1	Brai	n volumetric changes in the general population following the COVID-19 outbreak and	
2		lockdown	
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20			

21 Abstract

The COVID-19 outbreak introduced unprecedented health-risks, as well as pressure on the 22 financial, social, and psychological well-being due to the response to the outbreak 1^{-4} . Here, we 23 24 examined the manifestations of the COVID-19 outbreak on the brain structure in the healthy population, following the initial phase of the pandemic in Israel. We pre-registered our 25 hypothesis that the intense experience of the outbreak potentially induced stress-related brain 26 modifications⁵⁻⁸. Volumetric changes in n = 50 participants scanned before and after the 27 COVID-19 outbreak and lockdown, were compared with n = 50 control participants that were 28 scanned twice prior to the pandemic. The pandemic provided a rare opportunity to examine brain 29 plasticity in a natural experiment. We found volumetric increases in bilateral amygdalae, 30 putamen, and the anterior temporal cortices. Changes in the amygdalae diminished as time 31 elapsed from lockdown relief, suggesting that the intense experience associated with the 32 pandemic outbreak induced volumetric changes in brain regions commonly associated with 33 stress and anxiety $^{9-11}$. 34

35

36 Main text

Since early 2020, the world has been coping with the outbreak of the coronavirus disease 2019 (COVID-19) pandemic that has infected millions with devastating numbers of deaths globally. As an initial response to the first wave of the outbreak, countries closed their borders and implemented a series of ad-hoc laws and orders to restrict the spread of the disease. Countries with major outbreaks such as China, Italy, and Spain enforced stringent restriction of movement for a limited period, referred to here as 'lockdown'. Although lockdowns contributed to restricting the health risks of the outbreak¹², they also had a negative impact on the social, economic and psychological well-being of the general population, leading to one of the sharpest
declines in economic growth over the past decades^{1,2}. Lockdowns also led to high rates of stress
and anxiety which were attributed in large to the social and financial consequences of responding
to the health crisis⁴. It is now evident that the indirect consequences of the pandemic affected a
much larger proportion of the population, having an impact of no lesser gravity than the actual
health risks that were meant to be prevented^{3,13}.

In Israel, a strict lockdown period was issued from mid-March until the end of April. During its 50 peak, most unessential businesses were closed and civilians' movement for non-essential 51 destinations was restricted for a radius of 100 meters from their homes. Prior to COVID-19, the 52 country had experienced a period of peak economic prosperity¹⁴, which was interrupted by the 53 outbreak, leading to unprecedented unemployment rates (reaching nearly 30% of the work-force 54 in April 2020) and the collapse of several sectors such as aviation, tourism, and culture^{15,16}. The 55 outbreak period was characterized with acute uncertainty and increase in anxiety, regarding both 56 the health and socioeconomic effects of the pandemic 17 . 57

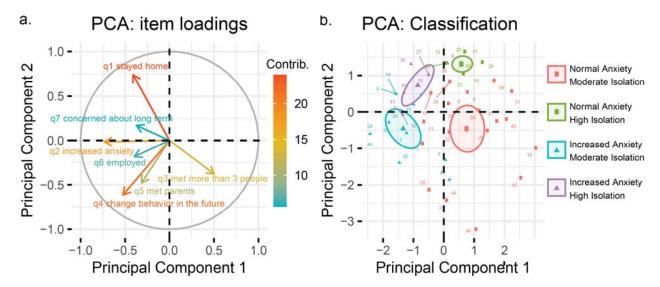
58 Over the past years several studies demonstrated brain plasticity detected using T1-weighted magnetic resonance imaging (MRI)¹⁸⁻²⁰. The current work was initiated as a reaction to the 59 outbreak of COVID-19 in Israel, aimed to study the structural brain plasticity in the general 60 population following a real-life event of global scale. For this purpose, we examined n = 50 test 61 group participants that were scanned with T1-weighted MRI prior to the outbreak and returned 62 for a follow-up scan after the lockdown period. The structural changes of the study group (before 63 versus after the outbreak) were compared to those of n = 50 control participants who were 64 scanned twice before the COVID-19 outbreak. The unique circumstances imposed due to the 65

66 COVID-19 lockdown created rare settings for a natural experiment to examine the effect of a
67 real-world intense event on brain plasticity.

All participants were healthy, without a history of neurological or psychiatric disorders, did not show COVID-19 symptoms, and were not diagnosed carrying the virus (see the methods section for further demographic information). The hypotheses and general design of the current study were pre-registered prior to the completion of data collection and were based on a small independent pilot sample with N = 16 participants (n = 8 participants in each group). The data and analysis codes are openly shared online (project page: https://osf.io/wu37z/; preregistration: https://osf.io/k6xhn/).

Prior to their follow-up MRI scan session, we asked participants of the post-lockdown test group 75 to fill in a short questionnaire regarding the lockdown period (see methods). Of the participants 76 77 who agreed to reply, 79.6% reported they did not leave their home for non-essential needs, 38.8% indicated an increased feeling of anxiety following the lockdown, 79.6% met no more 78 79 than 3 people, 34.7% anticipated that their future behavior will change after the lockdown, 80 44.9% did not meet with their parents at all, 42.9% indicated that their employment status was reduced to part-time or unemployment, 46.8% reported they were concerned about their personal 81 future well-being. In an exploratory principal component analysis (PCA), we found that two 82 principal components best explain the variability in the data, explaining together 42.58% of the 83 variance. The first component was highly loaded with increased feelings of anxiety, and the 84 85 second was related to items describing increased social isolation (Figure 1). These analyses (not 86 included in the pre-registration) indicate that the pandemic outbreak had a significant impact on the social and psychological well-being of most participants in our study. 87

88





90 **Figure 1.** Principal component analysis of COVID-19 questionnaire.

A principal component analysis (PCA) of the responses to the questionnaire revealed two main 91 themes characterized the participants. a. An increased feeling of anxiety dominated the first 92 93 principal component (x-axis), while the three items relating to social distancing (staying at home, meeting parents and meeting more than 3 people) contributed together to the second component 94 95 (y-axis). **b.** Visualization of participants dispersion across the two principal components and their categorization into binary anxiety and isolation groups. High isolation was defined as directional 96 response to all three social-isolation items described above. Points represent individual 97 98 participants.

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100 Based on our pilot study results and previous studies of stress-related morphological brain changes⁵⁻⁸ we hypothesized that the focus of volumetric changes will be observed mainly in the 101 amygdalae. The anatomical data were used as input for deformation and surface-based 102 morphometry (SBM) analysis using the CAT12 toolbox (http://www.neuro.uni-jena.de/cat/, 103 104 University of Jena) for SPM12 (http://www.fil.ion.ucl.ac.uk/spm/software/spm12/, Wellcome Trust Centre for Neuroimaging). The brain was segmented to 58 regions based on the cortical 105 and subcortical nuclei classifications of the Hammers atlas (Hammers et al., 2003). Following 106 107 surface reconstruction, each participant's individual gray matter volume was estimated for each of the 58 anatomically defined regions of interest (ROIs). This procedure accounted for the 108 longitudinal nature of the data, performing the analysis on both scans simultaneously. To avoid 109 voxel-based multiple comparisons, we performed a region-based analysis (following surface 110

111 projection to the Hammer atlas) and corrected for multiple comparisons using the Benjamini-112 Hochberg correction²¹ to control for false discovery rate (FDR; at p < 0.05 following correction). 113 Validation of this pipeline was performed using simulated data and by comparing the results with 114 other software (see methods).

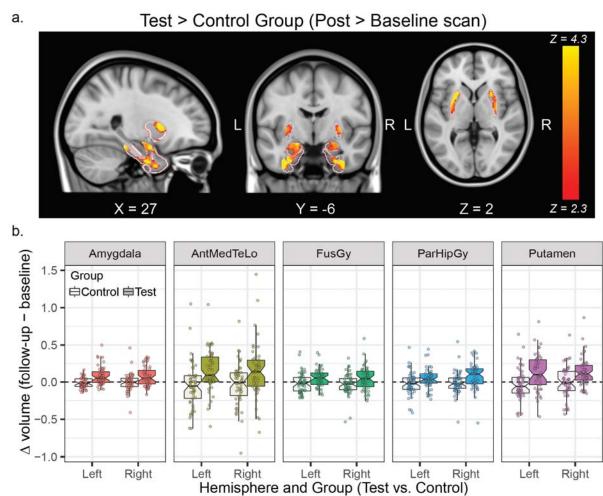
Using a linear mixed model, we examined volumetric changes, testing for regions with stronger 115 effects for the test group, compared to the control group. Examining an interaction effect of 116 117 session (baseline versus follow-up scans) and experimental group (test versus control) revealed ten anatomical brain regions (composed of bilateral five unique regions in both hemispheres) in 118 119 which volumetric increases were observed uniquely for the test group (Table 1 and Figure 2). Most prominently, as we expected and pre-registered, we found a robust effect in the bilateral 120 amygdalae. We also observed a significant effect bilaterally in the putamen, and in three 121 122 anatomical regions within the ventral anterior temporal cortex adjacent to each other, namely in 123 the medial part of the anterior temporal lobe, the fusiform gyrus, and the parahippocampal gyrus. 124 To examine the spatial distribution within significant ROIs, we performed an additional post-hoc 125 voxel-based analysis, which allowed us to visualize the changes within the significant ROIs (Figure 2a). Examining the post-hoc voxel-based results revealed that volumetric changes 126 occurred throughout the entire surface of bilateral amygdalae, while in the putamen the effects 127 occurred mainly in the dorsal area. In the ventral anterior temporal cortices, large connected 128 clusters of volumetric change spanned throughout the three adjacent temporal ROIs, thus 129 130 suggesting that the three ROIs shared a similar origin. In all ROIs, we ensured that the significant 131 effect was apparent for the test group but not for the control group (see methods), suggesting that the reported interaction effects originated from volumetric changes in the test group following 132 133 the COVID-19 outbreak and its related lockdown period (Figure 2b).

Region	Hemi- sphere	Interaction estimate (95% CI)	Interaction <i>p</i> (FDR adj.)	Session estimate (95% CI) ^a	Session <i>p</i> (FDR adj.)
Amygdala	Left	0.09 [0.05, 0.13]	2.4E ⁻⁵ (0.001)	0.08 [0.05, 0.11]	9.8E ⁻⁶ (2.1E ⁻⁴)
	Right	0.08 [0.03, 0.13]	0.003 (0.030)	0.08 [0.05, 0.11]	1.6E ⁻⁵ (2.3E ⁻⁴)
Putamen	Left	0.19 [0.09, 0.29]	4.1E ⁻⁴ (0.006)	0.13 [0.06, 0.2]	4.0E ⁻⁴ (0.002)
	Right	0.17 [0.08, 0.26]	2.4E ⁻⁴ (0.005)	0.14 [0.08, 0.2]	1.1E ⁻⁵ (2.1E ⁻⁴)
Anterior temporal lobe (medial part)	Left	0.25 [0.12, 0.38]	1.8E ⁻⁴ (0.005)	0.15 [0.07, 0.23]	4.7E ⁻⁴ (0.003)
	Right	0.21 [0.07, 0.35]	0.004 (0.030)	0.15 [0.05, 0.25]	0.004 (0.023)
Parahippocampal gyrus	Left	0.09 [0.03, 0.15]	0.006 (0.035)	0.04 [0, 0.08]	0.029 (0.085)
	Right	0.11 [0.04, 0.18]	0.003 (0.030)	0.08 [0.03, 0.13]	0.002 (0.009)
Fusiform gyrus	Left	0.08 [0.03, 0.13]	0.007 (0.036)	0.06 [0.03, 0.09]	3.8E ⁻⁴ (0.003)
	Right	0.11 [0.04, 0.18]	0.002 (0.022)	0.05 [0, 0.1]	0.044 (0.111)

134 Table 1. Surface based morphology analysis results

^a Session estimate examined the effect of baseline versus follow-up scan in the post-lockdown test group. This parameter was used to validate that the interaction effect observed between the group stemmed from a robust effect in the test group (see methods).

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140 Figure 2. Volumetric changes results.

An interaction effect for time (baseline versus follow-up scan) and group (test versus control) 141 was evaluated on segmented surfaces in an SBM analysis. Significant interaction effects were 142 143 observed bilaterally in the amygdala and putamen ROIs, as well as in three ventral temporal cortical ROIs. a. To examine spatial patterns within the identified ROIs, a post-hoc voxel-based 144 analysis was conducted within each ROI mask. Light red contours represent segmentation 145 borders of the ROIs. **b.** Individual distribution of the results in the control group (light colors) 146 and test group (dark colors). Box-plot center, hinges, and whiskers represent the median, 147 148 quartiles, and from the hinges, respectively. A notch of represent an estimated 95% confidence interval for comparing medians. Dots represent individual 149 150 participants. Abbreviated ROI names: AntMedTeLo = anterior temporal lobe (medial part); 151 FusGy = fusiform gyrus, ParHipGy = Parahippocampal gyrus.

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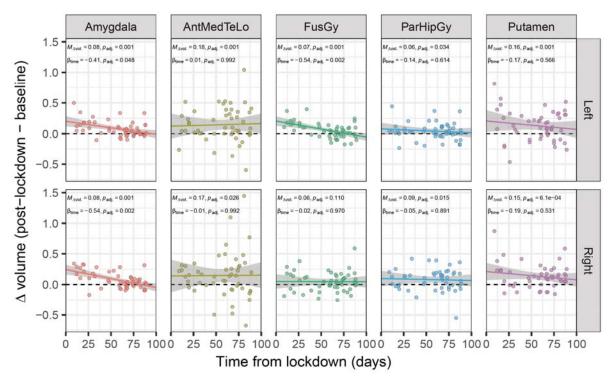
153 To evaluate and control for the effect of time between scans and time from lockdown, we

included in the model two additional covariates - the time between scans (TBS; which was

155 generally longer for the test group) and time following lockdown (TFL; see supplementary

156 methods for more details). The two covariates were not correlated with each other in our test

group sample (r = -0.106, t(48) = -0.74, p = 0.463). Our reported regions demonstrated 157 significant volumetric change above and beyond these covariates. After FDR correction, no 158 region showed an effect of TBS. However, we did find a negative effect of TFL in the two 159 amygdalae ROIs and the left fusiform gyrus, suggesting that the volumetric changes in these 160 regions moderated as time following lockdown elapsed. Based on these results we estimated the 161 time to decay as the estimated number of days from lockdown until volumetric changes returned 162 to normal levels, similar to those of the control group (left amygdala: $\beta_{\text{TFL}} = -0.41$, t(47) = -3.1, p 163 = 0.003, p_{adi} = 0.048, time to decay = 95 days; right amygdala: β_{TFL} = -0.54, t(47) = -4.38, p = 164 $6.7E^{-5}$, $p_{adi} = 0.002$, time to decay = 83 days; left fusiform gyrus: $\beta_{TFL} = -0.54$, t(47) = -4.44, p = -0.54165 $5.5E^{-5}$, $p_{adi} = 0.002$, time to decay = 82 days; Figure 3). 166



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168 **Figure 3.** Time following lockdown effect on volumetric changes.

169 The time from lockdown relief until the follow-up scan session (TFL) was introduced as addition 170 covariate to the model, revealing significant effect in the two amygdalae and left fusiform gyrus.

171 Points represent individual participants in the post-lockdown test, *p*-values were FDR adjusted

172 for multiple comparisons. Abbreviated ROIs: AntMedTeLo = anterior temporal lobe (medial

part); FusGy = fusiform gyrus, ParHipGy = Parahippocampal gyrus.

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Additional exploratory analyses examined the association of volumetric changes and the reported
experience during lockdown. We found no strong association between the two, as reported in the
supplementary results.

In conclusion, our study demonstrates that volumetric change patterns in the brain occurred following the COVID-19 initial outbreak period and restrictions. Previous studies demonstrated brain plasticity using T1-weighted MRI following planned interventions^{18–20}. The current work uniquely demonstrates stark structural brain plasticity following a major real-life event.

182 Our findings show changes in gray matter in the amygdala, putamen and ventral anterior temporal cortex. The changes in the amygdalae showed a temporal-dependent effect, related to 183 the time elapsed from lockdown but not the duration from the baseline scan. It should be noted 184 185 that although lockdown restrictions had initially reduced infection rates in Israel, just one month 186 after the lockdown was lifted, the number of infected cases started to rise again and reached higher number of active infected cases by the end of data collection, compared with the peak 187 numbers during the actual lockdown period (approximately 2,000 daily new cases by the end of 188 July versus under 750 new daily cases during the peak of the lockdown period in April²², see 189 190 supplementary Figure 1). This suggests that the effects observed in the current study are less 191 likely to be attributed to the concrete health risks of COVID-19, but rather to the first wave of the outbreak, characterized with perceived uncertainty. 192

193 The current study was in many aspects unplanned; thus we are left with only partial answers as 194 to which specific components of the COVID-19 outbreak led to the neural changes observed in 195 the healthy participants that took part in our study. The involvement of the amygdala may 196 suggest that stress and anxiety could be the source of the observed phenomenon, due to its well-197 recorded functional and structural associations^{5–11}. Nevertheless, it is hard to draw clear 198 conclusions as many aspects of life have changed in this time period, and could have potentially 199 affected different regions in the brain – from limiting social interactions, increased financial 200 stress, changes in physical activity, work routine, and many more. The limited behavioral data 201 collected in the current study did not provide a strong connection to the imaging results, and thus future work could try to better address the complex brain-behavioral associations in this real-life 202 experience. Nonetheless, our findings show healthy young adults, with no records of mental 203 204 health issues, were deeply affected by the outbreak of COVID-19. We suggest that policy makers 205 take into consideration the impact of their actions on the general well-being of the population 206 they seek to help, alongside the efficacy of disease prevention.

207

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215 Author contributions

T.Sa. wrote the manuscript with Y.A, assisted with the study dsign and analyzed the data. Data
was collected by A.C., R.G, S.O, G.R., D.R, and A.S. Free-surfer analysis was made by G.B-Z.
N.T performed VBM validation analysis. G.T. provided support and medical supervision. I.T.,
T.Sc. and Y.A. conceived the study, wrote the manuscript and supervised the study. Y.A

- 220 performed the preprocessing and analysis of imaging data. All Author contributed intellectually
- and reviewed the manuscript.

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223 **Competing interests**

224 The authors declare no competing financial interests.

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Supplementary Methods

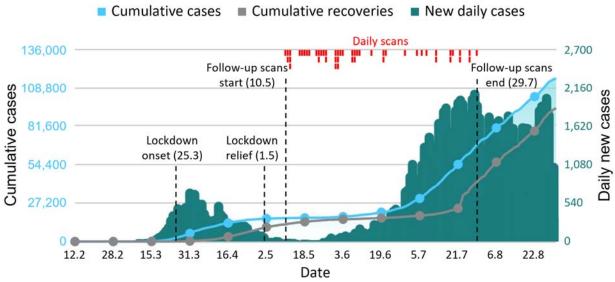
292 Codes and Data Accessibility

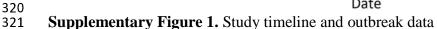
Our sample size, hypotheses and analyses plan were pre-registered on the Open Science Framework (OSF), soon after data collection began, but prior to completion of the data collection and data analysis (project page: <u>https://osf.io/wu37z/;</u> preregistration: <u>https://osf.io/k6xhn</u>). All behavioral processed imaging data along with the analysis codes are shared on the OSF project page. Uncorrected and small-volume corrected statistical maps of the voxel-based results described in the current work are available at https://neurovault.org/collections/8591/.

299 **Participants**

300 The study included two groups: A test group scanned before and after COVID-19 lockdown, and 301 a control group, scanned twice before COVID-19 lockdown. All participants had no background of neurological disorders, did not show symptoms for COVID-19 and were not diagnosed as 302 303 carriers of the virus. The study was approved by the ethics committee of Tel Aviv University and institutional review board (IRB) at the Sheba Tel-Hashomer medical center. Since the IRB 304 305 protocol allowed us to scan the participants several times over a long period of time, we were able to collect the data from participants who were scanned prior to COVID-19 outbreak and 306 307 invite them back for a follow-up scan as part of the longitudinal study they have agreed to take part in. Participants received monetary compensation for their time and gave their informed 308 consent to take part in a longitudinal experiment aimed to examine brain plasticity across several 309 310 sessions, which was initially not directly related to COVID-19 outbreak.

311 The test group comprised of n = 50 participants who were scanned before and after COVID-19 lockdown (Δ Time between scans: M = 309.3, SD = 207.5, range = 67 - 1460 days; Age: M =312 313 30.1, SD = 6.65, range = 21 - 48; Females: n = 20, prop. = 40%). The lockdown period began on March 25th, and was gradually relieved throughout late April. We mark here May 1st as the 314 lockdown relief date, as on this day an issued 100-meters movement limit for non-essential needs 315 was lifted. The test group data collection started as soon as lockdown relief took place, for a 316 period of approximately 3 months, until the end of July, 2020 (Δ Time from lockdown relief: M 317 = 57, SD = 24.62, range = 9 - 89 days; see Supplementary figure 1 for the study timeline). 318 319





On February 21st, 2020, the first COVID-19 case in Israel was recorded. Daily new cases are 322 presented on green bars (right y-axis), along with the cumulative number of cases and recoveries 323 (left y-axis). Data was retrieved and modified based on the Israeli Ministry of Health reports²². A 324 lockdown was issued on March 25th, which was gradually released until the removal of the 100-325 meter restriction on May 1st, marking lockdown onset and relief, respectively (shorter vertical 326 dashed line). MRI data of the test group was collected from May 10th to July 29th (longer vertical 327 328 dashed line). Red bars on top represent the number of participants scanned for the study each 329 day.

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As a control group, we used the data of n = 50 participants who were scanned twice using a
similar protocol before COVID-19 lockdown (\Delta Time between scans: M = 126.7, SD = 190.4,
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range = 21 - 886 days; Age: M = 27.3, SD = 5.63, range = 19 - 42; Females: n = 23, prop. =

334 46%).

335 The minimal sample size was determined and pre-registered (https://osf.io/uktsn), based on a 80% power analysis conducted using R 'pwr' package²³, on a pilot study with N = 16336 337 participants (n = 8 in each group). We decided to collect a minimum of n = 37 participants which should provide 80% to detect the group and session interaction effect with $\alpha = .05$, within both 338 339 the left and right Amygdala. We originally committed to collect n = 37 participants in each group, under the assumption it would be difficult to complete the sample due to COVID-19 340 341 limitations. Eventually, thanks to further relief in COVID-19 restrictions, we were able to extend the sample size to n = 50 in each group. 342

The results remain generally consistent, even when the data included only the first n = 37participant; demonstrating significant effects in the bilateral amygdalae, putamen, parahippocampal gyrus and the left anterior temporal lobe. In this smaller sample, we did not find significant effects in the right anterior temporal lobe, nor in the fusiform gyrus. We also found volumetric increase effects that were not identified using the full sample in the left nucleus accumbens, left cuneus, and left insula.

349 Imaging data

Acquisition protocol. Imaging data were acquired using a 3T Siemens Prisma scanner, with a 64-channel head coil. For the structural data, T1w high resolution (1-mm³) whole brain images were acquired with a magnetization prepared rapid gradient echo (MPRAGE) pulse sequence with repetition time (TR) of 2.53s, echo time (TE) of 2.88ms, flip angle (FA) = 7°, field-of-view (FOV) = $224 \times 224 \times 208$ mm, resolution = $1 \times 1 \times 1$. (see below).

Some participants were also scanned with diffusion-weighted echo-planar imaging (DW EPI) sequence and some with functional gradient-echo EPI (GE EPI) in a resting state scan. The analyses of these scans are beyond the scope of the current study.

358 Data processing and analysis. The T1w MPRAGE anatomical scans were used for a surface-359 based morphometry (SBM) analysis. From the images we estimated the pial and inner surfaces 360 of the cortex and projected those into the Hammers atlas (Hammers et al., 2003). Data were preprocessed in SPM12 (http://www.fil.ion.ucl.ac.uk/spm/software/spm12/, Wellcome Trust 361 362 Centre for Neuroimaging) and SPM based CAT12 (Computational Anatomy Toolbox 12; http://www.neuro.uni-jena.de/cat/, University of Jena) extension. We deployed the CAT12 363 surface-processing pipeline, which includes skull striping, a denoising filter²⁴ projection-based 364 thickness estimation²⁵, partial volume correction, and spatial normalization to MNI space. 365

366 Surface-based volumetric data of cortical and subcortical regions were segmented based on the Hammers Atlas, segmenting the volumetric data into 58 anatomically defined regions. Ten 367 368 additional ROIs of non-gray matter (ventricles, white matter, brain-stem and cerebellum ROIs) were excluded from statistical analyses. To evaluate the effect of lockdown on volumetric 369 370 imaging data we ran a mixed linear model on the data within each one of the 58 anatomical regions, examining the effect of session (baseline versus follow-up scan) and group interaction 371 372 (test versus control), controlling for time between scans (TBS) and time following lockdown (TFL) covariates. Both covariates were mean centered before they were added to the model. 373

374 To identify our regions of interest we included only regions that showed both a significant interaction effect (i.e. the volumetric difference between the two sessions was significantly 375 376 different for the test and control group), and a significant session effect within the test group (i.e. 377 a significant difference between the two sessions for the test group). Results were corrected for multiple comparisons using the false discovery rate (FDR) correction, based on the number of 378 brain regions tested²¹. Following the analysis pipeline, we identified ten significant ROIs: 379 bilateral amygdalae, putamen, parahippocampal gyrus, the medial part of the anterior temporal 380 lobe, and the fusiform gyrus. 381

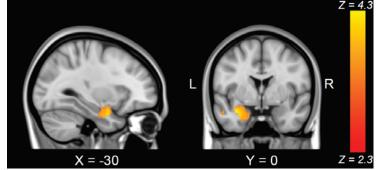
An additional ROI of the right inferior and middle temporal gyri showed a significant interaction 382 effect (interaction estimate = 0.17, 95% CI = [0.05, 0.29], $p = 6.1E^{-3}$, $p_{adi} = 0.035$), however 383 examining the test group separately, we could not identify a significant session effect (session 384 estimate = 0.04, 95% CI = [-0.02, 0.10], p = 0.259, $p_{adj.} = 0.424$). Therefore, it is harder to 385 interpret that this interaction effect stemmed from the test group. Thus, we did not include this 386 ROI as one of our significant ROIs. A less robust session effect within the test group was also 387 observed for the left parahippocampal gyrus (session estimate = 0.04 [0.00, 0.08], p = 0.029, p_{adi} . 388 = 0.085) and right fusiform gyrus ROI (session estimate = 0.05 [0.00, 0.10], p = 0.044, $p_{adj} =$ 389 0.111), however as these ROIs demonstrated strong interaction effects and their session effects 390 391 were significant before FDR correction, we decided to report them together with the other 392 significant regions. A similar procedure was used in the pre-registration (i.e. including only 393 regions that showed a significant interaction and an effect in the test group), however in the analysis of the pilot for the pre-registration, we used uncorrected results due to the small sample 394 size. 395

To examine the spatial distribution of our effect within significant ROIs, that were identified with the SBM analysis, we performed an additional post-hoc voxel-based analysis. We projected the data on a voxel-based map, effectively examining which voxels demonstrated an interaction effect. Then, we used anatomical masks of ROIs which were found to be significant in the SBM analysis, to visualize the results within these regions, similarly to a small-volume correction analysis (Figure 2a)

402 **Pipeline validation**

To validate the imaging processing protocol we used two approaches, before data collection was 403 completed (at the time of the pre-registration finalization). As many surface-based software 404 focus on analysis of cortical surfaces, rather than subcortical regions, we aimed to validate that 405 our SBM protocol using CAT12 could reliably identify subcortical morphological changes, such 406 as the ones we observed in the current study within the amygdala and putamen. To test the 407 detection capabilities of our protocol, we generated simulated data with volumetric changes in 408 the amygdala and ran the pipeline on the simulated data. A volumetric increase in the amygdala 409 was simulated using a hand-drawn polygon mask, surrounding the left amygdala on the original 410 T1w images. Within this polygon 3D mask, the signal intensity was artificially changed. 411 Following this procedure, both the original and modified T1w images underwent the same 412 413 CAT12 pipeline. The analysis was performed on 10 participants. We were able to identify the simulated volumetric changes within the subcortical amygdala nuclei (see supplementary Figure 414 415 2).

Simulated data in left amygdala



416

417 **Supplementary figure 2.** Data simulation

418 Sub-cortical changes in the left amygdala were simulated in ten participants. The SBM pipeline 419 applied in CAT12 identified the simulated change effect. Data was projected from surfaces back

- 420 to a voxel-based map for visualization in the current figure.
- 421

In an additional validation procedure, we reanalyzed our results with an additional analysis pipeline. Raw T1-weighted maps were preprocessed using FreeSurfer. Using this alternative analysis pipeline, we found similar results, including robust effects on the temporal cortical regions in addition to other cortical regions. It should be noted that the FreeSurfer pipeline analyze only cortical surface, thus the analysis did not include subcortical regions of the amygdala and putamen.

The results of both validations indicated our analysis pipeline could reliably identify regional volumetric estimations. Using CAT12 provided an advantage by performing a longitudinal analysis of subcortical regions, including the amygdala which was pre-hypothesized and of great importance in the current work.

432 Behavioral data

Data collection. To evaluate participants' experience in the peak days of the COVID-19 433 outbreak, we asked them to fill out a 7-items questionnaire regarding their experience of the 434 435 COVID-19 lockdown (see supplementary Table 1 for a description of the items). The questionnaires were filled out soon after the initiation of the study, when the lockdown's 436 stringent 100-meters limitation was lifted. Most participants filled out the questionnaire on the 437 day of the post-lockdown scan session, some filled it a few days before their second scanning 438 session. A total of n = 77 participants filled out the COVID-19 questionnaire and comprised the 439 440 potential pool of test group participants for the current study, out of which n = 50 were sampled 441 and scanned. One participant was scanned but did not complete the questionnaire, therefore this 442 participant's behavioral data were not used and analyses of the questionnaire were based on n =443 49 valid participants.

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Question	Possible Answers (prop.)	Binary outcome (prop.)
1. Did you stay home during the lockdown, except for essential needs / did not leave at all?	0 - no 1 - yes	0 - no (20.4%) 1 - yes (79.6%)
2. Did the lockdown increase your feeling of anxiety?	0 - no 1 - yes	0 - no (61.2%) 1 - yes (39.8%)
3. With how many people did you meet during the lockdown (including people you are living with at home)?	0 - none 1 - up to three people 2- up to five people 3 - up to ten people	0 - up to three (79.6%) 1 - more (20.4%)
4. Do you think your behavior will change following the lockdown?	0 - no 1 - yes	0 - no (65.3%) 1 - yes (34.7%)
5. How did your meeting with your parents' routine look like during the lockdown?	0 - same as before the lockdown1 - with precaution measurements:distancing, mask, etc.2 - did not visit at all	0 - as before or with precautions (44.9%) 1 - did not visit (55.1%)
6. What was your employment status during the lockdown?	 0 - same as before lockdown 1 - full time working from home 2 - part time working from home 3 - Furlough / unemployed 	0 - unemployed / part time (42.9%) 1 - same as before / full time from home (57.1%)
7. How concerned are you with the long-term effect of the lockdown, regarding yourself?	1 - 5 scale	0 - low, score 1-2 (53.1%) 1 - moderate-high, score 3-5 (46.9%)

446 Supplementary Table 1. COVID-19 lockdown questionnaire

447

Data analysis. Responses to the lockdown questionnaire were coded into binary responses, based on the sample median, splitting the sample into relatively similar sized groups for each item. To identify the main themes in the questionnaire which could be correlated with the imaging data, we performed a PCA analysis on the binarized data, using the "factoextra" R package²⁶. We found two principal components, which explained 42.6% of the variance in the sample data. These two components were extracted and correlated with the change in gray matter volumetric data in our regions of interest.

455

Supplementary results

As an exploratory analysis, we examined whether the volumetric brain changes were associated with the psychological constructs identified in our PCA analysis, based on participants' selfreports. We used two linear models aimed to explain the variability in each of the principal components, using the volumetric changes as our model features. Overall neither one of the two PCs were well associated with the volumetric changes (Principal component 1 model: R^2 = 0.205, F(10,38) = 0.98, p = 0.475; Principal component 2 model: $R^2 = 0.33$, F(10,38) = 1.89, p =0.077).

463 While we did find some sporadic ROIs showing significant contribution within the models 464 (measured as the significance of the ROI's β estimates), after FDR correction by the number of 465 features in the model, none of the ROIs demonstrated a significant association with the PCs ($p_{adj.}$ 466 > 0.05).

Finally, we examined correlation patterns of the volumetric change for all brain regions aiming 467 to identify shared change patterns across multiple ROIs. Hierarchical clustering of the correlation 468 469 matrices revealed different patterns between the two groups (Supplementary figure 3). In the test group, three principal groups of clusters could be identified - the first included the palladium, 470 471 hippocampus, amygdala, putamen, insula (all bilateraly), and right anterior cingulate cortex, the 472 second cluster included mostly occipito-temporal cortical and subcortical nuclei, and the third included highly correlated regions of the frontal, parietal and occipital cortices. All regions that 473 474 passed the statistical threshold of the SBM analysis (Table 1), except for the left fusiform gyrus, 475 were grouped closely together within the first two clusters, and showed low to negative correlations the ROIs of the third cluster. This analysis could suggest that the origin of the 476 volumetric change observed in the regions of the two clusters might be different. The regions 477 478 that appear in cluster 1 are often reported in the studies that explore brain changes following stress, anxiety or traumatic events^{5–8}, while the regions of the second cluster are less associated 479 with specific phenomena. 480

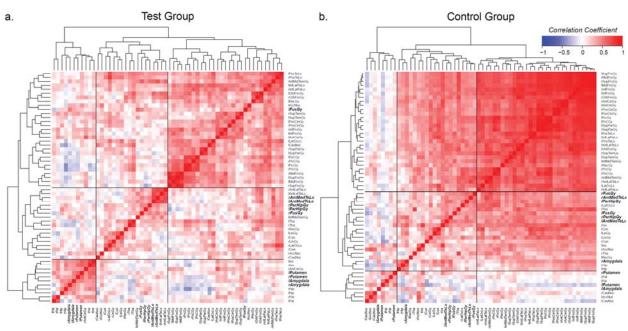
A different pattern was observed in the control group, where the correlation pattern demonstrated stronger volumetric synchrony, (Supplementary figure 3b). This could suggest that changes in the control group were much more affected by within-participant effects, rather than an exogenous effect (which could be the outbreak and lockdown in the test-group). It is important to note in this context that in addition to the experience of the lockdown, the participants in the test group also had longer time gaps between the two scanning sessions, which might provide an

487 alternative explanation for the stronger homogeneity in volumetric changes within the control

488 group.

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490



491 Supplementary figure . Volumetric changes correlation matrices

492 Correlation coefficients were calculated between the volumetric change values for all ROI pairs, 493 analyzed separately for the test (**a**) and the control group (**b**), clustered according to Euclidean 494 distances into dendrograms. In the test group, the first and second cluster contained the 495 subcortical nuclei (amygdala and putamen) and temporal ROI which were found significant in 496 the SBM interaction analysis, respectively (highlighted in italic bold font). In the control group, a 497 more homogeneous change pattern was observed with more robust correlation coefficients 498 between the ROIs. Pearson correlation coefficients are represented by the color scheme.