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Stages of granulovacuolar degeneration: their relation to Alzheimer's disease and chronic stress response

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Abstract Granulovacuolar degeneration (GVD) is characterized by the presence of vacuolar cytoplasmic lesions in nerve cells of the medial temporal lobe. These changes occur in older non-diseased individuals as well as in patients with Alzheimer's disease (AD), progressive supranuclear palsy (PSP), Pick's, sporadic Parkinson's (PD), and Guam diseases. We stained representative paraffin sections from all parts of the brain with anti-pTDP43, anti-CK18 or anti-CK1E from 14 non-demented elderly, 19 AD, 17 non-AD tauopathy, 9 sporadic PD, and 5 TDP43proteinopathy [amyotrophic lateral sclerosis (ALS) and frontotemporal lobar degeneration (FTLD)] cases. Our results showed five stages of GVD based on its distribution pattern: GVD began in the hippocampal subfields CA1, CA2, and the subiculum. In a second stage, entorhinal cortex, and CA4 neurons exhibited GVD. Additional neurons were involved in the temporal neocortex in stage 3, whereas the affection of the amygdala and/or the hypothalamus marked stage 4. A fifth and final stage was characterized by additional GVD in the cingulate cortex and occasionally in the frontal and parietal cortices as well as in the oral raphe and pedunculopontine tegmental nuclei. The GVD stages correlated with neurofibrillary tangle stages, Consortium to Establish a Registry for AD (CERAD) scores for neuritic plaque pathology, amyloid β-protein deposition phases, cerebral amyloid angiopathy stages, and clinical dementia rating (CDR) scores. No associations were seen between GVD stage and the presence of non-AD tauopathies, PD, ALS, or FTLD cases. In conclusion, GVD affects neurons in a hierarchical sequence that allows the distinction of five stages. The topographic distribution of GVD restricted to regions involved in response to chronic stress could indicate a link between GVD and chronically stressful influences. Moreover, the association of the GVD stages with those of AD-related pathology but not with other neurodegenerative disorders points to a possible role of GVD and the response to chronic stress in the pathogenesis of AD.

Keywords Granulovacuolar degeneration · Amyloid β-protein · Neurofibrillary tangles · τ -Protein · Dementia · Argyrophilic grain disease · Circadian clock · Chronic stress-response · pTDP43

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Introduction

Granulovacuolar degeneration (GVD) is marked by the presence of vacuolar changes in the cytoplasm of neurons of the medial temporal lobe (MTL) [4, 36, 49]. GVD lesions have also been described in the neocortex, hypothalamus, amygdala, and in the paramedian nuclei of the midbrain [37]. GVD-affected neurons were more frequently seen in Alzheimer's disease (AD) cases than in controls [3, 14]. Other authors did not find such a correlation between GVD and AD when comparing AD cases with non-demented



cases with AD-related pathology [46]. Down syndrome cases also exhibited GVD in age-related amounts rather than in relation to AD-related neurofibrillary tangle (NFT) pathology and synapse loss [5]. Moreover, GVD has been reported in association with other tauopathies, such as Guam disease [21], Pick's disease, and progressive supranuclear palsy (PSP) as well as in the synucleinopathies sporadic Parkinson's disease (PD) and dementia with Lewy bodies [35]. Thus, although GVD is often referred to as an "AD-related" histopathological phenomenon it is, nevertheless, presently unclear whether it is only associated with AD or with other tauopathies and with synucleinopathies as well.

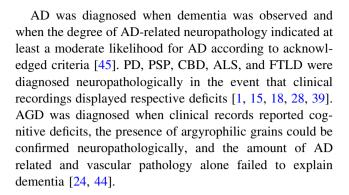
Recently, antibodies directed against the phosphorylated transactive-response DNA-binding protein TDP43 (pTDP43) and casein kinase 1 (CK1) isoforms δ or ϵ have been reported to stain GVD [17, 25–27, 35]. They can be used to study GVD pathology more easily than in hematoxylin–eosin or silver stained sections.

To determine the anatomical distribution of GVD in the human brain and to clarify its relationship to dementia and AD, we studied non-demented elderly, AD, non-AD tauopathy, sporadic PD, TDP43-proteinopathy cases for the distribution of GVD and its relationship to concomitant AD-related NFT and amyloid β -protein (A β) pathology as well as to sporadic PD, argyrophilic grain disease (AGD), PSP, corticobasal degeneration (CBD), amyotrophic lateral sclerosis (ALS), and frontotemporal lobar degeneration (FTLD).

Materials and methods

Study cohort

A total sample of 64 autopsy cases between 59 and 98 years of age covering all stages of AD-related NFT and AB pathology was investigated, including 14 non-demented elderly, 19 AD, 17 non AD tauopathy (9 AGD, 8 PSP or CBD), 9 PD, and 5 TDP43-proteinopathy (3 ALS and 2 FTLD) cases (Table 1). All autopsy brains were collected in accordance with local ethical committee guidelines and the federal law governing the use of human tissue in Germany and The Netherlands. Scores from the clinical dementia rating (CDR) [23] were observed retrospectively for 56 cases as previously described [42]. These data were also employed to determine whether individuals clinically met the DSM-IV criteria for dementia [2]. In three demented cases, the clinical records noted the diagnosis of dementia according to DSM-IV but did not provide sufficient information to determine the CDR score. In five non-demented cases, the clinical records did not provide sufficient information to determine the CDR score.



Neuropathology and immunohistochemistry

The brains of all individuals were fixed in a 4% aqueous formaldehyde solution for at least 3 weeks before undergoing neuropathological evaluation. Tissue blocks covering the regions listed in Table 2 were excised and embedded in paraffin or polyethylene glycol (PEG; Merck-Schuchardt, Hohenbrunn, Germany). The paraffin sections were cut at 10 μm , whereas the PEG-blocks were sectioned at 100 μm .

For the detection of GVD and other neurodegenerative lesions, silver staining methods and immunohistochemistry with the primary antibodies listed in Table 3 were applied. Primary antibodies were detected with biotinylated secondary antibodies and the ABC complex (Vectastain: Vector Laboratories, Burlingame, CA, USA). This reaction was subsequently visualized with 3,3-diaminobenzidine (DAB). Immunolabeled paraffin sections were counterstained with hematoxylin. Positive and negative controls were included to confirm immunostaining results. GVD was diagnosed when multiple vacuoles (2 aggregated vacuoles or more) with granular inclusions were detected with at least one GVD-detecting antibody (Table 3). Solitary autophagic vacuoles were not considered as GVD.

To confirm co-localization of CK1δ, CK1ε, and pTDP43 in GVD lesions, sections from the hippocampus of randomly chosen cases exhibiting GVD were used for double label immunofluorescence with anti-pTDP43 and either anti-CK1δ or anti-CK1ε. The primary antibodies were detected with carbocyanin Cy2 and Cy3 labeled secondary antibodies. Sections were mounted in Eukitt (O. Kindler; Freiburg, Germany) and cover-slipped. All tissue sections were viewed with a Leica DMLB or an Olympus BX 51 microscope. Digital photographs were obtained with Leica cameras (DC 500 or EC3).

Diagnosis of the stages in the development of NFTs (NFT stage) and the semiquantitative assessment of neuritic plaques [Consortium to Establish a Registry for AD (CERAD) score] were performed in accordance with published and recommended criteria [7, 9, 29, 45]. For staging of $A\beta$ pathology, we used a previously published



Table 1 List of autopsy cases studied

Case no.	Age	Sex	Diagnosis	CDR	PD stage	NFT stage	Aβ phase	CERAD	CAA stage	GVD stage
1	66	f	Control	0	n.a.	0	0	0	0	0
2	62	m	Control	0	n.a.	0	0	0	0	0
3	64	m	Control	0	n.a.	1	2	0	0	0
4	71	m	Control	0	n.a.	1	3	0	0	0
5	61	m	Control	0	n.a.	0	0	0	0	0
6	59	m	Control	0	0	0	0	0	0	0
7	71	m	Control	0	n.a.	1	2	0	1	1
8	69	f	Control	0	n.a.	0	0	0	0	1
9	74	m	Control	0	n.a.	1	0	0	0	2
10	83	f	Control	0	n.a.	3	1	0	1	2
11	72	m	Control	0	n.a.	3	2	0	0	3
12	83	f	Control	0	n.a.	4	3	1	2	3
13	85	m	Control	0	n.a.	3	2	1	0	4
14	63	f	Control	0	n.a.	3	4	1	0	4
15	73	f	AGD	0.5	n.a.	1	1	0	1	0
16	81	m	AGD	2	n.a.	2	3	0	0	1
17	77	m	AGD	0.5	n.a.	3	2	1	0	2
18	88	m	AGD	2	n.a.	2	3	1	1	3
19	79	m	AGD	2	n.a.	2	3	0	2	4
20	79	f	AGD-VD	2	n.a.	2	0	0	0	4
21	80	f	AGD, PD	1	3	2	3	0	0	0
22	79	m	AGD, PD	0	3	2	0	0	0	1
23	75	m	AGD, PSP, PD	2	4	3	2	0	1	0
24	78	f	CBD	3	n.a.	1	1	0	0	1
25 ^a	76	m	PSP	0	n.a.	0	2	0	0	0
26 ^a	69	f	PSP	0	n.a.	2	1	0	0	0
27 ^b	59	m	PSP	n.a.	n.a.	2	1	0	1	0
28 ^a	81	m	PSP	2	n.a.	4	4	2	3	2
29 ^a	76	m	PSP	1	n.a.	1	1	0	0	3
$30^{\rm b}$	83	m	PSP	n.a.	n.a.	3	2	0	0	4
31 ^a	84	f	PSP, AGD	0	n.a.	0	0	0	1	0
32	83	f	PD	n.a.	4	2	0	0	0	0
33	69	f	PD	n.a.	4	1	3	0	1	0
34	68	f	PD	2	5	1	3	0	1	0
35	77	m	PD	0	4	1	0	0	0	0
36 ^b	76	m	PD	n.a.	5	3	4	0	1	0
37	64	f	PD	0	4	3	0	0	1	1
38	68	m	PD	0.5	5	3	4	2	2	1
39	76	f	PD	0	4	2	0	0	0	1
40	77	m	PD	1	5	4	3	2	2	1
41	68	m	ALS	0	n.a.	2	2	0	1	2
42	75	m	ALS	0	n.a.	2	1	1	1	2
43	76	f	ALS	0	n.a.	2	0	0	0	2
44 ^c	76	f	FTLD-TDP	3	n.a.	0	3	0	2	0
45	84	m	FTLD-TDP, AD, DLB	3	6	6	4	3	2	5
46	64	f	AD	n.a.	n.a.	6	4	3	1	1
47	62	f	AD	3	n.a.	6	4	3	1	2
48	72	f	AD	1	n.a.	4	4	2	1	2



Table 1 continued

Case no.	Age	Sex	Diagnosis	CDR	PD stage	NFT stage	Aβ phase	CERAD	CAA stage	GVD stage
49	86	f	AD	3	n.a.	6	4	2	2	3
50	87	f	AD	3	n.a.	4	4	1	2	3
51	87	f	AD	0.5	n.a.	4	4	2	2	3
52	79	f	AD	n.a.	n.a.	4	3	2	1	4
53	82	m	AD	2	n.a.	3	3	2	1	4
54	89	f	AD	3	n.a.	5	4	3	2	4
55	98	f	AD	2	n.a.	4	4	3	2	4
56	68	f	AD	1	n.a.	6	4	2	1	4
57	78	m	AD	3	n.a.	4	4	1	2	4
58	83	f	AD	n.a.	n.a.	4	4	2	1	5
59	78	m	AD	1	n.a.	4	4	1	1	5
60	81	f	AD	3	n.a.	5	4	1	1	5
61	83	m	AD	3	n.a.	5	4	2	3	5
62	83	m	AD	3	n.a.	4	4	3	2	5
63	98	f	AD	0.5	n.a.	4	4	1	2	5
64	78	f	AD	3	n.a.	5	4	2	3	5

The table shows age in years, sex, neuropathological diagnosis, the degree of dementia (CDR score) [31], the PD stage according to Braak et al. [11], the stage of neurofibrillary tangle pathology according to Braak and Braak [9] (NFT stage), the $A\beta$ phase representing the distribution of $A\beta$ deposits in the subfields of the MTL [43], the CERAD score for the frequency of neuritic plaques according to Mirra et al. [29], the stages of CAA [40], and GVD

m Male, f female, (control) non-demented control, AD Alzheimer's disease, AGD argyrophilic grain disease, ALS amyotrophic lateral sclerosis, CBD corticobasal degeneration, DLB dementia with Lewy bodies, FTLD-TDP frontotemporolobar degeneration with TDP43-pathology, n.a. not assessed, PD Parkinson's disease, PSP progressive supranuclear palsy, VD vascular dementia

- ^a Indicates cases for which brain stem and diencephalic samples were not available
- b Indicates cases for which samples of the occipital lobe were not available

protocol for four phases of β -amyloidosis in the MTL [43]. Cerebral amyloid angiopathy (CAA) was diagnosed whenever vascular A β deposition was observed. The stage of the distribution of CAA-affected vessels throughout the brain was assessed according to published criteria [40, 41]. PD lesions in PD cases were staged according to previously published criteria [1, 11].

Statistical analysis

The presence or absence of GVD in a given brain region was assessed dichotomously with anti-pTDP43, anti-CK1 δ , and/or anti-CK1 ϵ . The brain areas listed in Table 2 were analyzed separately. Cluster analysis was performed to identify a staging pattern. The staging pattern was validated by testing the stage-related affection with GVD for each region studied with the Fisher's exact-test and subsequent trend-test. Partial correlation analysis controlled for age and gender was performed to describe the relationship between GVD stage and AD-related pathology as well as between GVD stage and the CDR score. Logistic regression analysis was performed controlling for age and gender to test whether AD was associated with the stages of GVD pathology.

Likewise, logistic regression was used to analyze the association between non-AD tauopathies (i.e., AGD, PSP, and CBD), PD, and TDP43-proteinopathies (ALS, FTLD) on the one hand and the GVD stages on the other. All statistical tests were calculated with the help of SPSS-statistical software, release 19.0 (Chicago, IL, USA). To correct for multiple testing Bonferroni correction was used.

Results

Anti-pTDP43 stained GVD similar to anti-CK1 δ and anti-CK1 ϵ (Fig. 1a–c). Positively stained material lacking the typical vacuolar pattern was not considered as GVD. Double label immunofluorescence confirmed the presence of CK1 δ and CK1 ϵ , both antibodies co-localizing pTDP43 in GVD lesions (Fig. 1d–i). With these antibodies, a similar distribution pattern throughout all investigated brain regions was observed, thereby indicating that these methods were interchangeable for GVD staging. A region-specific differential expression of these proteins in GVD lesions was not seen. Immunohistochemistry with the antibody against abnormal τ protein appeared less useful



^c Indicates cases for which samples of diencephalic regions and the parietal cortex were not available

Table 2 Frequencies of GVD prevalence in the brain regions studied at each stage of GVD

	GVD Stage 1	GVD Stage 2	GVD Stage 3	GVD Stage 4	GVD Stage 5
CA1/subiculum*	100.00%	100.00%	100.00%	100.00%	100.00%
entorhinal cortex (Brodmann area 28)*	0.00%	75.00%	100.00%	100.00%	100.00%
CA4*	0.00%	50.00%	80.00%	77.78%	100.00%
temporal cortex (Brodmann area 35/36)*	0.00%	0.00%	100.00%	100.00%	100.00%
NBM*	0.00%	0.00%	20.00%	55.56%	50.00%
Hypothalamus*	0.00%	0.00%	0.00%	88.89%	100.00%
Amygdala*	0.00%	0.00%	0.00%	77.78%	100.00%
cingulate gyrus (Brodmann area 24)*	0.00%	0.00%	0.00%	0.00%	75.00%
Oral raphe/ PPTg nuclei*	0.00%	0.00%	0.00%	0.00%	62.50%
Thalamus	0.00%	0.00%	0.00%	0.00%	37.50%
frontal cortex (Brodmann area 6)	0.00%	0.00%	0.00%	0.00%	12.50%
parietal cortex (Brodmann area 7)	0.00%	0.00%	0.00%	0.00%	12.50%
occipital cortex (Brodmann areas 17, 18, 19)	0.00%	0.00%	0.00%	0.00%	0.00%
basal ganglia	0.00%	0.00%	0.00%	0.00%	0.00%
cerebellum	0.00%	0.00%	0.00%	0.00%	0.00%
substantia nigra	0.00%	0.00%	0.00%	0.00%	0.00%
red nucleus	0.00%	0.00%	0.00%	0.00%	0.00%
brain stem motor neurons	0.00%	0.00%	0.00%	0.00%	0.00%
brain stem sensory neurons	0.00%	0.00%	0.00%	0.00%	0.00%
brain stem visceromotoric and -sensoric nuclei	0.00%	0.00%	0.00%	0.00%	0.00%
inferior olivary nucleus	0.00%	0.00%	0.00%	0.00%	0.00%
pontine nuclei	0.00%	0.00%	0.00%	0.00%	0.00%

It is shown which percent of cases in a given GVD stage exhibit GVD lesions in a given region. For visualization the frequencies are color-coded: 0% of the cases exhibit GVD lesions = white; 1-25% of the cases exhibit GVD lesions = 20% gray; 26-50% of the cases exhibit GVD lesions = 40% gray; 51-75% of the cases exhibit GVD lesions = 60% gray; 76-99% of the cases exhibit GVD lesions = 80% gray; all cases exhibit GVD lesions = 80% gray; all cases exhibit GVD lesions = 80% gray; 80%

Table 3 List of the conventional and immunohistochemical staining techniques used for the detection of neurodegenerative changes in this study

	GVD	NFT/ NT	Senile plaques	Lewy bodies	Glial τ inclusions	TDP43 threads/ NCI/Nucl. inclusions	Reference/clone, distributor, dilution, pretreatment
Gallyas		+			+		[8]
Campbell-Switzer			+	+			[8, 34]
Anti-abnormal τ protein	+	+			+		AT-8, Pierce-Endogen, USA, 1/1,000
Anti-amyloid β protein			+				4G8, Covance, Dedham, USA, 1/5,000, formic acid pretreatment
Anti-α synuclein				+			anti-syn-1; 1:2,000; Clone number 42, BD Biosciences, Mountain View, CA, USA
Anti-pTDP43	+					+	pS409/410-2, Cosmo Bio Co., Ltd, Tokyo, Japan, 1/10,000, microwave pretreatment
Anti-CK1δ	+						108, 1 polyclonal rabbit, [38], 1/200, microwave pretreatment
Anti-CK1ε	+						712 polyclonal rabbit [13], 1/600, microwave pretreatment
							H-60 polycloncal rabbit, Santa Cruz Biotechnology; Santa Cruz, CA, USA, 1/150, microwave pretreatment

The table indicates which lesions are detected with a given method. The detection of GVD with anti-abnormal phosphorylated τ antibodies is possible [16] but is less easy to analyze because other tauopathic lesions are stained as well as with this antibody

for GVD detection because this antibody also stained other τ lesions [16]. Therefore, the identification of GVD lesions was more difficult than with the other three antibodies.

The distribution of typical pTDP43- and CK1 δ / ϵ -positive GVD throughout the brain regions investigated

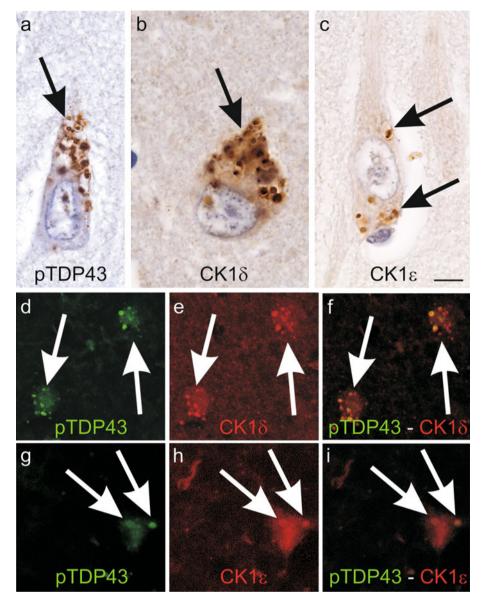
revealed a five-staged hierarchical pattern (Figs. 2, 3, 4, 5, 6; Table 2). The CA1/2-subiculum region always exhibited GVD whenever GVD was observed, regardless whether other brain regions were involved as well. Thus, GVD in the CA1/2 subiculum region in the absence of GVD elsewhere



^{*} indicates those regions that significantly showed differences in the distribution of GVD among the GVD stages (Fisher's exact-test:

^{*} p < 0.005) with a trend to increase with advancing GVD stage (trend-test: * p < 0.005)

Fig. 1 a–c GVD lesions in the subiculum-CA1 region were detected with anti-pTDP43 (arrow in a), with anti-CK1δ (arrow in b), and with anti-CK1ε (arrow in c). Double label immunofluorescence analyses revealed co-expression of pTDP43 and CK1δ (d–f arrows) and of pTDP43 and CK1ε (g–i arrows). (a, c case no. 20; b case no. 25; d–f case no. 29; g–i case no. 26). Calibration bar in c is valid for: a–c 8 μm; d–i 13 μm



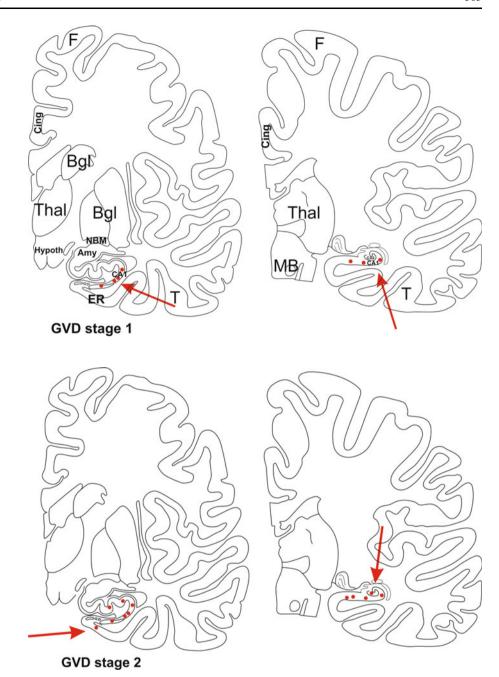
in the brain was referred to as stage 1 GVD (Fig. 2). In addition to GVD in the CA1/2 subiculum region, this pathology was also observed in the entorhinal cortex (Brodmann area 28) in neurons of layers pre- α , pre- β , pri- α , and pri-β and/or in the CA4 subfield of the Ammon's horn (Table 2). GVD in CA4 can be accompanied by GVD lesions in CA3. Stage 2 GVD cases are characterized by this pathology in the absence of further GVD pathology except for occasional GVD-affected neurons in the basal nucleus of Meynert (Fig. 3; Table 2). Expansion of this pathology into layers II/III and/or V of the temporal neocortex (Brodmann areas 35 and 36) (Table 2) was referred to as GVD stage 3 (Fig. 4). In GVD stage 4, GVD-containing neurons were also observed in the amygdala (amygdalohippocampal area, amygdalo-piriform area, central amygdaloid nucleus, basolateral and basomedial nuclear complexes) and/or in hypothalamic neurons of the tuberomammillary, paraventricular, supraoptic, ventromedial, dorsomedian, medial mammillary nuclei and/or nuclei of the anterior and lateral hypothalamic area (Fig. 5; Table 2). In the final stage, *GVD stage* 5, GVD was observed in layers II/III and V neurons of the cingulate gyrus (area 24) as well as in the oral raphe nuclei, the pedunculopontine tegmental nucleus, the superficial and medial thalamic nuclei. Occasionally, GVD lesions were also found in the frontal (Brodmann area 6) and parietal neocortex (Brodmann area 7) (Fig. 6; Table 2). As indicated in Table 2, the presence or absence of GVD in given brain regions resulted in significant differences among the GVD stages. The trend-test confirmed the hierarchical involvement of more and more brain regions (Table 2).

With advancing GVD stage, an increase was also noted in the NFT stage, the CERAD score for neuritic plaques, the phase of $A\beta$ deposition in the MTL, the CAA stage, and



Fig. 2 GVD stage 1: GVDaffected neurons (marked in red) were limited to the CA1subiculum regions (arrows). The right diagram depicts the forebrain hemisphere at the level of the entorhinal region and the hypothalamus, the left shows a section at the level of the lateral geniculate body and the substantia nigra. Anatomical orientation: Amy amygdala, Bgl basal ganglia, incl. Caudate nucleus, putamen and globus pallidus, CA1 CA1/subiculum sector of the hippocampus, CA4 CA4 sector of the hippocampus, cing cingulate gyrus, ER entorhinal cortex, F frontal cortex, Hypoth hypothalamus, MB midbrain, NBM basal nucleus of Meynert, T temporal cortex, Thal thalamus

Fig. 3 GVD stage 2: In addition to neurons affected in GVD stage 1 entorhinal and CA4 neurons exhibited GVD as well (arrows). GVD-affected neurons are marked in red. Basal nucleus of Meynert neurons only occasionally exhibited GVD-affected neurons and were, therefore, not included in this schematic representation because this region was not relevant for staging



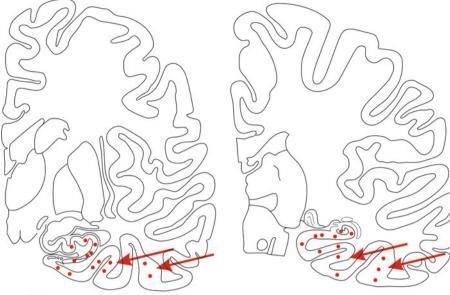
the degree of dementia as represented by the CDR score (Fig. 7a). GVD stage correlated with the NFT stage (partial correlation coefficient for age and gender: r=0.624, p<0.002), with the CERAD neuritic plaque score (partial correlation coefficient for age and gender: r=0.526, p<0.002), the A β phase for plaque deposition in the MTL (partial correlation coefficient for age and gender: r=0.454, p<0.002), CAA stage (partial correlation coefficient for age and gender: r=0.298, p=0.038), and the CDR score for dementia (Partial correlation coefficient for age and gender: r=0.347, p=0.020) (Fig. 7a). After exclusion of the AD cases, there was still a correlation with

NFT stage (partial correlation coefficient for age and gender: r = 0.462, p = 0.006) in the group of all non-AD cases whereas no significant correlation was observed for CERAD neuritic plaque score (p = 0.053), A β phase (p = 0.711), CAA stage (p = 0.999), and CDR score (p = 0.946).

Finally, logistic regression analysis controlled for age and gender revealed an association of AD with the GVD stage (Fig. 7b; odds ratio = 2.741, 95% CI: 1.59–4.724, p < 0.004). After exclusion of the AD cases, non-AD tauopathies (AGD, PSP and CBD), sporadic PD and TDP43-proteinopathies (ALS/FTLD) were not associated with the GVD stage in our sample (Fig. 7c; non-AD

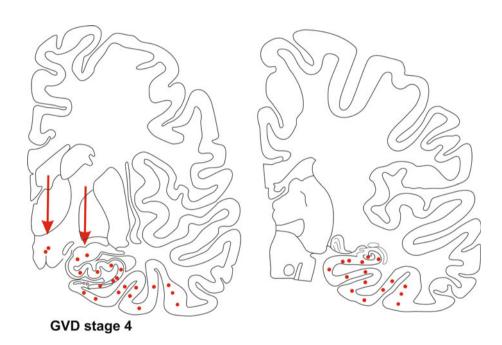


Fig. 4 GVD stage 3: Further GVD-affected neurons (marked in *red*) in addition to those already showing GVD in stage 2 were found in the temporal neocortex (area 36) (arrows). Basal nucleus of Meynert neurons only occasionally exhibited GVD-affected neurons and were, therefore, not included in this schematic representation because this region was not relevant for staging



GVD stage 3

Fig. 5 GVD stage 4: GVD-affected neurons (marked in red) were found in the amygdala and in the hypothalamus (arrows) in addition to those in areas already affected in stage 3. Basal nucleus of Meynert neurons only occasionally exhibited GVD-affected neurons and were, therefore, not included in this schematic representation because this region was not relevant for staging



tauopathies: p = 0.473; PD: p = 0.186; TDP43-proteinopathies: p = 0.989).

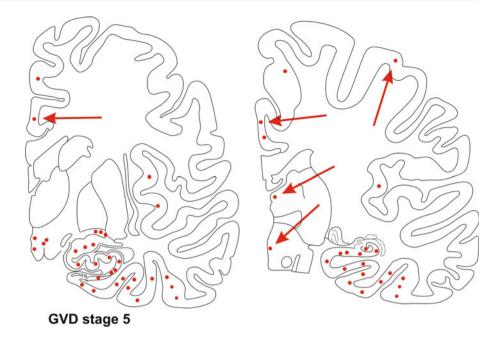
Discussion

Here, we show that GVD represents a pathological lesion in the brain that first develops in neurons of the CA1/2-subiculum region and, thereafter, expands in a predictable sequence into neurons of additional brain regions. This hierarchical distribution pattern makes it possible to distinguish five GVD stages (Figs. 2, 3, 4, 5, 6):

- Stage 1 GVD occurs only in the CA1/2-subiculum region.
- Stage 2 GVD is found in the CA1/2-subiculum region, and additionally in the entorhinal cortex and/or CA4; CA3, and the basal nucleus of Meynert can occasionally exhibit GVD-affected neurons in this stage.
- Stage 3 Stage 2 regions and the temporal neocortex display the presence of GVD.
- Stage 4 GVD is localized in stage 3 regions, subnuclei of the amygdala, and/or the hypothalamus.



Fig. 6 GVD stage 5: The final stage of GVD distribution showed GVD-affected neurons (marked in red) in the cingulate gyrus, other neocortical regions, the thalamus and the raphe nuclei and/or the pedunculopontine tegmental nucleus (PPTg; arrows). Neurons of the basal nucleus of Meynert were also affected



Stage 5 GVD is seen in the CA1/2-subiculum region, entorhinal cortex, CA4, temporal neocortex, subnuclei of the amygdala and the hypothalamus, oral raphe nuclei, pedunculopontine tegmental nucleus, cingulate gyrus, and occasionally in the superficial and medial nuclei of the thalamus, frontal and parietal neocortex.

This staging system confirms the findings of earlier studies that the CA1/2-subiculum region is a predilection site for GVD [4, 49] and, as reported by other groups [4, 37, 49, 50], that GVD, when present, occurs in the hippocampus, entorhinal, temporal and cingulate gyrus, hypothalamus, and amygdala. However, we also found GVD at additional sites, including the oral raphe nuclei, the pedunculopontine tegmental nucleus, superficial and medial thalamic nuclei, the basal nucleus of Meynert, and occasionally in other neocortical areas, whereby the visual cortex (Brodmann areas 17-19) appeared to be spared. The overall distribution pattern of GVD in our study corresponds to that described by other authors [50] and, as shown here, it also correlates with the distribution pattern of AD-related NFTs (NFT stage), A\beta plaques (A\beta phase), neuritic plaques (CERAD neuritic plaque score), CAA (CAA stage), and with the degree of dementia (CDR score). This correlation was not explained by age and gender effects since partial correlation analysis was appropriately controlled for age and gender. As such, it appears more than likely that GVD represents a fourth pathological hallmark of AD in addition to NFTs, $A\beta$ plaques and CAA.

Other neurodegenerative diseases such as PD, non-AD tauopathies (including AGD, PSP, CBD), and TPD43-

proteinopathies, such as ALS/FTLD were not associated with increased GVD stages although GVD (mean GVD stage 1.29; range 0-5) was observed in 42-80% of these cases similar to age-matched controls. Logistic regression models controlled for age and gender confirmed the association of GVD stage with AD. Therefore, the expansion of GVD pathology in its end stage distribution pattern appears to be AD related. An argument against this hypothesis is that GVD also occurs in up to 80% of cases with PSP, AGD, PD, ALS, and FTLD-TDP43. The extent, however, of the GVD lesions throughout the brain most frequently corresponded to early GVD stages, which could just as easily be explained by the presence of concomitant ADrelated pathology in these cases. Since nearly all individuals who died at the age of 40 years or older have been shown to exhibit at least minimal AD-like τ pathology in the brain stem [12] we were unable to find a single "pure" GVD case in the absence of early τ lesions. These latest results may provide additional evidence in favor of a link between GVD and AD. Another argument against GVD as an AD-related pathology is that vacuolar lesions in ballooned neurons have been described previously as GVD in a familial PD case [30]. Nonetheless, this study did not take into account that CK1δ and CK1ε failed to stain GVD within ballooned neurons in our CBD and AGD cases as well as in the sample reported by Schwab et al. [35]. Because we did not study Pick disease cases we cannot exclude an association of this disorder with the extent of GVD-affected neurons as represented by the GVD stage similar to that seen in AD cases.

When comparing studies about GVD it seems to be of major importance to compare the determination of GVD



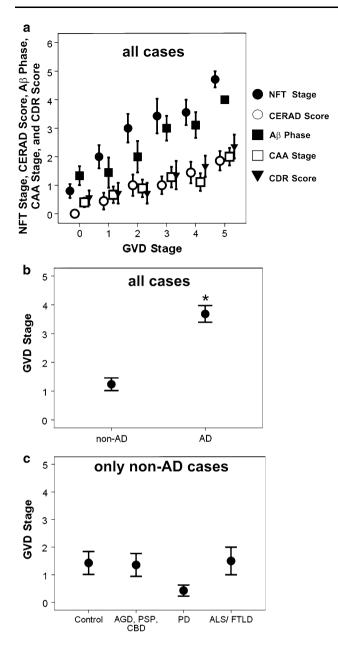
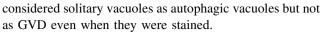


Fig. 7 a Relationship between GVD stage and neurofibrillary pathology (*NFT stage*), neuritic plaque pathology (*CERAD score*), Aβ pathology (*Aβ phase*), CAA (*CAA stage*), and the degree of dementia as represented by the CDR score. The increase of the GVD stage correlated with that of NFT stage, CERAD score, Aβ phase, CAA stage, and the degree of dementia as represented by the CDR score (partial correlation analysis controlled for ages and sex: r = 0.33-0.624; p < 0.05). **b** AD cases exhibited higher GVD stages than non-AD controls (*p < 0.004). **c** After exclusion of the AD cases non-AD tauopathy cases (covering AGD, PSP, and CBD), PD cases and TDP43-proteinopathy cases (covering ALS and FTLD-TDP) did not exhibit higher GVD stages than controls

used by the authors. Here, we determined GVD as pTDP43-, CK1δ- and CK1ε-positive accumulations of at least two aggregated vacuoles containing granular inclusions. We



The stages of GVD proposed here describe the biological process of a hierarchical expansion of this pathology throughout the brain. As such, the GVD stages describe a parameter that is related to the severity of GVD and independently of neuronal loss, which decreases, e.g., the number of CA1 neurons exhibiting GVD, in end-stage AD cases. Since $A\beta$ and τ targeting treatment strategies against AD have been unsuccessful to date [22, 32, 48] GVD presumably will be important for the understanding of ADrelated pathomechanisms as well as for the identification of alternative therapeutic targets. In contrast to this potential scientific relevance of GVD and GVD stages, GVD does not contribute to the clinical and neuropathological diagnosis of AD in addition to τ and A β pathology. Therefore, although we do not recommend staging or even diagnosis of GVD lesions for routine neuropathological use, we do recommend it as a parameter for neuropathological studies of neurodegenerative disorders as well as for brain banking purposes. Especially when studying NFT pathology, e.g., within the context of AD, biochemical analysis cannot distinguish between τ protein derived from NFTs and τ protein derived from GVD. Therefore, it is necessary to focus on regions with NFT-only pathology, i.e. regions that do not exhibit GVD when analyzing NFT-related τ pathology.

GVD is not directly linked to NFT pathology or AB plaque deposition. Two 62-year-old AD cases from our sample with NFT stage VI, Aβ-phase 4 but only GVD stages 1 and 2, respectively, support this hypothesis. A non-demented control case (No. 14) exhibiting GVD stage 4, NFT stage III, and Aβ phase 4 demonstrates, on the other hand, that GVD, NFT, and AB pathology alone do not explain dementia, as seen in AD case No. 53 with GVD stage 4, NFT stage III, and Aβ phase 3. Here, the presence or absence of CAA might explain dementia. Thus, GVD itself accounts for the development of late-onset AD, although other factors, such as age of onset, genetic background, NFTs, and parenchymal and vascular AB pathology, significantly contribute as well. It needs to be clarified in the future whether GVD distinguishes specific types of AD or represents a late-onset AD phenomenon. An argument favoring the latter hypothesis is the finding of Ball et al. [5] that GVD does not play a role in Down syndrome cases even when they show severe neurofibrillary pathology.

Neurons exhibiting GVD lesions in our study occurred in regions involved in response to chronic stress [47] and in the modulation of the sleep-wakefulness state [33]. GVD involved neurons of the top-down regulation circuit of the autonomic response to chronic stress, i.e., the hippocampus (CA1, CA2, CA4, subiculum), entorhinal cortex with



adjacent temporal cortex, cingulate gyrus, prefrontal cortex, amygdala, and hypothalamus. Each of these regions receives input from the locus coeruleus which is a major wakefulness-promoting nucleus [6, 33] that becomes activated in response to chronic stressors [47]. A central role for the locus coeruleus in the pathogenesis of AD has been discussed recently because this nucleus exhibits τ pathology very early in life [10, 12] and its degeneration has been shown to promote the development of AD-related lesions in experimental animal models of AD [19, 20]. Thus, one may speculate that chronically stressful influences and disturbed sleep-wakefulness modulation may play a role in the pathogenesis of AD. GVD appears to represent a morphological correlative for the alteration of nerve cells owing to chronic stress response. Further studies are required to verify this hypothesis. Single cell analysis of GVD-affected neurons as well as detection of chronic stress-induced inflammatory response may help to clarify this issue.

In summary, we show here that GVD lesions extend into increasing numbers of brain regions in an ordered sequence that allows the distinction of five stages. In this process, the expansion of GVD lesions correlates with the distribution of AD-related neuropathological lesions but not with other tauopathies (PSP, AGD, CBD), sporadic PD, and TDP43proteinopathies (ALS/FTLD-TDP)—an indication that GVD is associated with AD but not with neurodegeneration in general. In addition, the anatomical distribution pattern of GVD in the brain corresponds to regions involved in chronic stress response and in sleep-wakefulness modulation. Taken together, these points permit us to hypothesize that GVD anatomically related to chronic stress and its processing by the brain may contribute to the development of AD as a fourth factor in addition to the intraneuronal aggregation of τ protein, A β deposition in senile plaques, and CAA.

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