

Supplementary material of

Associations of brain structure with psychopathy

European Archives of Psychiatry and Clinical Neuroscience

Peter Pieperhoff¹, Lena Hofhansel^{2,3}, Frank Schneider⁴, Jürgen Müller⁶, Katrin Amunts^{1,8}, Sabrina Weber-Papen⁵, Carmen Weidler², Benjamin Clemens², Adrian Raine⁷, Ute Habel^{2,3}

¹ Institute of Neuroscience and Medicine (INM-1), Research Centre Jülich, Jülich, Germany

² Department of Psychiatry, Psychotherapy and Psychosomatics, Medical Faculty, RWTH Aachen University, Aachen, Germany

³ Institute of Neuroscience and Medicine (INM-10), Research Centre Jülich, Jülich, Germany

⁴ Department of History, Philosophy and Ethics of Medicine, School of Medicine, Heinrich-Heine-University Düsseldorf, Germany

⁵ University Hospital Düsseldorf, Heinrich Heine University Düsseldorf, Düsseldorf, Germany

⁶ Department of Psychiatry, Forensic Psychiatry, Human Medical Center, Georg August University Göttingen, Germany

⁷ Departments of Criminology, Psychiatry, and Psychology, University of Pennsylvania, Philadelphia, PA, USA

⁸ C. and O. Vogt Institute for Brain Research, University Hospital Düsseldorf, Heinrich Heine University Düsseldorf, Düsseldorf, Germany

Corresponding author: Peter Pieperhoff (Email: p.pieperhoff@fz-juelich.de)

Methods

Magnetic resonance imaging

The following MR scanners and sequence parameters had been applied in each subsample:

- (1) Data set MU (Müller et al. 2008) was recorded at the University Hospital in Göttingen, Germany with a 1.5 Tesla Siemens Symphony scanner (Siemens Medical Systems, Erlangen, Germany). T1-weighted images were acquired with a magnetization-prepared rapid acquisition gradient echo image (MPRAGE) sequence with a voxel size of $1 \times 1 \times 1.08$ mm (matrix size 256×192 , TR = 11.08 ms, TE = 4 ms, flip angle = 15°).
- (2) Data set (FS, UH) (Schneider et al. 2000) was recorded at the Forschungszentrum Jülich with a 1.5 Tesla Magnetom Vision Scanner (Siemens Medical Systems, Erlangen, Germany). A MPRAGE sequence with a voxel size of $0.9 \times 0.9 \times 1.25$ mm and $0.9 \times 0.9 \times 1.33$ mm was used (matrix size: 256×256 , TR = 704, TE = 4.4 ms).
- (3) Data set (UH, LH) (Hofhansel et al. 2020) was acquired using a 3 Tesla Siemens PRISMA MR scanner (Siemens Medical Systems, Erlangen, Germany), located in the Medical Faculty of RWTH Aachen University. T1-weighted structural images were obtained by means of a three-dimensional MPRAGE sequence (voxel size: $1 \times 1 \times 1$ mm, matrix size: 256×256 , TR = 2300ms, TE = 2.98ms, flip angle = 9°).

Measurement of the intracranial volume (ICV)

The intracranial volume (ICV) of each subject was measured by manual segmentation. First, the MR images were affinely transformed into the space of the reference brain so that the transformed brains had nearly the same extent and alignment. The contour of the intracranial vault was manually delineated on every tenth sagittal section, as well as on the three most lateral sections, using the program ITK-SNAP. The sizes of the segmented mask sections were calculated and linearly interpolated for sections in between. The summed section sizes yielded the ICV in the space of the reference brain, which was rescaled by the determinant of the affine transformation to yield the actual ICV.

Results of region-based analysis

The brain regions are arranged in a hierarchical atlas system with the whole brain on the highest level, and cortical areas and subcortical nuclei at the lowest level. The latter are from the Julich-Brain atlas (Amunts et al. 2020), complemented by maps of nuclei and cerebellar lobules of the AAL3 atlas (Tzourio-Mazoyer et al. 2002, Rolls et al. 2020), and masks of pons, mesencephalon and cerebellar white matter. Regions on upper levels are formed by combination of their child-regions. For every region within the hierarchy the test-statistic t_{uncorr} is calculated by evaluation of the respective statistical model with the region's volume data. The multiple-testing correction follows the procedure described in (Bogomolov et al. 2020).

All regions which have the same *parent region* form a *region family*. The p-value of such a family can be calculated by direct statistical evaluation of the parent region's volume data, which results in the uncorrected p-value p_{uncorr} . Alternatively, this p-value can be calculated by the Simes p-value (Simes 1986) of the family's member regions, see main text for details. Then within each family, the multiple-testing correction of the p-values of its members is calculated using the Benjamini-Hochberg method for controlling the false discovery rate (FDR)(Benjamini and Hochberg 1995), yielding the corrected p-value p_{fdr} . When p_{fdr} is below the specified FDR threshold, the multiple-testing criterion is fulfilled. However, according to (Bogomolov et al. 2020) the FDR threshold q_{set} has to be reduced by the proportion of "significant findings" or "discoveries" in the upper region. This procedure must only be applied within a region family, when its parent region was found to have a significant effect.

The following tables show the results of the atlas branches down to every region, where at least the uncorrected p-value is < 0.05 . The column "sign" indicates columns with significant findings. Significance of multiple-testing corrected p-value: * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$.

u = uncorrected p-value ≤ 0.05 .

Table S1: Associations between PCL-R dimension 1 and regions of hierarchical brain atlas

REGION	t_uncorr	p_simes	p_fdr	sign	q_set
BRAIN	0.79	0.9730	0.9730		0.0500
└─ BRAIN_L	0.78	0.5660	0.9730		0.0500
└─┬─ DIENCEPHALON_L	1.12	0.1420	0.5660		0.0000
└─┬─┬─ THALAMUS_L	1.09	0.4310	0.4310		0.0000
└─┬─┬─┬─ DORSAL THALAMUS_L	1.03	0.2880	0.4310		0.0000
└─┬─┬─┬─┬─ MEDIAL_GROUP_L	0.52	0.0957	0.2880		0.0000
└─┬─┬─┬─┬─ THALAMUS-MV_L	2.24	0.0319	0.0957	u	0.0000
└─┬─ TELECEPHALON_L	1.39	0.5510	0.7210		0.0000
└─┬─┬─ CEREBRAL CORTEX_L	1.14	0.5510	0.5510		0.0000
└─┬─┬─┬─ FRONTAL LOBE_L	0.71	0.5710	0.6850		0.0000
└─┬─┬─┬─┬─ MEDIAL ORBITOFRONTAL CORTEX_L	-0.92	0.0594	0.6240		0.0000
└─┬─┬─┬─┬─┬─ AREA-FO2_L	-2.45	0.0198	0.0594	u	0.0000
└─┬─┬─┬─┬─┬─ SUPERIOR FRONTAL SULCUS_L	1.90	0.1370	0.8870		0.0000
└─┬─┬─┬─┬─┬─ AREA-SFS1_L	2.07	0.0465	0.1370	u	0.0000
└─┬─┬─┬─┬─┬─ VENTRAL PRECENTRAL GYRUS_L	1.44	0.0272	0.5710	u	0.0000
└─┬─┬─┬─┬─┬─ AREA-6V1_L	2.89	0.0068	0.0272	u	0.0000
└─┬─┬─ INSULA_L	0.40	0.0918	0.5510		0.0000
└─┬─┬─┬─ AGRANULAR INSULA_L	1.93	0.0306	0.0918	u	0.0000
└─┬─┬─┬─ AREA-IA3_L	2.73	0.0102	0.0306	u	0.0000
└─┬─┬─ LIMBIC LOBE_L	1.29	0.3670	0.5510		0.0000
└─┬─┬─┬─ HIPPOCAMPAL FORMATION_L	2.50	0.1840	0.3670		0.0000
└─┬─┬─┬─┬─ AREA-EC_L	2.30	0.0283	0.1840	u	0.0000
└─┬─┬─┬─┬─ HIPPOCAMPUS-CA1_L	2.08	0.0459	0.1840	u	0.0000
└─┬─┬─┬─┬─ HIPPOCAMPUS-SUBC_L	1.73	0.1040	0.2360		0.0000
└─┬─┬─┬─┬─┬─ HIPPOCAMPUS-SUBC.PROS_L	2.33	0.0261	0.1040	u	0.0000
└─┬─┬─┬─┬─ PIRIFORM CORTEX_L	1.99	0.1360	0.3670		0.0000
└─┬─┬─┬─┬─┬─ AREA-PIRT_L	2.10	0.0679	0.1360		0.0000
└─┬─┬─┬─┬─┬─ AREA-PIRT.TU_L	2.19	0.0358	0.0679	u	0.0000
└─┬─┬─┬─ TEMPORAL LOBE_L	2.07	0.2220	0.5510		0.0000
└─┬─┬─┬─┬─ GAPMAP-TEMPORAL-TO-PARIETAL_L	2.44	0.0202	0.2220	u	0.0000
└─┬─┬─ CEREBRAL NUCLEI_L	1.56	0.3380	0.5510		0.0000
└─┬─┬─┬─ AMYGDALA_L	1.69	0.4550	0.4550		0.0000
└─┬─┬─┬─┬─ AMYGDALA-CM_L	1.99	0.0852	0.4550		0.0000
└─┬─┬─┬─┬─┬─ AMYGDALA-CM.AAA_L	2.30	0.0284	0.0852	u	0.0000
└─ BRAIN_R	0.77	0.9730	0.9730		0.0500
└─┬─ TELECEPHALON_R	1.14	0.9730	0.9730		0.0000
└─┬─┬─ CEREBRAL CORTEX_R	0.89	0.9730	0.9730		0.0000
└─┬─┬─┬─ FRONTAL LOBE_R	-0.07	0.3280	0.9730		0.0000
└─┬─┬─┬─┬─ MEDIAL ORBITOFRONTAL CORTEX_R	-1.76	0.0156	0.3280	u	0.0000
└─┬─┬─┬─┬─┬─ AREA-FO1_R	-3.00	0.0052	0.0156	u	0.0000
└─┬─┬─┬─┬─┬─ AREA-FO2_R	-2.41	0.0218	0.0327	u	0.0000
└─┬─┬─┬─┬─ MIDDLE FRONTAL GYRUS_R	-0.43	0.1670	0.9370		0.0000
└─┬─┬─┬─┬─┬─ AREA-8V2_R	-2.12	0.0418	0.1670	u	0.0000
└─┬─┬─┬─┬─ SUPERIOR FRONTAL GYRUS_R	-1.27	0.1780	0.9370		0.0000
└─┬─┬─┬─┬─┬─ AREA-8D2_R	-2.09	0.0446	0.1780	u	0.0000
└─┬─┬─┬─┬─ SUPERIOR FRONTAL SULCUS_R	-1.18	0.1270	0.9370		0.0000
└─┬─┬─┬─┬─┬─ AREA-6D3_R	-2.25	0.0317	0.1270	u	0.0000

Table S1: Associations between PCL-R dimension 1 and regions of the hierarchical brain atlas. p-values of higher-level regions where calculated using the Simes method (see above and main text).

Table S2: Associations between PCL-R dimension 2 and regions of hierarchical brain atlas

BRAIN	REGION	t_uncorr	p_simes	p_fdr	sign	q_set
BRAIN		-1.86	0.0240	0.0240	*	0.0500
└─	BRAIN_L	-1.66	0.0136	0.0240	*	0.0500
└─	└─ DIENCEPHALON_L	-2.10	0.0034	0.0136	**	0.0500
└─	└─ └─ SUBTHALAMUS_L	-3.44	0.0017	0.0034	**	0.0375
└─	└─ └─ └─ SUBTHALAMUS-STN_L	-3.44	0.0017	0.0017	**	0.0375
└─	└─ └─ THALAMUS_L	-2.06	0.0135	0.0135	*	0.0375
└─	└─ └─ └─ DORSAL THALAMUS_L	-2.04	0.0090	0.0135	**	0.0375
└─	└─ └─ └─ └─ ANTERIOR GROUP_L	-2.57	0.0018	0.0090	**	0.0250
└─	└─ └─ └─ └─ └─ THALAMUS-AM_L	-3.82	0.0006	0.0018	***	0.0050
└─	└─ └─ └─ └─ └─ THALAMUS-AV_L	-3.25	0.0027	0.0041	**	0.0050
└─	└─ └─ └─ └─ INTRALAMINAR GROUP_L	-2.19	0.0508	0.0635		0.0250
└─	└─ └─ └─ └─ └─ THALAMUS-CL_L	-2.34	0.0254	0.0508	u	0.0050
└─	└─ └─ └─ └─ └─ THALAMUS-SPF_L	-2.46	0.0195	0.0508	u	0.0050
└─	└─ └─ └─ └─ └─ MEDIAL GROUP_L	-2.38	0.0408	0.0635	u	0.0250
└─	└─ └─ └─ └─ └─ THALAMUS-MD_L	-2.17	0.0376	0.0564	u	0.0050
└─	└─ └─ └─ └─ └─ THALAMUS-PV_L	-2.61	0.0136	0.0408	u	0.0050
└─	└─ └─ └─ └─ VENTRAL GROUP_L	-2.40	0.0257	0.0635	u	0.0250
└─	└─ └─ └─ └─ └─ THALAMUS-VAMC_L	-2.96	0.0057	0.0257	u	0.0050
└─	└─ └─ └─ └─ └─ THALAMUS-VA_L	-2.84	0.0077	0.0257	u	0.0050
└─	└─ └─ └─ └─ └─ THALAMUS-VIM_L	-2.16	0.0380	0.0633	u	0.0050
└─	└─ └─ └─ └─ └─ THALAMUS-VLA_L	-2.61	0.0136	0.0340	u	0.0050
└─	└─ └─ └─ └─ └─ THALAMUS-VM_L	-2.84	0.0077	0.0257	u	0.0050
└─	└─ └─ └─ └─ └─ THALAMUS-VPMP_L	-2.23	0.0332	0.0633	u	0.0050
└─	└─ └─ └─ VENTRAL THALAMUS_L	-2.29	0.0050	0.0135	**	0.0375
└─	└─ └─ └─ └─ THALAMUS-ZI_L	-3.28	0.0025	0.0050	**	0.0250
└─	└─ MESENCEPHALON_L	-1.93	0.0357	0.0476	*	0.0500
└─	└─ └─ MIDBRAIN TEGMENTUM_L	-2.48	0.0357	0.0357	*	0.0375
└─	└─ └─ └─ NUCLEUS RUBER PARVOCELL PART_L	-2.49	0.0180	0.0357	*	0.0375
└─	└─ └─ └─ MIDBRAIN-NRP_L	-2.49	0.0180	0.0180	*	0.0250
└─	└─ └─ └─ SUBSTANTIA NIGRA_L	-2.53	0.0238	0.0357	*	0.0375
└─	└─ └─ └─ MIDBRAIN-SNC_L	-2.36	0.0246	0.0246	*	0.0250
└─	└─ └─ └─ MIDBRAIN-SNR_L	-2.67	0.0119	0.0238	*	0.0250
└─	└─ METENCEPHALON_L	-2.26	0.0099	0.0198	**	0.0500
└─	└─ └─ CEREBELLUM_L	-2.11	0.0099	0.0099	**	0.0375
└─	└─ └─ └─ CEREBELLAR CORTEX_L	-2.01	0.0348	0.0522	u	0.0375
└─	└─ └─ └─ └─ CEREBELLUM ANT LOBE_L	-2.97	0.0116	0.0348	u	0.0125
└─	└─ └─ └─ └─ └─ Cerebellum_3_L	-2.68	0.0116	0.0116	u	0.0000
└─	└─ └─ └─ └─ └─ Cerebellum_4_5_L	-2.95	0.0059	0.0116	u	0.0000
└─	└─ └─ └─ └─ CEREBELLUM POST LOBE_L	-1.67	0.1330	0.1330		0.0125
└─	└─ └─ └─ └─ └─ Cerebellum_6_L	-2.05	0.0489	0.1410	u	0.0000
└─	└─ └─ └─ └─ └─ Cerebellum_9_L	-2.47	0.0190	0.1330	u	0.0000
└─	└─ └─ └─ CEREBELLUM VERMIS	-2.69	0.0946	0.1330		0.0125
└─	└─ └─ └─ └─ Vermis_3	-2.14	0.0403	0.0946	u	0.0000
└─	└─ └─ └─ └─ Vermis_4_5	-2.20	0.0348	0.0946	u	0.0000
└─	└─ └─ └─ └─ Vermis_6	-2.55	0.0156	0.0946	u	0.0000
└─	└─ └─ └─ └─ Vermis_8	-2.06	0.0473	0.0946	u	0.0000
└─	└─ └─ CEREBELLAR NUCLEI_L	-2.37	0.0578	0.0578		0.0375
└─	└─ └─ └─ DENTATE NUCLEUS_L	-2.19	0.0454	0.0578	u	0.0125
└─	└─ └─ └─ CEREBELLUM-NDENTD_L	-2.06	0.0476	0.0476	u	0.0000
└─	└─ └─ └─ CEREBELLUM-NDENTV_L	-2.39	0.0227	0.0454	u	0.0000
└─	└─ └─ CEREBELLAR WM_L	-3.18	0.0033	0.0099	**	0.0375
└─	└─ PONS_L	-3.01	0.0062	0.0099	**	0.0375
└─	└─ └─ PONTINE TEGMENTUM_L	-2.31	0.0274	0.0274	*	0.0375
└─	└─ └─ VENTRAL PONS_L	-3.20	0.0031	0.0062	**	0.0375
└─	└─ TELENCEPHALON_L	-0.90	0.2080	0.2080		0.0500
└─	└─ └─ CEREBRAL CORTEX_L	-0.72	0.2080	0.2080		0.0375
└─	└─ └─ └─ FRONTAL LOBE_L	-1.70	0.3680	0.3680		0.0000
└─	└─ └─ └─ └─ DORSAL PRECENTRAL GYRUS_L	-2.42	0.0186	0.3680	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-6D1_L	-2.77	0.0093	0.0186	u	0.0000
└─	└─ └─ └─ └─ LATERAL ORBITOFRONTAL CORTEX_L	-1.10	0.1800	0.5390		0.0000
└─	└─ └─ └─ └─ └─ AREA-FO6_L	-2.09	0.0449	0.1800	u	0.0000
└─	└─ └─ └─ └─ MEDIAL ORBITOFRONTAL CORTEX_L	-1.08	0.0668	0.3680		0.0000
└─	└─ └─ └─ └─ └─ AREA-FO1_L	-2.33	0.0262	0.0668	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-FO2_L	-2.09	0.0445	0.0668	u	0.0000
└─	└─ └─ INSULA_L	-2.32	0.0468	0.2080	u	0.0000
└─	└─ └─ └─ DYSGRANULAR INSULA_L	-2.02	0.1470	0.2210		0.0000
└─	└─ └─ └─ └─ AREA-ID4_L	-2.52	0.0168	0.1470	u	0.0000
└─	└─ └─ └─ └─ AREA-ID5_L	-2.28	0.0295	0.1470	u	0.0000
└─	└─ └─ └─ GRANULAR INSULA_L	-2.89	0.0156	0.0468	u	0.0000
└─	└─ └─ └─ └─ AREA-IG1_L	-2.72	0.0104	0.0156	u	0.0000

	REGION	t_uncorr	p_simes	p_fdr	sign	q_set
	AREA-IG2_L	-2.44	0.0206	0.0206	u	0.0000
	AREA-IG3_L	-2.75	0.0097	0.0156	u	0.0000
	LIMBIC LOBE_L	0.95	0.1100	0.2210		0.0000
	HIPPOCAMPAL FORMATION_L	1.09	0.0552	0.1100		0.0000
	AREA-EC_L	2.89	0.0069	0.0552	u	0.0000
	PIRIFORM CORTEX_L	2.25	0.0384	0.1100	u	0.0000
	AREA-PIRT_L	2.66	0.0192	0.0384	u	0.0000
	AREA-PIRT.TIT_L	2.47	0.0192	0.0192	u	0.0000
	AREA-PIRT.TU_L	2.62	0.0135	0.0192	u	0.0000
	OCCIPITAL LOBE_L	-0.68	0.3590	0.3680		0.0000
	LATERAL OCCIPITAL CORTEX_L	-2.05	0.0897	0.3590		0.0000
	AREA-HOC4LP_L	-2.27	0.0299	0.0897	u	0.0000
	PARIETAL LOBE_L	-0.57	0.3510	0.3680		0.0000
	INFERIOR PARIETAL LOBULE_L	-0.13	0.1330	0.3510		0.0000
	AREA-PFCM_L	-2.47	0.0190	0.1330	u	0.0000
	INTRAPARIETAL SULCUS_L	-2.00	0.1760	0.3510		0.0000
	AREA-HIP3_L	-2.10	0.0439	0.1760	u	0.0000
	AREA-HIP4_L	-2.32	0.0272	0.1760	u	0.0000
	PARIETAL OPERCULUM_L	-0.05	0.0602	0.3510		0.0000
	AREA-OP2_L	-2.27	0.0301	0.0602	u	0.0000
	AREA-OP4_L	2.54	0.0163	0.0602	u	0.0000
	TEMPORAL LOBE_L	1.31	0.0693	0.2080		0.0000
	COLLATERAL SULCUS_L	2.92	0.0063	0.0693	u	0.0000
	AREA-COS1_L	2.92	0.0063	0.0063	u	0.0000
	FUSIFORM GYRUS_L	-0.37	0.1380	0.2280		0.0000
	AREA-FG5_L	2.31	0.0275	0.1380	u	0.0000
	GAPMAP-TEMPORAL-TO-PARIETAL_L	2.05	0.0491	0.2280	u	0.0000
	HESCHL GYRUS_L	-1.95	0.0738	0.2280		0.0000
	AREA-TE-1.1_L	-2.36	0.0246	0.0738	u	0.0000
	SUPERIOR TEMPORAL GYRUS_L	-0.93	0.0981	0.2280		0.0000
	AREA-TE-2.2_L	-2.23	0.0327	0.0981	u	0.0000
	CEREBRAL NUCLEI_L	-1.20	0.1670	0.2080		0.0375
	BASAL GANGLIA_L	-1.09	0.0558	0.1670		0.0000
	Pallidum_L	-2.77	0.0093	0.0558	u	0.0000
	VENTRAL PALLIDUM_L	-2.41	0.0217	0.0651	u	0.0000
	VENTRALPALLIDUM-VP_L	-2.41	0.0217	0.0217	u	0.0000
	BRAIN_R	-1.99	0.0240	0.0240	*	0.0500
	DIENCEPHALON_R	-1.28	0.0136	0.0272	*	0.0500
	SUBTHALAMUS_R	-2.90	0.0068	0.0136	**	0.0500
	SUBTHALAMUS-STN_R	-2.90	0.0068	0.0068	**	0.0250
	THALAMUS_R	-1.27	0.0660	0.0660		0.0500
	DORSAL THALAMUS_R	-1.25	0.0450	0.0675	u	0.0250
	ANTERIOR GROUP_R	-3.04	0.0117	0.0450	u	0.0000
	THALAMUS-AM_R	-3.11	0.0039	0.0117	u	0.0000
	THALAMUS-AV_R	-2.84	0.0078	0.0117	u	0.0000
	MEDIAL GROUP_R	-0.98	0.0180	0.0450	u	0.0000
	THALAMUS-PV_R	-2.94	0.0060	0.0180	u	0.0000
	VENTRAL GROUP_R	-1.90	0.0395	0.0658	u	0.0000
	THALAMUS-VAMC_R	-2.83	0.0079	0.0395	u	0.0000
	THALAMUS-VA_R	-3.10	0.0040	0.0395	u	0.0000
	THALAMUS-VIA_R	-2.50	0.0179	0.0447	u	0.0000
	THALAMUS-VM_R	-2.51	0.0173	0.0447	u	0.0000
	VENTRAL THALAMUS_R	-1.34	0.0220	0.0660	u	0.0250
	THALAMUS-ZI_R	-2.70	0.0110	0.0220	u	0.0000
	MESENCEPHALON_R	-2.14	0.0303	0.0404	*	0.0500
	MIDBRAIN TEGMENTUM_R	-2.64	0.0303	0.0303	*	0.0500
	NUCLEUS RUBER PARVOCELL PART_R	-2.45	0.0202	0.0303	*	0.0500
	MIDBRAIN-NRP_R	-2.45	0.0202	0.0202	*	0.0333
	SUBSTANTIA NIGRA_R	-2.67	0.0144	0.0303	*	0.0500
	MIDBRAIN-SNC_R	-2.43	0.0207	0.0207	*	0.0333
	MIDBRAIN-SNR_R	-2.87	0.0072	0.0144	**	0.0333
	METENCEPHALON_R	-2.80	0.0060	0.0240	**	0.0500
	CEREBELLUM_R	-2.63	0.0039	0.0060	**	0.0500
	CEREBELLAR CORTEX_R	-2.46	0.0792	0.0792		0.0500
	CEREBELLUM ANT LOBE_R	-2.37	0.0396	0.0792	u	0.0167
	Cerebellum 4_5_R	-2.45	0.0198	0.0396	u	0.0000
	CEREBELLUM POST LOBE_R	-2.30	0.0805	0.0805		0.0167
	Cerebellum 6_R	-2.66	0.0121	0.0805	u	0.0000
	Cerebellum 9_R	-2.29	0.0288	0.0805	u	0.0000
	Cerebellum Crus1_R	-2.21	0.0345	0.0805	u	0.0000
	CEREBELLAR NUCLEI_R	-2.64	0.0437	0.0655	u	0.0500
	CEREBELLUM-NINTERP_R	-2.10	0.0437	0.0437	u	0.0167
	DENTATE NUCLEUS_R	-2.45	0.0292	0.0437	u	0.0167
	CEREBELLUM-NDENTD_R	-2.24	0.0322	0.0322	u	0.0000
	CEREBELLUM-NDENTV_R	-2.58	0.0146	0.0292	u	0.0000

REGION	t_uncorr	p_simes	p_fdr	sign	q_set
FASTIGIAL_NUCLEUS_R	-2.10	0.0433	0.0437	u	0.0167
CEREBELLUM-NFAST_R	-2.10	0.0433	0.0433	u	0.0000
CEREBELLAR_WM_R	-3.52	0.0013	0.0039	**	0.0500
PONS_R	-3.08	0.0060	0.0060	**	0.0500
PONTINE_TEGMENTUM_R	-2.51	0.0175	0.0175	*	0.0500
VENTRAL_PONS_R	-3.21	0.0030	0.0060	**	0.0500
TELENCEPHALON_R	-1.14	0.0450	0.0450	*	0.0500
CEREBRAL_CORTEX_R	-0.81	0.7590	0.7590		0.0500
FRONTAL_LOBE_R	-1.73	0.3890	0.7590		0.0250
FRONTAL_OPERCULUM_R	-1.25	0.2630	0.8190		0.0000
AREA-OP5_R	-2.10	0.0438	0.2630	u	0.0000
MEDIAL_ORBITOFRONTAL_CORTEX_R	-1.86	0.0201	0.3890	u	0.0000
AREA-FO1_R	-2.90	0.0067	0.0201	u	0.0000
AREA-FO2_R	-2.05	0.0483	0.0725	u	0.0000
SUPERIOR_FRONTAL_SULCUS_R	-2.90	0.0522	0.3890		0.0000
AREA-SFG3_R	-2.35	0.0253	0.0522	u	0.0000
AREA-SFS1_R	-2.33	0.0261	0.0522	u	0.0000
INSULA_R	-1.88	0.2880	0.7590		0.0250
DYSGRANULAR_INSULA_R	-1.77	0.0960	0.2880		0.0000
AREA-ID4_R	-2.76	0.0096	0.0960	u	0.0000
OCCIPITAL_LOBE_R	-1.24	0.5060	0.7590		0.0250
LATERAL_OCCIPITAL_CORTEX_R	-1.56	0.1500	0.5060		0.0000
AREA-HOC5_R	-2.04	0.0499	0.1500	u	0.0000
TEMPORAL_LOBE_R	1.37	0.3870	0.7590		0.0250
COLLATERAL_SULCUS_R	2.20	0.0352	0.3870	u	0.0000
AREA-COS1_R	2.20	0.0352	0.0352	u	0.0000
FUSIFORM_GYRUS_R	0.72	0.1260	0.4140		0.0000
AREA-FG5_R	2.35	0.0252	0.1260	u	0.0000
CEREBRAL_NUCLEI_R	-2.13	0.0225	0.0450	*	0.0500
AMYGDALA_R	-0.45	0.5020	0.5020		0.0250
AMYGDALA-MF_R	-1.90	0.0864	0.5020		0.0083
AMYGDALA-MF.LM_R	-2.11	0.0432	0.0864	u	0.0000
AMYGDALA-SF_R	-0.50	0.1430	0.5020		0.0083
AMYGDALA-SF.AHI_R	-2.06	0.0478	0.1430	u	0.0000
BASAL_FOREBRAIN_R	-1.85	0.0237	0.0356	u	0.0250
SUBLENTICULAR_BASAL_FOREBRAIN_R	-2.83	0.0079	0.0237	u	0.0083
BASALFOREBRAIN-CH-4_R	-2.83	0.0079	0.0079	u	0.0000
BASAL_GANGLIA_R	-2.03	0.0075	0.0225	**	0.0250
N_Acc_R	-2.22	0.0333	0.0500	u	0.0083
Pallidum_R	-3.28	0.0025	0.0075	**	0.0083
VENTRAL_PALLIDUM_R	-3.34	0.0022	0.0075	**	0.0083
VENTRALPALLIDUM-VP_R	-3.34	0.0022	0.0022	**	0.0028
VENTRAL_STRIATUM_R	-2.95	0.0082	0.0164	u	0.0083
VENTRALSTRIATUM-ACBL_R	-3.19	0.0032	0.0082	u	0.0028
VENTRALSTRIATUM-FUCD_R	-2.22	0.0339	0.0452	u	0.0028
VENTRALSTRIATUM-FUP_R	-3.10	0.0041	0.0082	u	0.0028

Table S2: Associations between PCL-R dimension 2 and regions of the hierarchical brain atlas. p-values of higher-level regions were calculated using the Simes method (see above and main text).

Table S3: Associations between PCL-R total and regions of hierarchical brain atlas

	REGION	t_uncorr	p_simes	p_fdr	sign	q_set
BRAIN		-0.84	0.6000	0.6000		0.0500
└─	BRAIN_L	-0.72	0.6000	0.6000		0.0500
└─	└─ METENCEPHALON_L	-2.18	0.1500	0.6000		0.0000
└─	└─ └─ CEREBELLUM_L	-2.17	0.1110	0.1500		0.0000
└─	└─ └─ └─ CEREBELLAR CORTEX_L	-2.16	0.1820	0.2230		0.0000
└─	└─ └─ └─ └─ CEREBELLUM POST LOBE_L	-2.06	0.1410	0.1820		0.0000
└─	└─ └─ └─ └─ └─ Cerebellum_8_L	-2.23	0.0329	0.1410	u	0.0000
└─	└─ └─ └─ └─ └─ CEREBELLUM VERMIS	-1.98	0.1820	0.1820		0.0000
└─	└─ └─ └─ └─ └─ └─ Vermis_8	-2.39	0.0228	0.1820	u	0.0000
└─	└─ └─ └─ └─ └─ CEREBELLAR WM_L	-2.17	0.0369	0.1110	u	0.0000
└─	└─ TELECEPHALON_L	0.02	0.6010	0.8020		0.0000
└─	└─ └─ CEREBRAL CORTEX_L	-0.02	0.3010	0.6010		0.0000
└─	└─ └─ └─ FRONTAL LOBE_L	-0.78	0.1700	0.3010		0.0000
└─	└─ └─ └─ └─ MEDIAL ORBITOFRONTAL CORTEX_L	-1.55	0.0081	0.1700	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-FO1_L	-2.89	0.0067	0.0100	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-FO2_L	-3.25	0.0027	0.0081	u	0.0000
└─	└─ └─ └─ LIMBIC LOBE_L	1.57	0.0544	0.3010		0.0000
└─	└─ └─ └─ └─ HIPPOCAMPAL FORMATION_L	2.28	0.0272	0.0544	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-EC_L	3.16	0.0034	0.0272	u	0.0000
└─	└─ └─ └─ └─ └─ PIRIFORM CORTEX_L	2.40	0.0152	0.0544	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-PIRT_L	2.71	0.0076	0.0152	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-PIRT.TIT_L	2.37	0.0240	0.0240	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-PIRT.TU_L	3.11	0.0038	0.0076	u	0.0000
└─	└─ └─ └─ PARIETAL LOBE_L	-0.14	0.2000	0.3010		0.0000
└─	└─ └─ └─ └─ INFERIOR PARIETAL LOBULE_L	-0.13	0.2450	0.5440		0.0000
└─	└─ └─ └─ └─ └─ AREA-PFCM_L	-2.20	0.0350	0.2450	u	0.0000
└─	└─ └─ └─ └─ └─ PARIETO-OCCIPITAL SULCUS_L	2.22	0.0334	0.2000	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-HPO1_L	2.22	0.0334	0.0334	u	0.0000
└─	└─ └─ └─ TEMPORAL LOBE_L	1.67	0.1280	0.3010		0.0000
└─	└─ └─ └─ └─ COLLATERAL SULCUS_L	2.23	0.0324	0.1280	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-COS1_L	2.23	0.0324	0.0324	u	0.0000
└─	└─ └─ └─ └─ GAPMAP-TEMPORAL-TO-PARIETAL_L	2.45	0.0196	0.1280	u	0.0000
└─	└─ └─ └─ └─ SUPERIOR TEMPORAL GYRUS_L	-1.19	0.0348	0.1280	u	0.0000
└─	└─ └─ └─ └─ AREA-TE-2.2_L	-2.67	0.0116	0.0348	u	0.0000
└─	BRAIN_R	-0.95	0.5290	0.6000		0.0500
└─	└─ METENCEPHALON_R	-2.23	0.1320	0.5290		0.0000
└─	└─ └─ CEREBELLUM_R	-2.19	0.1320	0.1320		0.0000
└─	└─ └─ └─ CEREBELLAR CORTEX_R	-2.16	0.0882	0.1320		0.0000
└─	└─ └─ └─ └─ CEREBELLUM POST LOBE_R	-2.15	0.0441	0.0882	u	0.0000
└─	└─ └─ └─ └─ └─ Cerebellum_6_R	-2.64	0.0126	0.0441	u	0.0000
└─	└─ └─ └─ └─ └─ Cerebellum_Crus1_R	-2.87	0.0070	0.0441	u	0.0000
└─	└─ TELECEPHALON_R	-0.30	0.3020	0.6050		0.0000
└─	└─ └─ CEREBRAL CORTEX_R	-0.28	0.1510	0.3020		0.0000
└─	└─ └─ └─ FRONTAL LOBE_R	-1.45	0.0252	0.1510	u	0.0000
└─	└─ └─ └─ └─ FRONTAL OPERCULUM_R	-1.32	0.1210	0.6130		0.0000
└─	└─ └─ └─ └─ └─ AREA-OP5_R	-2.44	0.0202	0.1210	u	0.0000
└─	└─ └─ └─ └─ └─ MEDIAL ORBITOFRONTAL CORTEX_R	-2.31	0.0012	0.0252	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-FO1_R	-3.92	0.0004	0.0012	u	0.0000
└─	└─ └─ └─ └─ └─ AREA-FO2_R	-3.24	0.0028	0.0042	u	0.0000
└─	└─ └─ └─ └─ POSTERIOR SFG_R	-2.09	0.0447	0.3130	u	0.0000
└─	└─ └─ └─ └─ AREA-6MA_R	-2.09	0.0447	0.0447	u	0.0000
└─	└─ └─ └─ └─ SUPERIOR FRONTAL GYRUS_R	-2.53	0.0124	0.1300	u	0.0000
└─	└─ └─ └─ └─ AREA-8D2_R	-3.19	0.0031	0.0124	u	0.0000
└─	└─ └─ └─ └─ SUPERIOR FRONTAL SULCUS_R	-2.74	0.1460	0.6130		0.0000
└─	└─ └─ └─ └─ AREA-SFG3_R	-2.18	0.0365	0.1460	u	0.0000
└─	└─ └─ INSULA_R	-1.27	0.4380	0.8780		0.0000
└─	└─ └─ └─ DYSGRANULAR INSULA_R	-1.34	0.1460	0.4380		0.0000
└─	└─ └─ └─ AREA-ID4_R	-2.58	0.0146	0.1460	u	0.0000
└─	└─ └─ LIMBIC LOBE_R	0.61	0.6050	0.8780		0.0000
└─	└─ └─ └─ HIPPOCAMPAL FORMATION_R	2.07	0.2360	0.6050		0.0000
└─	└─ └─ └─ └─ HIPPOCAMPUS-CA1_R	2.28	0.0295	0.2360	u	0.0000
└─	└─ CEREBRAL NUCLEI_R	-0.60	0.7610	0.7610		0.0000
└─	└─ └─ BASAL GANGLIA_R	-0.60	0.4840	0.7610		0.0000
└─	└─ └─ └─ VENTRAL STRIATUM_R	-1.51	0.1640	0.4910		0.0000
└─	└─ └─ └─ └─ VENTRALSTRIATUM-ACBL_R	-2.13	0.0409	0.1640	u	0.0000

Table S3: Associations between PCL-R total score and regions of the hierarchical brain atlas. p-values of higher-level regions where calculated using the Simes method (see above and main text).

Figure S4: Associations between PCL-R total and brain regions

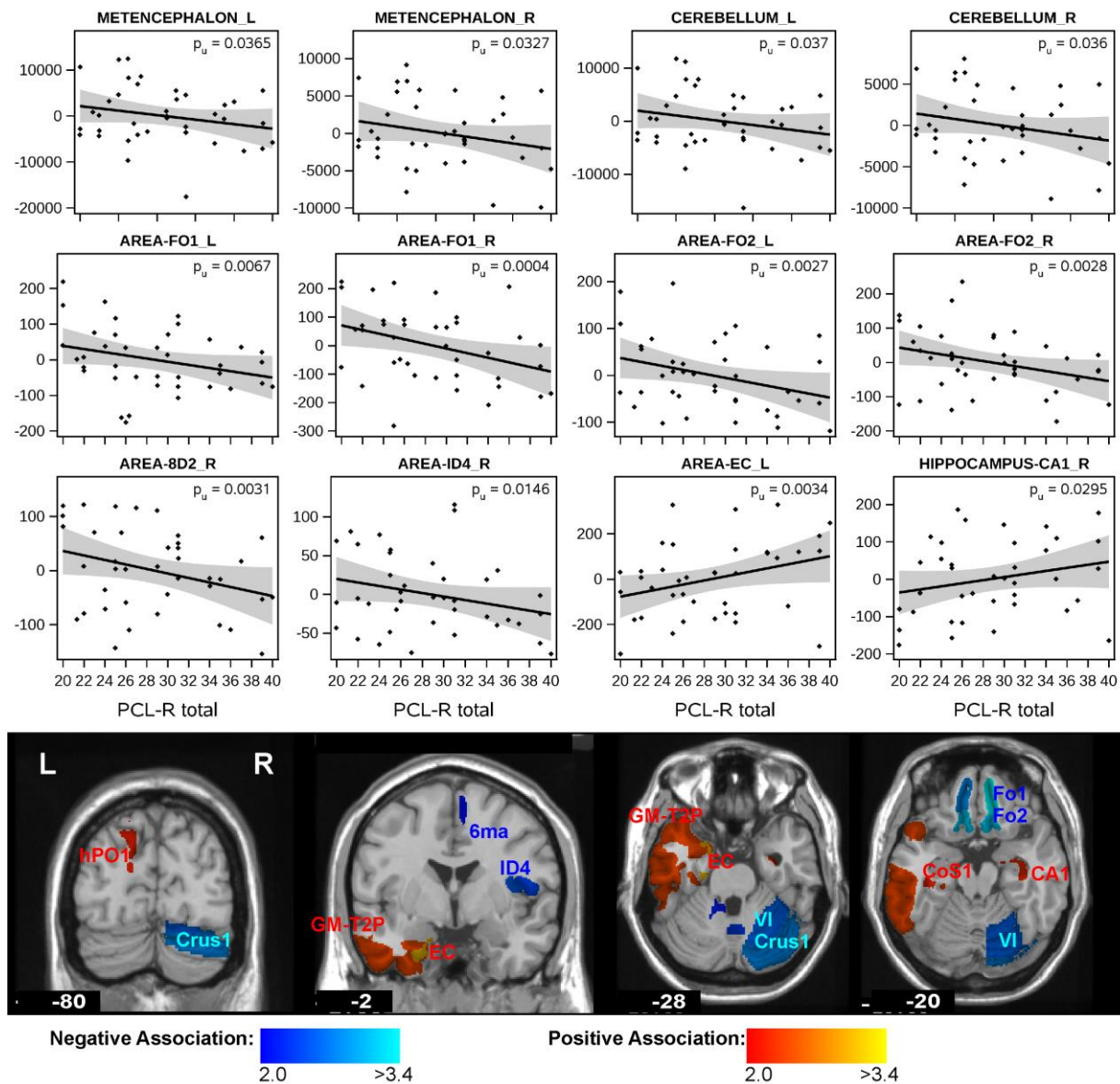


Figure S4: Associations of PCL-R total with volumes of brain regions in psychopathic subjects with PCL-R total ≥ 20 . Volume data were adjusted for study (acquisition site), age and intra-cranial volume. The scatter-plots show the individual brain data points, the regression line with its confidence limits and the p-value (uncorr.). The bottom row shows the brain regions, where an association with $p_{\text{uncorr}} < 0.05$ had been found, with colors corresponding to the statistical score (student's t). Blue-cyan: Negative association, red-yellow: positive association.

Table S5: Group differences between psychopathic subjects and controls in regions of the hierarchical atlas

REGION		t_uncorr	p_uncorr	p_fdr	sign	q_set
BRAIN		-2.96	0.0042	0.0042	**	0.0500
└─ BRAIN_L		-2.95	0.0043	0.0061	**	0.0500
└─┬─ METENCEPHALON_L		-1.87	0.0662	0.1320		0.0500
└─┬─┬─ CEREBELLUM_L		-1.96	0.0542	0.1080		0.0000
└─┬─┬─┬─ CEREBELLAR CORTEX_L		-1.96	0.0537	0.0805		0.0000
└─┬─┬─┬─┬─ CEREBELLUM POST LOBE_L		-2.20	0.0310	0.0930	u	0.0000
└─┬─┬─┬─┬─┬─ Cerebellum_6_L		-2.20	0.0310	0.0933	u	0.0000
└─┬─┬─┬─┬─ CEREBELLUM VERMIS		-0.38	0.7040	0.8320		0.0000
└─┬─┬─┬─┬─┬─ Vermis_10		-2.43	0.0176	0.1410	u	0.0000
└─┬─┬─┬─┬─ CEREBELLAR NUCLEI_L		-2.07	0.0423	0.0805	u	0.0000
└─┬─┬─┬─┬─┬─ DENTATE NUCLEUS_L		-1.90	0.0612	0.1840		0.0000
└─┬─┬─┬─┬─┬─ CEREBELLUM-NDENTD_L		-2.05	0.0436	0.0872	u	0.0000
└─┬─ TELENCEPHALON_L		-2.46	0.0162	0.0648	u	0.0500
└─┬─┬─ CEREBRAL CORTEX_L		-2.83	0.0061	0.0122	u	0.0000
└─┬─┬─┬─ FRONTAL LOBE_L		-2.61	0.0111	0.0666	u	0.0000
└─┬─┬─┬─┬─ GAPMAP-FRONTAL-TO-OCCIPITAL_L		-2.51	0.0142	0.2980	u	0.0000
└─┬─┬─┬─┬─ MIDDLE FRONTAL GYRUS_L		-1.78	0.0789	0.3260		0.0000
└─┬─┬─┬─┬─┬─ AREA-8V2_L		-2.15	0.0353	0.1410	u	0.0000
└─┬─┬─┬─┬─┬─ SUPERIOR FRONTAL GYRUS_L		-1.88	0.0636	0.3260		0.0000
└─┬─┬─┬─┬─┬─┬─ AREA-SFG2_L		-2.20	0.0309	0.0724	u	0.0000
└─┬─┬─┬─┬─┬─┬─ AREA-SFG4_L		-2.13	0.0362	0.0724	u	0.0000
└─┬─┬─┬─┬─┬─ SUPERIOR FRONTAL SULCUS_L		-1.62	0.1090	0.3260		0.0000
└─┬─┬─┬─┬─┬─┬─ AREA-6D3_L		-2.31	0.0236	0.0944	u	0.0000
└─┬─┬─┬─ LIMBIC LOBE_L		-1.95	0.0548	0.1110		0.0000
└─┬─┬─┬─┬─ HIPPOCAMPAL FORMATION_L		-1.78	0.0796	0.3180		0.0000
└─┬─┬─┬─┬─┬─ HIPPOCAMPUS-SUBC_L		-1.97	0.0531	0.1890		0.0000
└─┬─┬─┬─┬─┬─┬─ HIPPOCAMPUS-SUBC.PRES_L		-2.10	0.0390	0.1160	u	0.0000
└─┬─┬─ OCCIPITAL LOBE_L		-0.96	0.3400	0.3400		0.0000
└─┬─┬─┬─ DORSAL OCCIPITAL CORTEX_L		-2.06	0.0431	0.1720	u	0.0000
└─┬─┬─┬─┬─ AREA-HOC4D_L		-2.44	0.0173	0.0516	u	0.0000
└─┬─┬─┬─┬─ AREA-HOC6_L		-2.16	0.0344	0.0516	u	0.0000
└─┬─┬─ PARIETAL LOBE_L		-1.56	0.1220	0.1840		0.0000
└─┬─┬─┬─ INFERIOR PARIETAL LOBULE_L		-0.95	0.3460	0.4800		0.0000
└─┬─┬─┬─┬─ AREA-PGP_L		-2.16	0.0339	0.2370	u	0.0000
└─┬─┬─┬─┬─ INTRAPARIETAL SULCUS_L		-2.79	0.0067	0.0402	u	0.0000
└─┬─┬─┬─┬─┬─ AREA-HIP1_L		-3.89	0.0002	0.0016	u	0.0000
└─┬─┬─┬─┬─┬─ AREA-HIP2_L		-2.70	0.0087	0.0232	u	0.0000
└─┬─┬─┬─┬─┬─ AREA-HIP3_L		-2.75	0.0075	0.0232	u	0.0000
└─┬─┬─┬─┬─ PARIETAL OPERCULUM_L		1.06	0.2930	0.4800		0.0000
└─┬─┬─┬─┬─┬─ AREA-OP1_L		2.44	0.0174	0.0696	u	0.0000
└─┬─┬─┬─ TEMPORAL LOBE_L		-1.95	0.0555	0.1110		0.0000
└─┬─┬─┬─┬─ GAPMAP-FRONTAL-TO-TEMPORAL-II_L		-2.04	0.0450	0.1240	u	0.0000
└─┬─┬─┬─┬─┬─ OCCIPITO-TEMPORAL SULCUS_L		2.57	0.0122	0.1240	u	0.0000
└─┬─┬─┬─┬─┬─┬─ AREA-OTS1_L		2.57	0.0122	0.0122	u	0.0000
└─┬─┬─┬─┬─┬─ PARAHIPPOCAMPAL GYRUS_L		-2.31	0.0235	0.1240	u	0.0000
└─┬─┬─┬─┬─┬─┬─ AREA-PH2_L		-2.00	0.0494	0.0741	u	0.0000
└─┬─┬─┬─┬─┬─┬─ AREA-PH3_L		-2.49	0.0152	0.0456	u	0.0000
└─┬─┬─┬─┬─┬─ SUPERIOR TEMPORAL SULCUS_L		-2.16	0.0338	0.1240	u	0.0000
└─┬─┬─┬─┬─┬─┬─ AREA-STS2_L		-2.06	0.0435	0.0809	u	0.0000
└─┬─┬─ CEREBRAL NUCLEI_L		-0.53	0.6000	0.6000		0.0000
└─┬─┬─┬─ AMYGDALA_L		-1.53	0.1300	0.3890		0.0000
└─┬─┬─┬─┬─ AMYGDALA-CM_L		-1.52	0.1340	0.2350		0.0000
└─┬─┬─┬─┬─┬─ AMYGDALA-CM.ME_L		-2.06	0.0426	0.1280	u	0.0000
└─┬─┬─┬─┬─┬─ AMYGDALA-LB_L		-1.29	0.2010	0.2820		0.0000
└─┬─┬─┬─┬─┬─┬─ AMYGDALA-LB.BM_L		-2.29	0.0251	0.1000	u	0.0000
└─┬─┬─┬─┬─┬─ AMYGDALA-MF_L		-2.07	0.0418	0.2350	u	0.0000
└─┬─┬─┬─┬─┬─┬─ AMYGDALA-MF.ICM_L		-2.22	0.0296	0.0592	u	0.0000
└─ BRAIN_R		-2.83	0.0061	0.0061	**	0.0500
└─┬─ DIENCEPHALON_R		-0.58	0.5630	0.5630		0.0500
└─┬─┬─ THALAMUS_R		-0.59	0.5540	0.9080		0.0125
└─┬─┬─┬─ METATHALAMUS_R		-1.68	0.0969	0.2910		0.0000
└─┬─┬─┬─┬─ LATERAL GENICULATE BODY_R		-2.14	0.0356	0.0712	u	0.0000
└─┬─┬─┬─┬─┬─ METATHALAMUS-CGL_R		-2.14	0.0356	0.0356	u	0.0000
└─┬─ TELENCEPHALON_R		-2.65	0.0098	0.0392	**	0.0500
└─┬─┬─ CEREBRAL CORTEX_R		-3.10	0.0028	0.0056	**	0.0125
└─┬─┬─┬─ FRONTAL LOBE_R		-1.75	0.0843	0.1010		0.0063
└─┬─┬─┬─┬─ DORSAL PRECENTRAL GYRUS_R		-1.89	0.0626	0.3290		0.0000
└─┬─┬─┬─┬─┬─ AREA-6D2_R		-2.00	0.0493	0.0986	u	0.0000

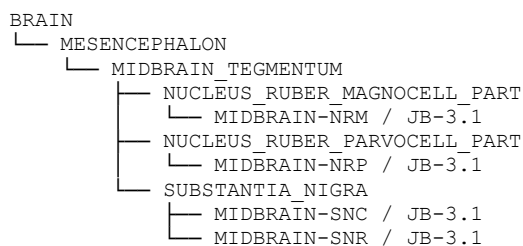
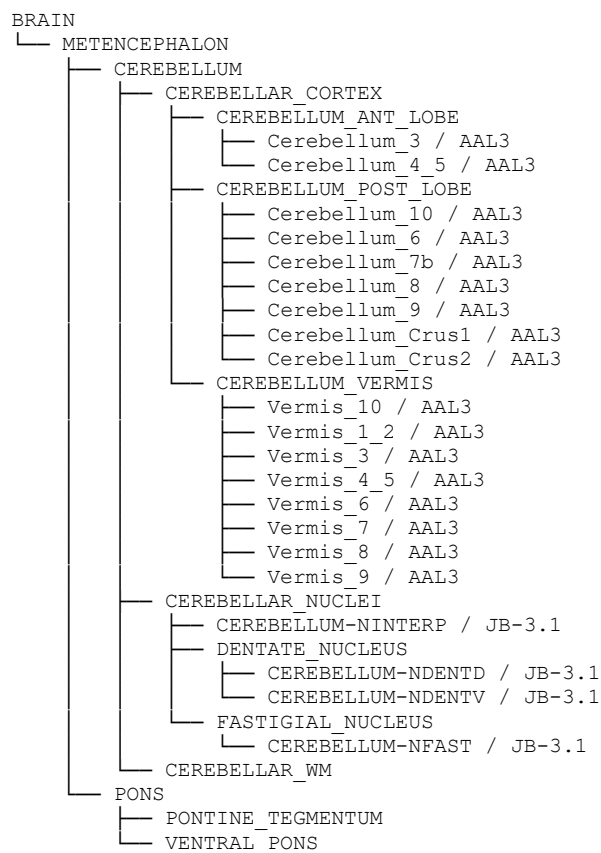
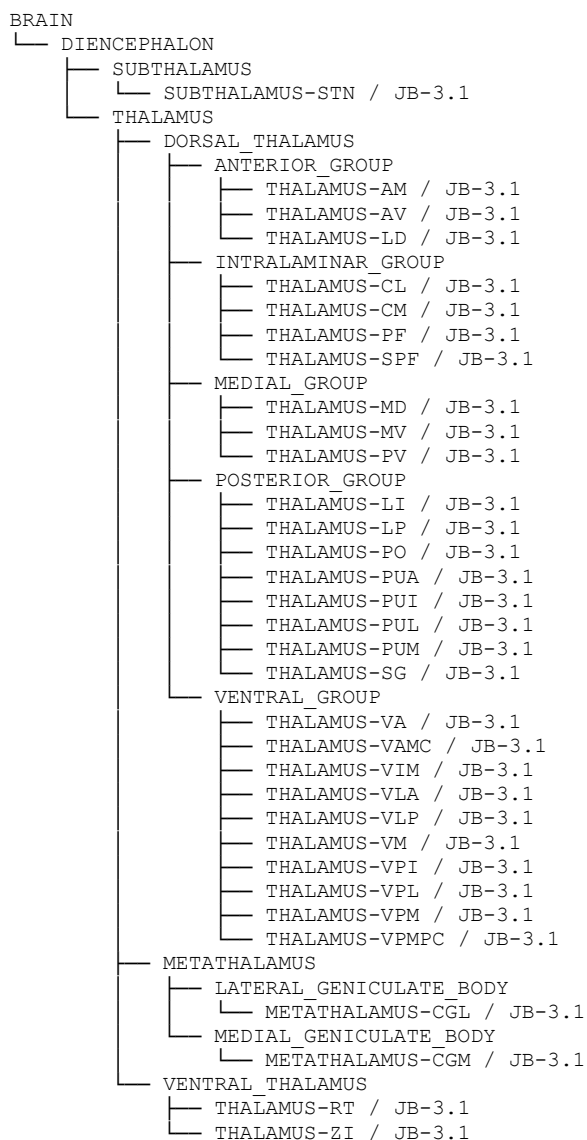
REGION	t_ uncorr	p_ uncorr	p_ fdr	sign	q_ set
└ GAPMAP-FRONTAL-TO-OCCIPITAL_R	-2.89	0.0051	0.0672	u	0.0000
└ INFERIOR FRONTAL SULCUS_R	-1.47	0.1460	0.4700		0.0000
└ AREA-IFS1_R	-2.03	0.0458	0.2160	u	0.0000
└ MESIAL PRECENTRAL GYRUS_R	-2.81	0.0064	0.0672	u	0.0000
└ AREA-6MP_R	-2.81	0.0064	0.0064	u	0.0000
└ MIDDLE FRONTAL GYRUS_R	-1.16	0.2490	0.5820		0.0000
└ AREA-MFG4_R	-2.32	0.0230	0.0920	u	0.0000
└ POSTERIOR SFG_R	-2.58	0.0121	0.0847	u	0.0000
└ AREA-6MA_R	-2.58	0.0121	0.0121	u	0.0000
└ SUPERIOR FRONTAL SULCUS_R	-1.43	0.1570	0.4700		0.0000
└ AREA-6D3_R	-2.04	0.0454	0.1820	u	0.0000
└ INSULA_R	-2.54	0.0132	0.0396	u	0.0063
└ DYSGRANULAR INSULA_R	-2.46	0.0161	0.0483	u	0.0000
└ AREA-ID4_R	-2.17	0.0331	0.1650	u	0.0000
└ AREA-ID9_R	-2.20	0.0309	0.1650	u	0.0000
└ LIMBIC LOBE_R	-2.91	0.0048	0.0288	u	0.0063
└ FRONTAL CINGULATE GYRUS_R	-1.33	0.1870	0.2490		0.0000
└ AREA-33_R	-2.14	0.0355	0.2260	u	0.0000
└ HIPPOCAMPAL FORMATION_R	-3.17	0.0022	0.0088	u	0.0000
└ HIPPOCAMPUS-CA1_R	-2.17	0.0332	0.1330	u	0.0000
└ HIPPOCAMPUS-SUBC_R	-4.37	0.0000	0.0000	u	0.0000
└ HIPPOCAMPUS-SUBC.PAS_R	-2.60	0.0112	0.0112	u	0.0000
└ HIPPOCAMPUS-SUBC.PRES_R	-3.37	0.0012	0.0016	u	0.0000
└ HIPPOCAMPUS-SUBC.PROS_R	-4.09	0.0001	0.0002	u	0.0000
└ HIPPOCAMPUS-SUBC.SUB_R	-4.14	0.0001	0.0002	u	0.0000
└ PIRIFORM CORTEX_R	-1.60	0.1130	0.2260		0.0000
└ AREA-PIRTB_R	-2.28	0.0255	0.0510	u	0.0000
└ AREA-PIRTB.TBD_R	-2.26	0.0272	0.0369	u	0.0000
└ AREA-PIRTB.TBV_R	-2.13	0.0369	0.0369	u	0.0000
└ PARIETAL LOBE_R	-2.24	0.0280	0.0466	u	0.0063
└ INFERIOR PARIETAL LOBULE_R	-1.54	0.1270	0.2540		0.0000
└ AREA-PFCM_R	-2.12	0.0375	0.2630	u	0.0000
└ INTRAPARIETAL SULCUS_R	-1.94	0.0564	0.2540		0.0000
└ AREA-HIP3_R	-2.07	0.0424	0.3390	u	0.0000
└ SUPERIOR PARIETAL LOBULE_R	-1.55	0.1260	0.2540		0.0000
└ AREA-7PC_R	-2.64	0.0103	0.0721	u	0.0000
└ TEMPORAL LOBE_R	-2.20	0.0311	0.0466	u	0.0063
└ GAPMAP-FRONTAL-TO-TEMPORAL-II_R	-2.74	0.0077	0.0847	u	0.0000
└ TEMPORO-PARIETAL JUNCTION_R	-2.31	0.0240	0.1320	u	0.0000
└ AREA-TPJ_R	-2.31	0.0240	0.0240	u	0.0000
└ CEREBRAL NUCLEI_R	-0.74	0.4600	0.4600		0.0125
└ AMYGDALA_R	-2.24	0.0281	0.0843	u	0.0063
└ AMYGDALA-CM_R	-1.99	0.0499	0.1160	u	0.0000
└ AMYGDALA-CM.CE_R	-2.08	0.0413	0.1240	u	0.0000
└ AMYGDALA-IF_R	-1.43	0.1570	0.1830		0.0000
└ AMYGDALA-IF.LD_R	-2.13	0.0370	0.1010	u	0.0000
└ AMYGDALA-LB_R	-1.86	0.0668	0.1170		0.0000
└ AMYGDALA-LB.BL_R	-2.25	0.0276	0.1100	u	0.0000
└ AMYGDALA-VTM_R	-2.39	0.0196	0.1160	u	0.0000
└ AMYGDALOSTRIATAL TZ_R	-2.03	0.0459	0.1160	u	0.0000
└ AMYGDALA-ASTR_R	-2.03	0.0459	0.0459	u	0.0000

Table S5: Group differences between psychopathic subjects and controls in regions of the hierarchical brain atlas. p-values of all regions were calculated by direct evaluation of the statistical model with each region's volume data (see above and main text). The test statistic t_{uncorr} belongs to the comparison $V_{\text{PS}} - V_{\text{C}}$, thus a negative value indicates that the region volume was smaller in the brains of psychopathic subjects than in controls' brains.

Table S6: Hierarchical anatomical atlas

JB-3.1: Julich-Brain Atlas 3.1 (Amunts et al. 2020), <https://www.ebrains.eu/tools/human-brain-atlas>

AAL3: AAL3-Atlas (Tzourio-Mazoyer et al. 2002, Rolls et al. 2020)



```

BRAIN
├── TELECEPHALON
│   ├── CEREBRAL_CORTEX
│   │   ├── FRONTAL LOBE
│   │   │   ├── DORSAL_PRECENTRAL_GYRUS
│   │   │   │   ├── AREA-6D1 / JB-3.1
│   │   │   │   └── AREA-6D2 / JB-3.1
│   │   │   ├── FRONTAL OPERCULUM
│   │   │   │   ├── AREA-OP10 / JB-3.1
│   │   │   │   ├── AREA-OP5 / JB-3.1
│   │   │   │   ├── AREA-OP6 / JB-3.1
│   │   │   │   ├── AREA-OP7 / JB-3.1
│   │   │   │   ├── AREA-OP8 / JB-3.1
│   │   │   │   └── AREA-OP9 / JB-3.1
│   │   │   ├── FRONTAL_POLE
│   │   │   │   ├── AREA-FP1 / JB-3.1
│   │   │   │   └── AREA-FP2 / JB-3.1
│   │   │   ├── FRONTO-MARGINAL_SULCUS
│   │   │   │   └── AREA-MFG2 / JB-3.1
│   │   │   ├── GAPM-FRONTAL-I.1 / JB-3.1
│   │   │   ├── GAPM-FRONTAL-I.2 / JB-3.1
│   │   │   ├── GAPM-FRONTAL-II / JB-3.1
│   │   │   ├── GAPM-FRONTAL-TO-OCCIPITAL / JB-3.1
│   │   │   ├── GAPM-FRONTAL-TO-TEMPORAL-I / JB-3.1
│   │   │   ├── INFERIOR_FRONTAL_GYRUS
│   │   │   │   ├── AREA-44 / JB-3.1
│   │   │   │   └── AREA-45 / JB-3.1
│   │   │   ├── INFERIOR_FRONTAL_SULCUS
│   │   │   │   ├── AREA-IFJ1 / JB-3.1
│   │   │   │   ├── AREA-IFJ2 / JB-3.1
│   │   │   │   ├── AREA-IFS1 / JB-3.1
│   │   │   │   ├── AREA-IFS2 / JB-3.1
│   │   │   │   ├── AREA-IFS3 / JB-3.1
│   │   │   │   └── AREA-IFS4 / JB-3.1
│   │   │   ├── LATERAL_ORBITOFRONTAL_CORTEX
│   │   │   │   ├── AREA-FO4 / JB-3.1
│   │   │   │   ├── AREA-FO5 / JB-3.1
│   │   │   │   ├── AREA-FO6 / JB-3.1
│   │   │   │   └── AREA-FO7 / JB-3.1
│   │   │   ├── MEDIAL_ORBITOFRONTAL_CORTEX
│   │   │   │   ├── AREA-FO1 / JB-3.1
│   │   │   │   ├── AREA-FO2 / JB-3.1
│   │   │   │   └── AREA-FO3 / JB-3.1
│   │   │   ├── MESIAL_PRECENTRAL_GYRUS
│   │   │   │   └── AREA-6MP / JB-3.1
│   │   │   ├── MFG_IFS
│   │   │   │   └── AREA-MFG5 / JB-3.1
│   │   │   ├── MIDDLE_FRONTAL_GYRUS
│   │   │   │   ├── AREA-8V1 / JB-3.1
│   │   │   │   ├── AREA-8V2 / JB-3.1
│   │   │   │   ├── AREA-MFG1 / JB-3.1
│   │   │   │   └── AREA-MFG4 / JB-3.1
│   │   │   ├── POSTERIOR_SFG
│   │   │   │   └── AREA-6MA / JB-3.1
│   │   │   ├── PRECENTRAL_GYRUS
│   │   │   │   ├── AREA-4A / JB-3.1
│   │   │   │   └── AREA-4P / JB-3.1
│   │   │   ├── SUPERIOR_FRONTAL_GYRUS
│   │   │   │   ├── AREA-8D1 / JB-3.1
│   │   │   │   ├── AREA-8D2 / JB-3.1
│   │   │   │   ├── AREA-SFG2 / JB-3.1
│   │   │   │   └── AREA-SFG4 / JB-3.1
│   │   │   ├── SUPERIOR_FRONTAL_SULCUS
│   │   │   │   ├── AREA-6D3 / JB-3.1
│   │   │   │   ├── AREA-SFG3 / JB-3.1
│   │   │   │   ├── AREA-SFS1 / JB-3.1
│   │   │   │   └── AREA-SFS2 / JB-3.1
│   │   │   ├── VENTRAL_PRECENTRAL_GYRUS
│   │   │   │   ├── AREA-6R1 / JB-3.1
│   │   │   │   ├── AREA-6V1 / JB-3.1
│   │   │   │   ├── AREA-6V2 / JB-3.1
│   │   │   │   └── AREA-6V3 / JB-3.1

```

```

BRAIN
├── TELECEPHALON
│   ├── CEREBRAL_CORTEX
│   │   ├── INSULA
│   │   │   ├── AGRANULAR_INSULA
│   │   │   │   ├── AREA-IA1 / JB-3.1
│   │   │   │   ├── AREA-IA2 / JB-3.1
│   │   │   │   └── AREA-IA3 / JB-3.1
│   │   │   ├── DYSGRANULAR_INSULA
│   │   │   │   ├── AREA-ID1 / JB-3.1
│   │   │   │   ├── AREA-ID10 / JB-3.1
│   │   │   │   ├── AREA-ID2 / JB-3.1
│   │   │   │   ├── AREA-ID3 / JB-3.1
│   │   │   │   ├── AREA-ID4 / JB-3.1
│   │   │   │   ├── AREA-ID5 / JB-3.1
│   │   │   │   ├── AREA-ID6 / JB-3.1
│   │   │   │   ├── AREA-ID7 / JB-3.1
│   │   │   │   ├── AREA-ID8 / JB-3.1
│   │   │   │   └── AREA-ID9 / JB-3.1
│   │   │   ├── GRANULAR_INSULA
│   │   │   │   ├── AREA-IG1 / JB-3.1
│   │   │   │   ├── AREA-IG2 / JB-3.1
│   │   │   │   └── AREA-IG3 / JB-3.1

```

```

BRAIN
├── TELECEPHALON
│   ├── CEREBRAL_CORTEX
│   │   ├── LIMBIC LOBE
│   │   │   ├── FRONTAL_CINGULATE_GYRUS
│   │   │   │   ├── AREA-25
│   │   │   │   │   ├── AREA-25.25A / JB-3.1
│   │   │   │   │   └── AREA-25.25P / JB-3.1
│   │   │   │   ├── AREA-33 / JB-3.1
│   │   │   │   ├── AREA-P24AB
│   │   │   │   │   ├── AREA-P24AB.P24A / JB-3.1
│   │   │   │   │   └── AREA-P24AB.P24B / JB-3.1
│   │   │   │   ├── AREA-P24C
│   │   │   │   │   ├── AREA-P24C.PD24CD / JB-3.1
│   │   │   │   │   ├── AREA-P24C.PD24CV / JB-3.1
│   │   │   │   │   └── AREA-P24C.PV24C / JB-3.1
│   │   │   │   ├── AREA-P32 / JB-3.1
│   │   │   │   ├── AREA-S24
│   │   │   │   │   ├── AREA-S24.S24A / JB-3.1
│   │   │   │   │   └── AREA-S24.S24B / JB-3.1
│   │   │   │   └── AREA-S32 / JB-3.1
│   │   │   ├── HIPPOCAMPAL_FORMATION
│   │   │   │   ├── AREA-EC / JB-3.1
│   │   │   │   ├── HIPPOCAMPUS-CA1 / JB-3.1
│   │   │   │   ├── HIPPOCAMPUS-CA2 / JB-3.1
│   │   │   │   ├── HIPPOCAMPUS-CA3 / JB-3.1
│   │   │   │   ├── HIPPOCAMPUS-DG / JB-3.1
│   │   │   │   ├── HIPPOCAMPUS-HATA / JB-3.1
│   │   │   │   ├── HIPPOCAMPUS-SUBC
│   │   │   │   │   ├── HIPPOCAMPUS-SUBC.PAS / JB-3.1
│   │   │   │   │   ├── HIPPOCAMPUS-SUBC.PRES / JB-3.1
│   │   │   │   │   ├── HIPPOCAMPUS-SUBC.PROS / JB-3.1
│   │   │   │   │   └── HIPPOCAMPUS-SUBC.SUB / JB-3.1
│   │   │   │   └── HIPPOCAMPUS-TRS / JB-3.1
│   │   │   ├── OLFACTORY_CORTEX
│   │   │   │   ├── BASALFOREBRAIN-TU / JB-3.1
│   │   │   │   └── BASALFOREBRAIN-TUTI / JB-3.1
│   │   │   ├── PIRIFORM_CORTEX
│   │   │   │   ├── AREA-PIRT
│   │   │   │   │   ├── AREA-PIRT.TIT / JB-3.1
│   │   │   │   │   └── AREA-PIRT.TU / JB-3.1
│   │   │   │   ├── AREA-PIRTB
│   │   │   │   │   ├── AREA-PIRTB.TBD / JB-3.1
│   │   │   │   │   └── AREA-PIRTB.TBV / JB-3.1

```

```

BRAIN
├── TELECEPHALON
│   ├── CEREBRAL_CORTEX
│   │   └── OCCIPITAL_LOBE
│   │       ├── DORSAL_OCCIPITAL_CORTEX
│   │       │   ├── AREA-HOC3D / JB-3.1
│   │       │   ├── AREA-HOC4D / JB-3.1
│   │       │   └── AREA-HOC6 / JB-3.1
│   │       ├── LATERAL_OCCIPITAL_CORTEX
│   │       │   ├── AREA-HOC4LA / JB-3.1
│   │       │   ├── AREA-HOC4LP / JB-3.1
│   │       │   └── AREA-HOC5 / JB-3.1
│   │       ├── OCCIPITAL_CORTEX
│   │       │   ├── AREA-HOC1 / JB-3.1
│   │       │   └── AREA-HOC2 / JB-3.1
│   │       └── VENTRAL_OCCIPITAL_CORTEX
│   │           ├── AREA-HOC3V / JB-3.1
│   │           └── AREA-HOC4V / JB-3.1

```

```

BRAIN
├── TELECEPHALON
│   ├── CEREBRAL_CORTEX
│   │   └── PARIETAL_LOBE
│   │       ├── INFERIOR_PARIETAL_LOBULE
│   │       │   ├── AREA-PF / JB-3.1
│   │       │   ├── AREA-PFCM / JB-3.1
│   │       │   ├── AREA-PFM / JB-3.1
│   │       │   ├── AREA-PFOP / JB-3.1
│   │       │   ├── AREA-PFT / JB-3.1
│   │       │   ├── AREA-PGA / JB-3.1
│   │       │   └── AREA-PGP / JB-3.1
│   │       ├── INTRAPARIETAL_SULCUS
│   │       │   ├── AREA-HIP1 / JB-3.1
│   │       │   ├── AREA-HIP2 / JB-3.1
│   │       │   ├── AREA-HIP3 / JB-3.1
│   │       │   ├── AREA-HIP4 / JB-3.1
│   │       │   ├── AREA-HIP5 / JB-3.1
│   │       │   ├── AREA-HIP6 / JB-3.1
│   │       │   ├── AREA-HIP7 / JB-3.1
│   │       │   └── AREA-HIP8 / JB-3.1
│   │       ├── PARIETAL_OPERCULUM
│   │       │   ├── AREA-OP1 / JB-3.1
│   │       │   ├── AREA-OP2 / JB-3.1
│   │       │   ├── AREA-OP3 / JB-3.1
│   │       │   └── AREA-OP4 / JB-3.1
│   │       ├── PARIETO-OCCIPITAL_SULCUS
│   │       │   └── AREA-HPO1 / JB-3.1
│   │       ├── POSTCENTRAL_GYRUS
│   │       │   ├── AREA-1 / JB-3.1
│   │       │   ├── AREA-2 / JB-3.1
│   │       │   ├── AREA-3A / JB-3.1
│   │       │   └── AREA-3B / JB-3.1
│   │       └── SUPERIOR_PARIETAL_LOBULE
│   │           ├── AREA-5CI / JB-3.1
│   │           ├── AREA-5L / JB-3.1
│   │           ├── AREA-5M / JB-3.1
│   │           ├── AREA-7A / JB-3.1
│   │           ├── AREA-7M / JB-3.1
│   │           ├── AREA-7P / JB-3.1
│   │           └── AREA-7PC / JB-3.1

```

```

BRAIN
├── TELECEPHALON
│   ├── CEREBRAL_CORTEX
│   │   └── TEMPORAL_LOBE
│   │       ├── COLLATERAL_SULCUS
│   │       │   └── AREA-COS1 / JB-3.1
│   │       ├── FUSIFORM_GYRUS
│   │       │   ├── AREA-FG1 / JB-3.1
│   │       │   ├── AREA-FG2 / JB-3.1
│   │       │   ├── AREA-FG3 / JB-3.1
│   │       │   ├── AREA-FG4 / JB-3.1
│   │       │   └── AREA-FG5 / JB-3.1
│   │       ├── GAPM-FRONTAL-TO-TEMPORAL-II / JB-3.1
│   │       ├── GAPM-TEMPORAL-TO-PARIETAL / JB-3.1
│   │       ├── HESCHL_GYRUS
│   │       │   ├── AREA-TE-1.0 / JB-3.1
│   │       │   ├── AREA-TE-1.1 / JB-3.1
│   │       │   └── AREA-TE-1.2 / JB-3.1
│   │       ├── OCCIPITO-TEMPORAL_SULCUS
│   │       │   └── AREA-OTS1 / JB-3.1
│   │       ├── PARAHIPPOCAMPAL_GYRUS
│   │       │   ├── AREA-PH1 / JB-3.1
│   │       │   ├── AREA-PH2 / JB-3.1
│   │       │   └── AREA-PH3 / JB-3.1
│   │       ├── SUPERIOR_TEMPORAL_GYRUS
│   │       │   ├── AREA-TE-2.1 / JB-3.1
│   │       │   ├── AREA-TE-2.2 / JB-3.1
│   │       │   └── AREA-TE-3 / JB-3.1
│   │       ├── SUPERIOR_TEMPORAL_SULCUS
│   │       │   ├── AREA-ST1 / JB-3.1
│   │       │   └── AREA-ST2 / JB-3.1
│   │       ├── TEMPORAL_INSULA
│   │       │   ├── AREA-TEI / JB-3.1
│   │       │   └── AREA-TI / JB-3.1
│   │       └── TEMPORO-PARIETAL_JUNCTION
│   │           └── AREA-TPJ / JB-3.1

```

```

BRAIN
├── TELECEPHALON
│   └── CEREBRAL_NUCLEI
│       ├── AMYGDALA
│       │   ├── AMYGDALA-CM
│       │   │   ├── AMYGDALA-CM.AAA / JB-3.1
│       │   │   ├── AMYGDALA-CM.CE / JB-3.1
│       │   │   └── AMYGDALA-CM.ME / JB-3.1
│       │   ├── AMYGDALA-IF
│       │   │   ├── AMYGDALA-IF.ICE / JB-3.1
│       │   │   ├── AMYGDALA-IF.IOL / JB-3.1
│       │   │   └── AMYGDALA-IF.LD / JB-3.1
│       │   ├── AMYGDALA-LB
│       │   │   ├── AMYGDALA-LB.BL / JB-3.1
│       │   │   ├── AMYGDALA-LB.BM / JB-3.1
│       │   │   ├── AMYGDALA-LB.LA / JB-3.1
│       │   │   └── AMYGDALA-LB.PL / JB-3.1
│       │   ├── AMYGDALA-MF
│       │   │   ├── AMYGDALA-MF.ICM / JB-3.1
│       │   │   └── AMYGDALA-MF.LM / JB-3.1
│       │   ├── AMYGDALA-SF
│       │   │   ├── AMYGDALA-SF.AHI / JB-3.1
│       │   │   ├── AMYGDALA-SF.APIR / JB-3.1
│       │   │   └── AMYGDALA-SF.VCO / JB-3.1
│       │   ├── AMYGDALA-VTM / JB-3.1
│       │   └── AMYGDALOSTRIATUM_TZ
│       │       └── AMYGDALA-ASTR / JB-3.1
│       ├── BASAL_FOREBRAIN
│       │   ├── BASALFOREBRAIN-BST / JB-3.1
│       │   ├── BASALFOREBRAIN-CH-123 / JB-3.1
│       │   └── SUBLENTICULAR_BASAL_FOREBRAIN
│       │       └── BASALFOREBRAIN-CH-4 / JB-3.1
│       └── BASAL_GANGLIA
│           ├── Caudate / AAL3
│           ├── N_Acc / AAL3
│           ├── Pallidum / AAL3
│           ├── Putamen / AAL3
│           ├── VENTRAL_PALLIDUM
│           │   └── VENTRALPALLIDUM-VP / JB-3.1
│           └── VENTRAL_STRIATUM
│               ├── VENTRALSTRIATUM-ACBL / JB-3.1
│               ├── VENTRALSTRIATUM-ACBM / JB-3.1
│               ├── VENTRALSTRIATUM-FUCD / JB-3.1
│               └── VENTRALSTRIATUM-FUP / JB-3.1

```

References

- Amunts, K., H. Mohlberg, S. Bludau and K. Zilles (2020). Julich-Brain: A 3D probabilistic atlas of the human brain's cytoarchitecture. *Science* **369**(6506): 988-992.
- Benjamini, Y. and Y. Hochberg (1995). Controlling the False Discovery Rate: A Practical and Powerful Approach to Multiple Testing. *Journal of the Royal Statistical Society: Series B (Methodological)* **57**(1): 289-300.
- Bogomolov, M., C. B. Peterson, Y. Benjamini and C. Sabatti (2020). Hypotheses on a tree: new error rates and testing strategies. *Biometrika* **108**(3): 575-590.
- Hofhansel, L., C. Weidler, M. Votinov, B. Clemens, A. Raine and U. Habel (2020). Morphology of the criminal brain: gray matter reductions are linked to antisocial behavior in offenders. *Brain Structure and Function* **225**(7): 2017-2028.
- Müller, J. L., S. Gänßbauer, M. Sommer, K. Döhnelt, T. Weber, T. Schmidt-Wilcke and G. Hajak (2008). Gray matter changes in right superior temporal gyrus in criminal psychopaths. Evidence from voxel-based morphometry. *Psychiatry Research: Neuroimaging* **163**(3): 213-222.
- Rolls, E. T., C.-C. Huang, C.-P. Lin, J. Feng and M. Joliot (2020). Automated anatomical labelling atlas 3. *NeuroImage* **206**: 116189.
- Schneider, F., U. Habel, C. Kessler, S. Posse, W. Grodd and H.-W. Müller-Gärtner (2000). Functional Imaging of Conditioned Aversive Emotional Responses in Antisocial Personality Disorder. *Neuropsychobiology* **42**(4): 192-201.
- Simes, R. J. (1986). An improved Bonferroni procedure for multiple tests of significance. *Biometrika* **73**(3): 751-754.
- Tzourio-Mazoyer, N., B. Landeau, D. Papathanassiou, F. Crivello, O. Etard, N. Delcroix, B. Mazoyer and M. Joliot (2002). Automated anatomical labeling of activations in SPM using a macroscopic anatomical parcellation of the MNI MRI single-subject brain. *NeuroImage* **15**(1): 273-289.