O1 (solid), O2 (dashed), and the mean trend (gray). It can be
278 observed that the volume variation is always lower than 20%. The trend is variable between the two
279 operators. This aspect may be justified by the difficulty to clearly distinguish the exact borders of the
280 structures in the *vivo* phase.

281 B) Volume variation between the *post mortem* and *ex_vivo_1* (after 1 month) phases. Volume
282 variations were obtained on each structure for O1 (solid), O2 (dashed), and the mean trend (gray). It
283 can be noted that the agreement between the two operators is higher than in the previous phase, with
284 most structures experiencing an increase in volume from the *post mortem* to *ex vivo* phase. This is
285 particularly interesting since the shift of the contrast after the time in formalin qualitatively seems to
286 improve the visualization of the smallest structures.

287 C) Volume variation between the *in vivo* and *ex_vivo_1* phases. It can be noted that also in this case
288 the operators agree and a trend similar to the previous figure is observed across the structures, i.e.
289 most of the smallest structures appeared more clearly identifiable, thus justifying an increase of the
290 volume.

291 D) Volume variation between the *ex_vivo_1* and *ex_vivo_12* phases. It can be easily observed that
292 the volume variation is relatively low between the two phases, with an overall agreement from both
293 operators, thus suggesting that the time spent in formalin does not significantly influence the volume
294 variation.

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299 As far as it regards the across-operator variability of the segmentations, also in this case, they
300 seem reproducible in the various phases, i.e. the % variation of the volumes globally follows a similar
301 trend for the two operators. Specifically, for the *in vivo/ post mortem* comparison (Fig 5A), the
302 volumetric variation observed in the various structures is close to 12%, fluctuating at most around
303 20%, and thus can be considered low. Notably, in some structures such as the Lateral and Medial
304 Geniculate nucleus, the volume increased by 18-20%. In this comparison (*in vivo* vs *post mortem*),
305 especially for the *in vivo* images, both operators experienced some difficulties to delineate the borders
306 of the investigated structures. They reported that this was independent of the structures' dimension,
307 i.e. it applied also to the largest structures. This applies to the *in vivo* phase while in the *post mortem/*
308 *ex vivo* comparison, the majority of the structures showed a higher contrast leading to more
309 reproducible segmentations across operators, the highest observed variability ranging around 10%
310 (Fig 5B-D). However, we also observed for some structures, a large increase in volume. This was not
311 expected since the formalin fixations are reported to lead to shrinking and reduced tissue volumes
312 (24). This effect was particularly evident for Substantia Nigra (average increase of around 70% Fig
313 5B, C), followed by Globus Pallidus (around 30%) and Geniculate (around 20%). In general, for both
314 *ex vivo* phases, all the operators described an increased contrast to detect the borders of the various
315 structures, and the smallest structures were described to be sensitively easier to be segmented.
316 Considering the previous findings, this suggests that the increase of contrast seems to be ascribed to
317 the *ex-vivo* condition and thus the effect of fixation. Now, to study if this change remains stable over
318 time, we compared the two *ex vivo* conditions (*ex_vivo_1* and *ex_vivo_12*). As it can be seen in Fig
319 5D, the highest changes fluctuate around 20%. This suggests that the increased time spent in formalin
320 did not influence significantly the volumes of the various structures. Basically, in this period the
321 structures' volumes remained stable. This is an important finding suggesting that the segmentation
322 can be performed even after a significant amount of time from the formalin fixation. In order to
323 understand if these volume changes were induced by an overall shrinkage or inflation of the brain,
324 we performed the coregistration of the data acquired at the different time points.

325 As an example, in Fig 6, we report the borders of the brain extracted (for a representative slice) from
326 *in-vivo* data overlaid to the T1w images obtained *ex-vivo* (central panel).

327

328 **Fig 6. Data coregistration.** T1w *in vivo* data are used as the reference to coregister the *ex-vivo* data
329 with a 6-parameter coregistration approach (no scaling included). As an example, a representative
330 slice is reported (A). The *in vivo* brain contours are overlaid to the *ex-vivo* data after the coregistration
331 (B). It can be noted a good agreement. Analogously, the *in-vivo* brain contours overlaid on the *ex vivo*
332 12 months show that the rigid coregistration successfully aligned the data (C). Since no scaling was
333 involved in the data transformation, this suggests that the brain did not experience any significant
334 inflation/shrinkage.

335

336 It can be noted that a rigid co-registration with 6 parameters, thus excluding any scaling factor,
337 was able to coregister the brain successfully. Moreover, as it can be noted in Fig 6 (left panel), the
338 sample applies to the coregistration of the *ex-vivo* data after 12 months. The fact that no scaling factor
339 was needed to coregister the data, seems to suggest that the brain did not experiment with any
340 significant inflation/deflation over time. Therefore, the changes observed in the volumes were likely
341 due to a change in the signal contrast.

342 So far, the two samples of dogs have been treated separately. However, an interesting point would be
343 if the longitudinal changes observed on a single dog hold also for the sample of 11 dogs. To address
344 this aspect, in a future study we will perform the same longitudinal study on a sample of dogs.
345 However, with the data available at this stage, we tried to assess if there were statistical differences
346 between the volumes obtained from the single dog and the 11-dog sample. To this aim, we considered
347 the distribution of the volumes, structure by structure, obtained from the sample and we tested
348 whether the volumes obtained from the single dog (after 12 months in formalin) belonged to the same
349 distribution or not, i.e. if they were statistically different.

350 As it can be seen in Table 2, where we report the 95% confidence intervals and the test outcomes,
 351 apart from Ventricles, Right Hippocampus and Right Substantia Nigra all structures were not
 352 statistically different.

353

354 **Table 2. Comparison between the measurements obtained on the group of 11 dogs (entire head)**
 355 **after 12 months in formalin and the ones obtained on the single brain dog after 12 months in**
 356 **formalin**

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		Mean_11 dogs	CI_11 dogs	Mean_exvivo12 months	t test
Ventricles	T1	931.37	593.76-1251.2	1620.53	*
Caudate	T2 L	588.22	479.48-690.26	498,13	
Caudate	T2 R	586.75	472-681.36	496.75	
Hippocampus	T2 L	611.68	516.1-728.84	530.83	
Hippocampus	T2 R	611.95	533.14-690.48	524.04	*
Sub Nigra	T2 L	49.43	33.2-63.7	37.88	
Sub Nigra	T2 R	56.09	39.8-70.58	36.86	*
Lat Geniculate	T2 L	67.37	49.38-92.34	74.06	
Lat Geniculate	T2 R	70.97	53.84-89.58	68.00	
Med Geniculate	T2 L	83.41	54.88-112.1	65.03	
Med Geniculate	T2 R	89.37	63.14-109.16	70.11	
Putamen	T2 L	76.70	58.3-95.64	67.76	
Putamen	T2 R	70.81	51.62-912	69.14	
Globus Pallidus	T2 L	57.62	41.34-73.86	56.49	
Globus Pallidus	T2 R	59.90	36.72-72.48	57.63	

375 Mean values between the operators are reported for each structure for both the group of the 11 dogs
376 and the single brain dog after 12 months in formalin. A t-test revealed that only for the Ventricles,
377 Hippocampus right and Sub Nigra R a significant difference was obtained (marked as *) between
378 the two distributions.

379

380 The volumes obtained from the single dog, after 12 months in formalin, seem to be consistent
381 with the volumes obtained from the 11 dogs' sample. The mast majority, namely 80% of the
382 structures, considered for the longitudinal study, belonged to the same distribution of the 11 dogs.
383 This suggests that the considerations on the longitudinal changes observed on a single dog might hold
384 also for the larger sample. Although encouraging, this point needs to be validated with a larger sample
385 in a future study.

386

387

388 **Discussion**

389

390 This work has been conceived with a dual purpose. First, we assessed the feasibility of brain
391 segmentations on selected structures in the brain kept in formalin for one year, focusing on the
392 reproducibility of the segmentations as a function of the operator and their intrinsic variability within
393 the sample. This study was performed on a homogeneous sample of 11 dogs. Then, one dog (not part
394 of the previous sample) was used for the longitudinal evaluation of the effect of death and fixation
395 on MRI properties of the brain. We obtained that after one year in formalin the segmentations seem
396 reliable and mostly reproducible across operators. Of note, we observed that the time spent in formalin
397 increased the contrast for some specific structures, such as the Substantia Nigra.

398 The impact of Beagle cerebral models on MRI translational studies is currently increasing. For this
399 reason, several groups are interested in characterizing brain structures. To this aim, in [11,13,14,25]
400 canine MRI based atlases have been developed. However, at the current stage, these are either based
401 on a heterogeneous group of dogs, i.e. a mixed breed sample [3], or on just Beagle dogs, but
402 considering only WM, GM, and CSF, and not specific structures [14]. Further, while these atlases
403 were obtained from alive dogs, in [11] a mixed-breed atlas was obtained by coregistering *in vivo* with
404 *ex vivo* data. In this case, brains were extracted from the skull and fixed in 10% buffered formalin.
405 Compared to these works, in our study, for the first time to our knowledge, selected brain structures
406 have been segmented on brains fixed in formalin to provide their volumetric evaluation. This allows
407 assessing the reproducibility of such structures, an aspect that has not been previously evaluated. This
408 point is important since manual segmentation is still the more accurate approach, being automatic
409 tools not yet available for these applications [26]. Our findings suggest that the segmentation
410 procedure on this kind of data can be considered overall reproducible since only slight statistical
411 differences were detected, apart from the ventricles. This could be due to the fact that ventricles are
412 fluid filled structures, and they may not be clearly distinguished from the rest of brain CSF.

413 Furthermore, the fact that some structures are characterized by higher volume variations than others
414 may rely on an intrinsic individual variability. This aspect is particularly interesting, since single
415 structures may be the target of specific experiments, and to determine that there may be an intrinsic
416 variability could be crucial, e.g. in disentangling a pathology/drug induced effect vs a physiological
417 variation. Next to strictly experimental purposes, since the estimation of volumes can assist in the
418 detection and monitoring of the progression of brain disease/damage, the accuracy and reproducibility
419 of MRI segmentation are important also in the clinical field [27]. In fact, nowadays manual
420 segmentation for brain structures or lesions is considered the gold standard in terms of precision, even
421 if it may result in longer analysis times compared to a more “automatic” approach [28,29]. This is in
422 line with [14,22], where the mentioned brain templates were obtained from manually segmented data.
423 Compared to histopathological analyses, the MRI based segmentation of formalin fixed brains has
424 several advantages. First, structural abnormalities can be assessed within the entire brain without
425 altering the original structures. Second, the formalin fixation allows to analyze the data multiple times
426 in any plane to focus on specific areas [19,30] by preserving the structure of the tissues. In fact, after
427 death, the brain undergoes microbial degradation, autolysis, and breakdown of cell membranes. The
428 chemical fixation tends to preserve the macromolecular structure providing the longitudinal stability
429 required for extensive scanning times. Nevertheless, due to death and fixation, a series of artifacts
430 and changes are expected. As far as it regards eventual longitudinal studies, it is known that the
431 formalin fixation may alter the relaxation times. This is due to the induced tissue dehydration,
432 crosslinking, and reduced transmembrane water exchange [30] that lead to a T1/T2 shortening. This
433 results in a higher spatial resolution in terms of borders visualization, for technical details see for
434 example [19,31]. This is in line with what we observed in the *ex vivo* phases, where both operators
435 found an increase of the volumes, especially for the smallest structures, that were judged easier to be
436 segmented. To support this interpretation, we carefully checked the brain volumes by co-registering
437 the brains across the three phases. We obtained no significant changes. Therefore, this apparent
438 increase of volume did not correspond to a real increase in the size of the structure but to an increase

439 of contrast in the structures' borders allowing a more accurate segmentation. Evidently, before
440 fixation, the structures' volumes were under-estimated due to low contrast. This is in line with
441 previous findings, see for example [32] where it was reported that the qualitative image evaluation
442 significantly improved after fixation, and the structures segmentations were described to be easier
443 than *in vivo* images.

444 Another effect of formalin fixation that has been reported in the literature is the tissue shrinkage
445 which may not be homogeneous among the various brain structures, especially when MRI is
446 performed after fixation, see [22]. For example, it has been observed that ventricles may experience
447 filling or emptying according to pressures from the surrounding tissue, and shrinkage of surrounding
448 tissue may not always be paired with an expansion of the ventricles [33] with those tissues
449 experiencing a “positive formalin effect”, characterized by a swelling effect caused by the osmotic
450 pressure of the formalin solution [34]. These aspects should be taken into proper considerations when
451 volumetric evaluations are performed on brains in *post mortem* or *ex vivo* conditions. These effects
452 may relate to the observed variability of the ventricles in our data.

453 Finally, according to our findings, the amount of time (12 months) spent in formalin seems not to
454 influence the volumes of the structures: their percentage variation did not exceed 10%. This aspect is
455 important since it suggests that the same brain can be potentially used for several studies, even after
456 some time, without the risk of significant structural changes. This observation is consistent with data
457 proposed by other studies since previous investigations on the relationship between MR volumetric
458 measurements performed in-vivo and ex-vivo report that the brain structures remain relatively stable
459 for 6 months post-mortem [35]. This applies also to human medicine where it has been reported that
460 fixation leads to no significant leaching of iron in long term storage [36] and that WM components,
461 including myelin, are all well preserved [37].

462

463 To summarize, based on our findings *post mortem* MRI based segmentation seems to be a useful and
464 accurate tool that allows longitudinal studies. However, especially for the observed longitudinal
465 variations, these findings need to be further validated on a larger sample.

466

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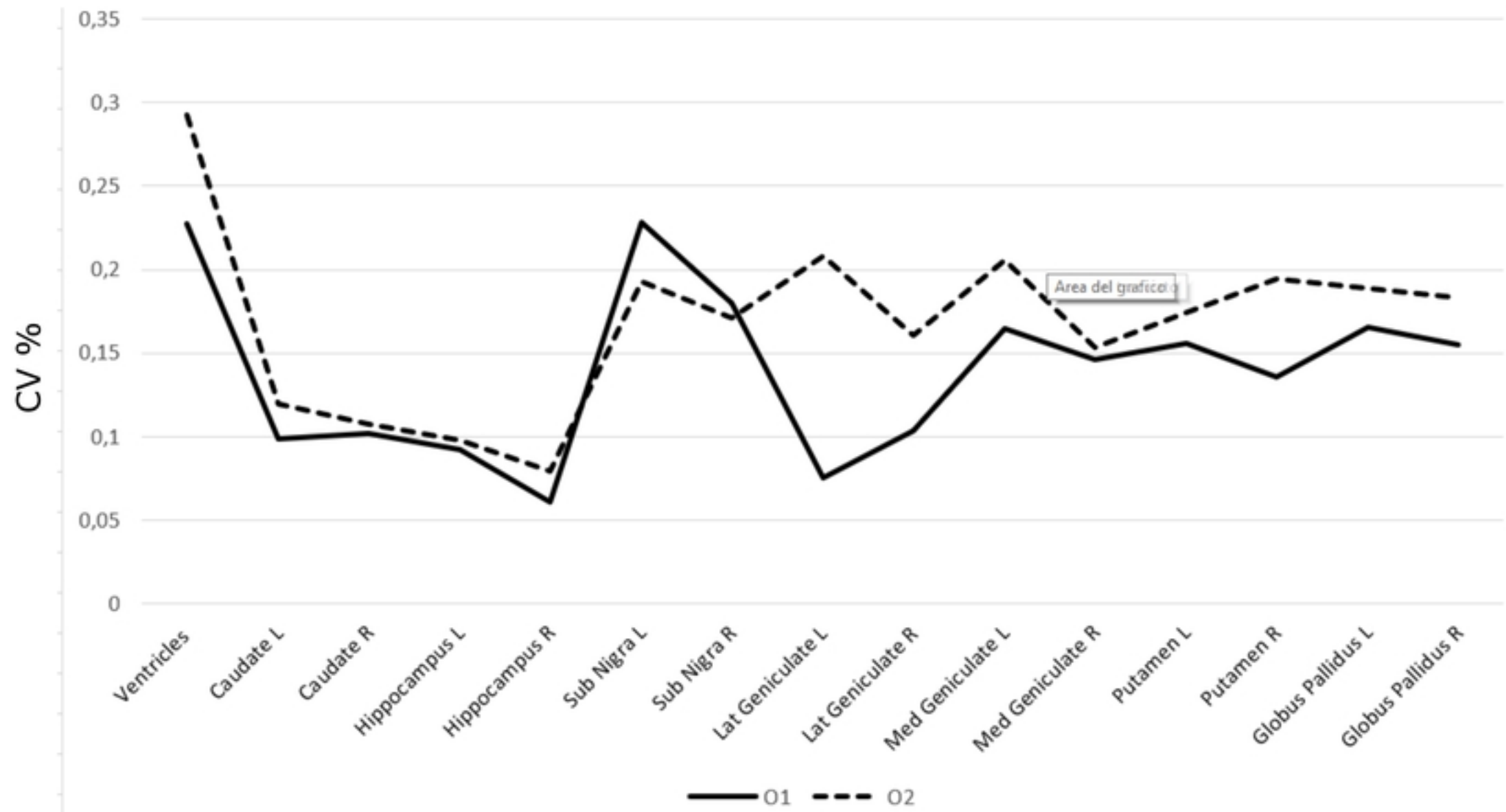


Figure 1

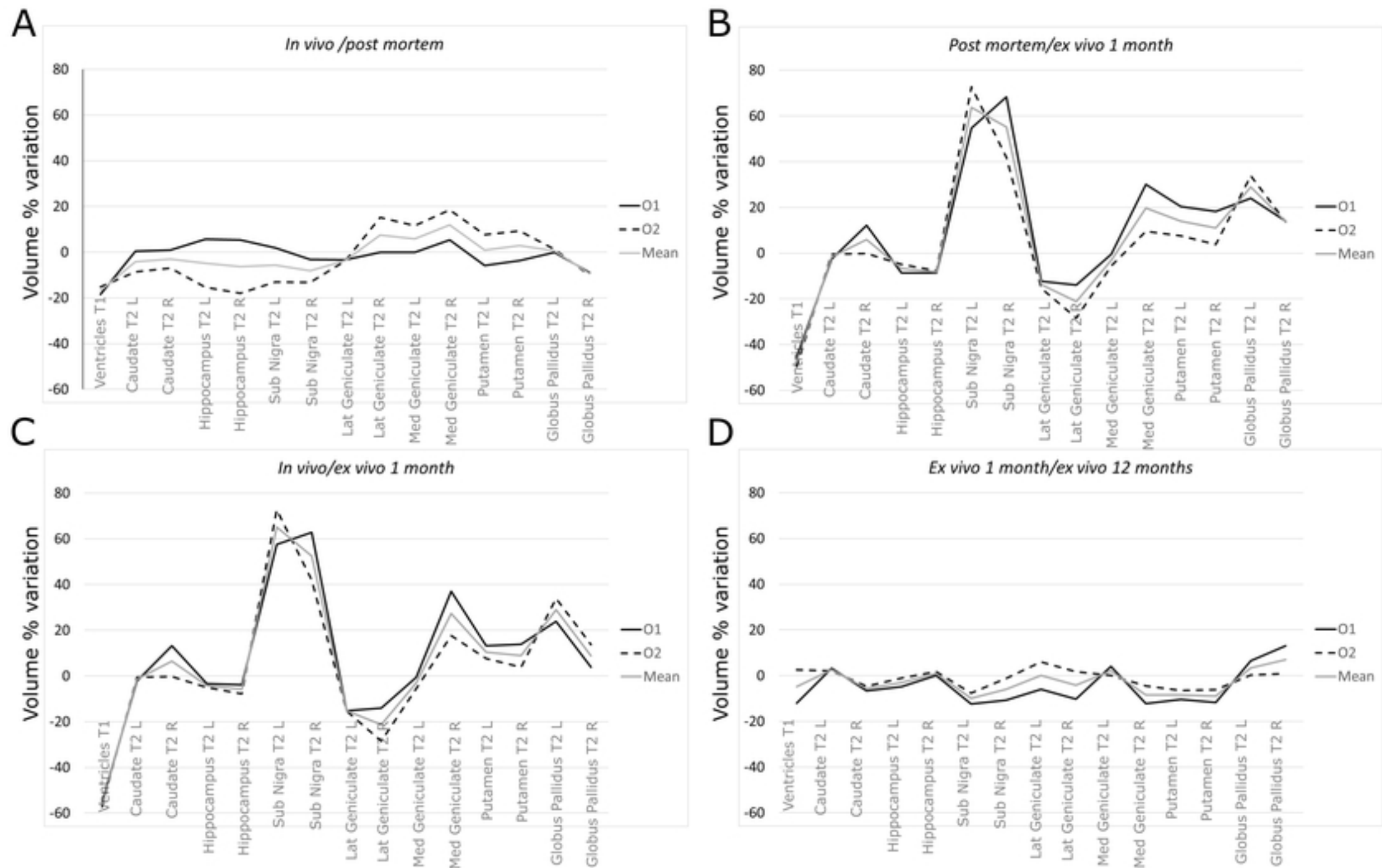
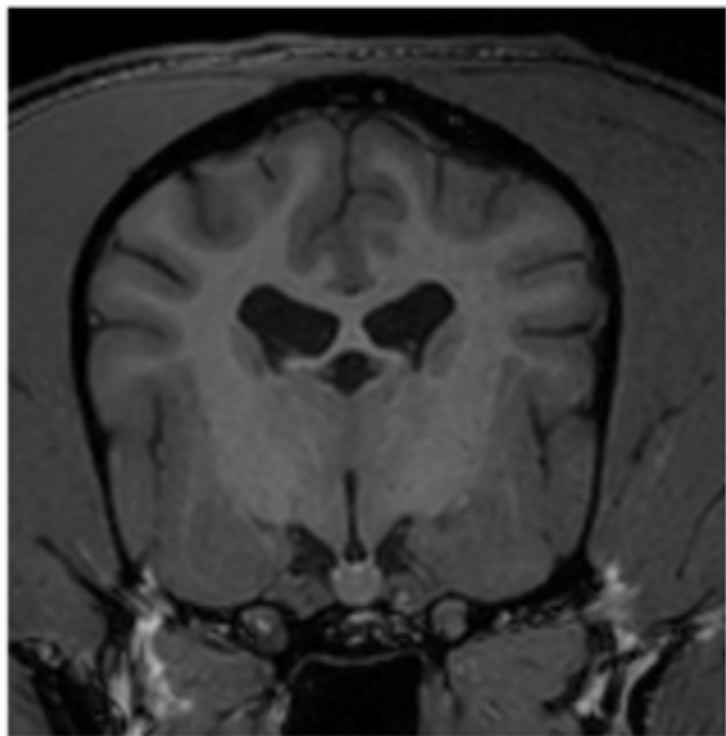
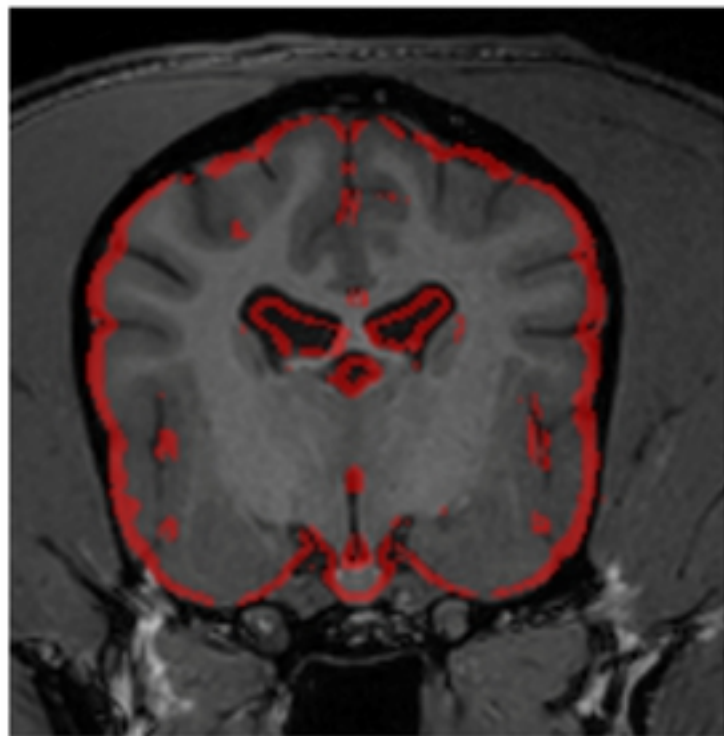


Figure 5

Brain MRI T1w In-Vivo



Rigid Coregistration Overlay



Brain MRI T1w Ex-Vivo + EDGE (RED)

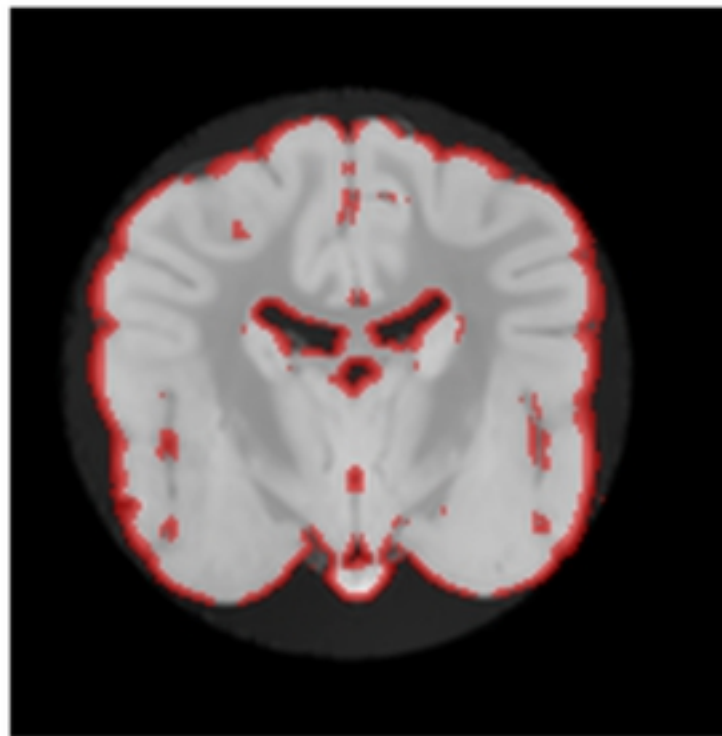


Figure 6

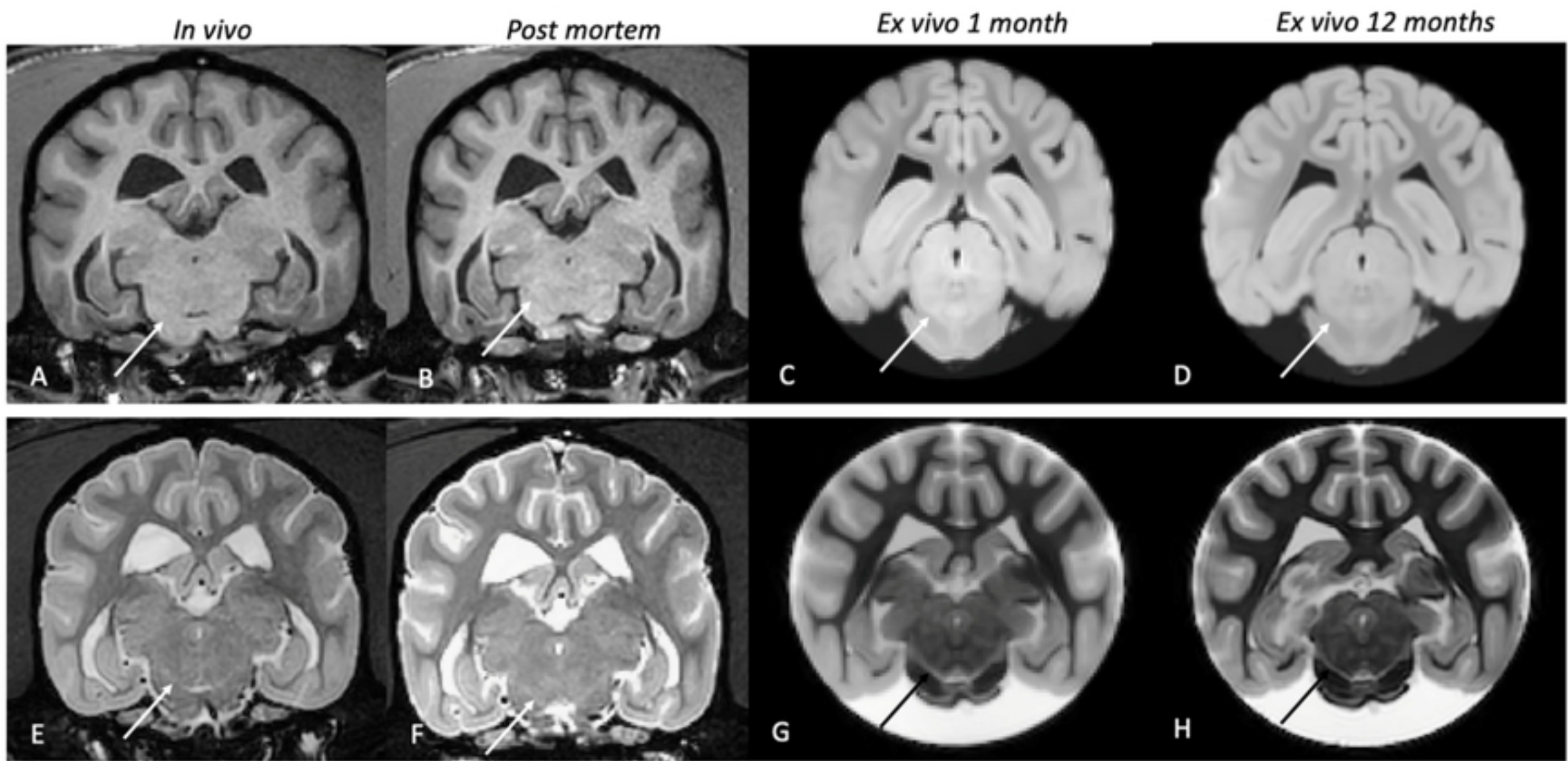


Figure 4

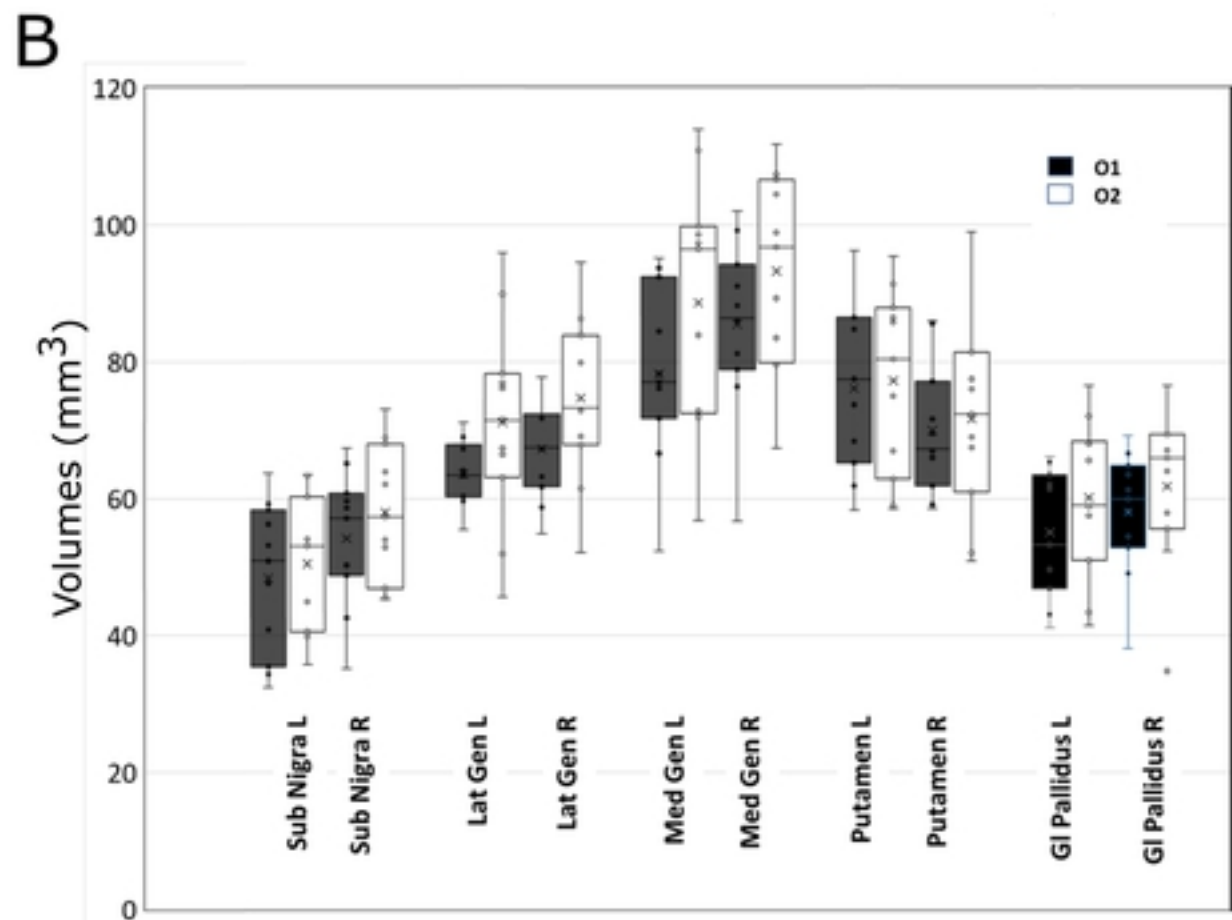
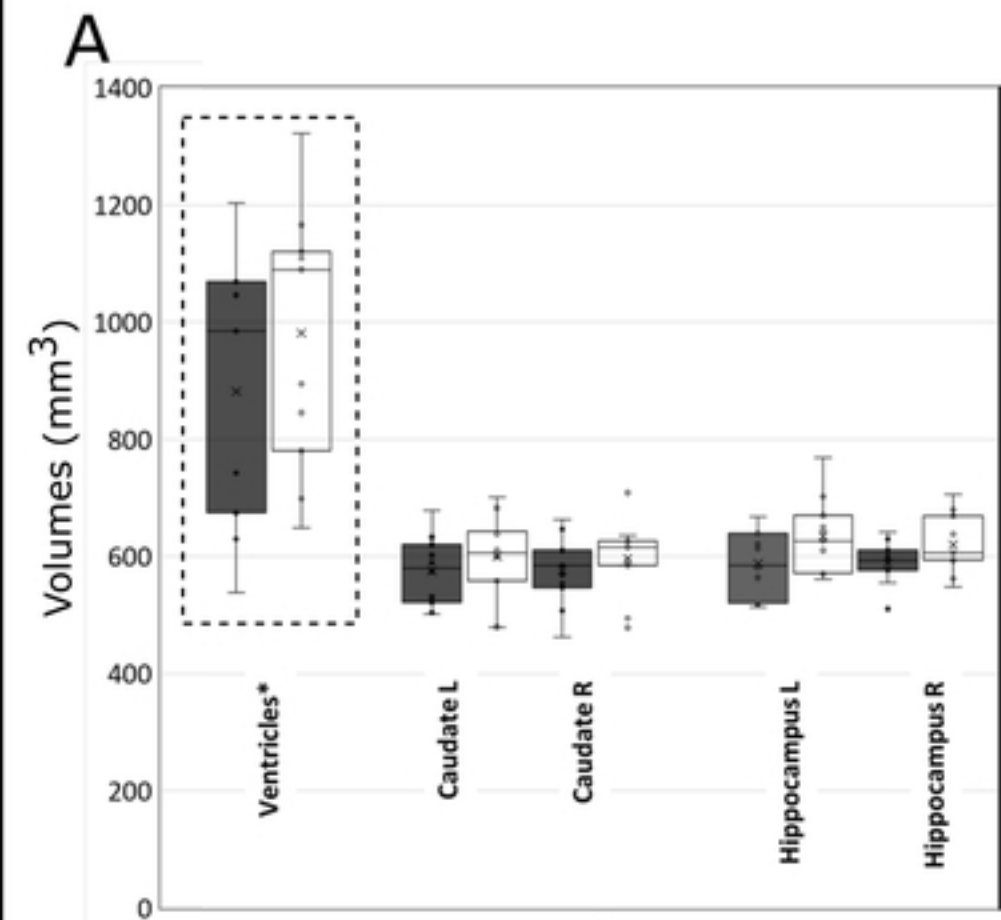


Figure 3

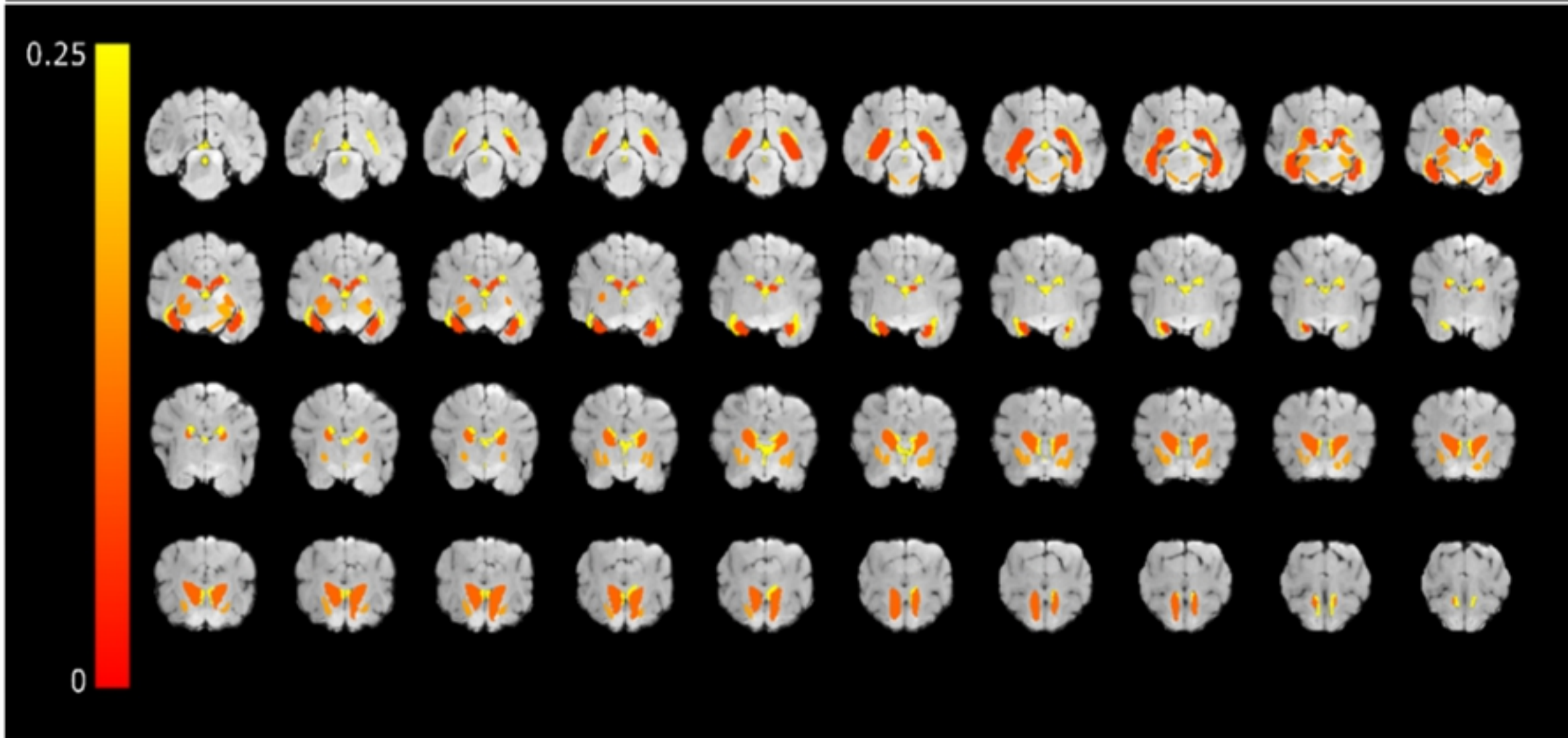


Figure 2