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# Autoantibody profiles in Alzheimer's, Parkinson's, and dementia with Lewy bodies: altered IgG affinity and IgG/IgM/IgA responses to alpha-synuclein, amyloid-beta, and tau in disease-specific pathological patterns

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## Abstract

**Background** Alzheimer's disease (AD) and Parkinson's disease (PD) are leading neurodegenerative disorders marked by protein aggregation, with AD featuring amyloid-beta (A $\beta$ ) and tau proteins, and PD alpha-synuclein (Syn). Dementia with Lewy bodies (DLB) often presents with a mix of these pathologies. This study explores naturally occurring autoantibodies (nAbs), including Immunoglobulin (Ig)G, IgM, and IgA, which target Syn, A $\beta$  and tau to maintain homeostasis and were previously found altered in AD and PD patients, among others.

**Main text** We extended this investigation across AD, PD and DLB patients investigating both the affinities of IgGs and levels of IgGs, IgMs and IgAs towards Syn, A $\beta$  and tau utilizing chemiluminescence assays. We confirmed that AD and PD patients exhibited lower levels of high-affinity anti-A $\beta$  and anti-Syn IgGs, respectively, than healthy controls. AD patients also showed diminished levels of high-affinity anti-Syn IgGs, while anti-tau IgG affinities did not differ significantly across groups. However, DLB patients exhibited increased anti-Syn IgG but decreased anti-Syn IgM levels compared to controls and PD patients, with AD patients showing a similar pattern. Interestingly, AD patients had higher anti-A $\beta$  IgG but lower anti-A $\beta$  IgA levels than DLB patients. DLB patients had reduced anti-A $\beta$  IgM levels compared to controls, and anti-tau IgG levels were lower in AD than PD patients, who had reduced anti-tau IgM levels compared to controls. AD patients uniquely showed higher anti-tau IgA levels. Significant correlations were observed between clinical measures and nAbs, with negative correlations between anti-Syn IgG affinity and levels in DLB

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patients and a positive correlation with anti-Syn IgA levels in PD patients. Disease-specific changes in nAb levels and affinity correlations were identified, highlighting altered immune responses.

**Conclusion** This study reveals distinctive nAb profiles in AD, DLB, and PD, pinpointing specific immune deficiencies against pathological proteins. These insights into the autoreactive immune system's role in neurodegeneration suggest nAbs as potential markers for vulnerability to protein aggregation, offering new avenues for understanding and possibly diagnosing these conditions.

**Keywords** Alzheimer's disease, Dementia with lewy bodies, Parkinson's disease, Naturally occurring autoantibodies, Alpha-synuclein, Amyloid-beta, Tau

## Background

Neurodegenerative diseases are mainly characterized by the pathological accumulation of specific proteins, which play a pivotal role in disease progression. Alzheimer's disease (AD) is characterized by abnormal accumulation of extracellular amyloid-beta (A $\beta$ ) and intracellular tau [1], while Parkinson's disease (PD) is characterized by abnormal intracellular accumulation of alpha-synuclein (Syn) [2]. Dementia with Lewy bodies (DLB) is characterized by increased Lewy body pathology by disease definition, but also shares pathologies with both AD, including A $\beta$  plaques and tau neurofibrillary tangles, in up to 76% of cases. In contrast, non-dementia PD patients share pathology with AD in fewer cases (7–10%) [3–6]. Although still debated, the consensus emphasizes that the aggregation and toxicity of intermediate toxic seed structures of these pathogenic proteins are considered to be key in disease initiation and progression [7–9].

Naturally occurring autoantibodies (nAbs) are a distinct set of antibodies that recognize self- and non-self-antigens without prior immunization and play a pivotal role in immune clearance of neoepitopes, aggregated and misfolded protein [10]. Although they likely cannot reach the intracellular compartment, they contribute to the engulfment of dying cells and aid in their clearance, while also surveilling the extracellular space, inhibiting the transmission of pathological proteins from cell to cell. They have been found in large amounts in healthy individuals as well in aberrant levels in patients with neurodegenerative diseases such as AD, PD, DLB, and other neurological disorders (summarized in Table S1) [11]. Previous studies have shown alterations in the levels and affinity of nAbs against Syn, A $\beta$ , and tau in these diseases, suggesting that dysfunction in the immune clearance of pathological proteins may play an considerable role in the development of neurodegenerative diseases [12, 13]. Generally, there is a consistent pattern observed in the levels and functionality of nAbs in neurodegenerative diseases. Early PD and DLB are characterized by increased levels of anti-Syn nAbs. On the other hand, AD patients, in general, exhibit reduced levels of anti-A $\beta$  nAbs, while no significant differences are observed in anti-tau nAbs (Table S1). Most studies have

predominantly focused on IgG nAbs, under the assumption that immune responses following class-switching are of primary importance. However, significant immune functions are also found in the IgM and IgA antibody classes. IgM nAbs, often regarded as the immune system's "first responders," can rapidly react to alterations in pathological proteins or result in depletion of inhibitors for protein aggregation [14, 15]. IgA's on the other hand play a crucial role in mucosal and gut immunity, which has been implicated as a potential mediator in the pathogenesis of neurodegenerative disorders, such as PD and AD [16, 17]. Furthermore, studies evaluating the functionality of nAbs have revealed important insights. PD patients have been found to have reduced affinity of anti-Syn autoantibodies in both plasma and cerebrospinal fluid (CSF), respectively [18, 19]. This reduction in affinity is also observed in prodromal phases of PD and the atypical parkinsonian disorder multiple system atrophy (MSA) [20]. Similarly, AD patients exhibit reduced affinity for anti-A $\beta$  nAbs [21]. The precise role of nAbs in neurodegenerative disorders, however, remains a subject of ongoing debate and whether these differences in levels, specificity, and efficacy between healthy individuals and those with PD, AD or DLB, suggest that they may contribute to disease onset or progression. However, promising results have been obtained in preclinical animal models, where nAbs have been evaluated in terms of passive immunization (reviewed by [22]). More recently, positive results have been reported in clinical trials for AD using donanemab and lecanemab, both of which target A $\beta$  structures [23, 24].

Here, we evaluated the repertoire of high affinity Immunoglobulin (Ig)G nAbs specific to Syn, A $\beta$ , and tau in AD, PD, and DLB patients compared to healthy controls. We also investigated the levels of nAbs of different classes (Immunoglobulin (Ig)G, IgM, and IgA). Understanding the connection between nAbs and protein pathology could provide valuable insight into disease mechanisms and identify potential targets for therapeutic treatment.

Materials and methods

Demographics

A total of 235 plasma samples were collected from three different biobanks for this study (Table 1 and 2). (1) The samples included 38 PD and 15 DLB patients samples, and 29 control samples from the Bispebjerg Movement Disorder Biobank (BMDDB) at the Department of Neurology, Bispebjerg-Frederiksberg Hospital, Copenhagen University Hospital, Denmark (2) 69 AD and 31 DLB patient samples were obtained from the Danish Dementia BioBank (DDBB), Rigshospitalet, Copenhagen University Hospital, Denmark, and (3) 12 PD patients and 41 controls from the research-biobank at the Centre for Neuroscience and Stereology, Bispebjerg-Frederiksberg Hospital, Copenhagen University Hospital, Denmark. Only cases that met the international criteria for probable disease were included in the study [25–28]. The healthy control individuals had no central nervous system conditions, immunological disorders, or ongoing immunomodulatory treatment. All participants provided written consent for inclusion in the biobanks, adhering to the World Medical Association Declaration of Helsinki.

Syn/A /tau competition electrochemiluminescence immunoassay (ECLIA)

The affinity of anti- Syn/A /tau nAbs was assessed based on a competitive antigen-antibody reaction, whereby increasing antigen concentrations in the fluid phase facilitated distinguishable repertoires of high-affinity and low-affinity antibody fractions, previously developed

in-house [18]. In this study the assay was adapted and optimized for A and tau. In brief, 96-well mesoscale discovery (MSD) plates were coated overnight at 4 °C with antigens ( Syn: 20 ng/mL (rPeptide, #S-1001), standard small spot MSD plate (MSD, #L45XA); A<sub>1–42</sub>: 1 µg/mL (rPeptide, #A-1002), standard small spot MSD plate (MSD, #L45XA); tau: 1 ng/mL (rPeptide, #T-1001), high bind plate (MSD, #L15XB)) in ice-cold 0.1 M carbonate buffer, pH 8.5 (Sigma Aldrich, #C3041). Next, the plates were blocked for 1 h at 800 rpm ( Syn: PBS+BSA 3% (Sigma Aldrich, #05482); A : Intercept™ Blocking Buffer in PBS (LI-COR, #927-90001); tau: ROTI®Block1X (Carl Roth, #A151). Meanwhile, plasma samples were diluted ( Syn: 1:200; A /tau: 1:100) in PBS+BSA-0.1% (Sigma Aldrich, #05482) and preincubated with the antigen ( Syn: 1000 nM/2 nM/0.2 nM, 0 nM; A : 600 nM/6 nM/0.06 nM/0.0006 nM/0 nM; tau: 100 nM/1 nM/0.01 nM/0 nM) for 1 h before adding onto a newly washed antigen-coated plate (5 times with PBS+0.05%-Tween-20 (Sigma Aldrich, #P7949)) and incubated for 1 h at 800 rpm. After an additional washing step (5 times with PBS+0.05%-Tween-20 (Sigma Aldrich, #P7949)), SULFO-tag goat anti-human (1:10,000; MSD, #R32AJ-1) in PBS+BSA-0.1% (Sigma Aldrich, #05482) was added and eventually incubated for 1 h at 800 rpm. Finally, the plate was washed (5 times with PBS+0.05%-Tween-20) and Read Buffer T (1:2 (MSD, #R92TC)) was added upon reading the plate immediately before the MSD Sector Imager/QuickPlex SQ 120 Reader (MSD, LLC, USA). The percentage of max binding for each sample and pool was calculated as follows:

Table 1 Demographic and clinical data

	AD (N= 69)	PD (N= 50)	DLB (N= 46)	NC (N= 70)	p values
Age [years]	70.4 (8.1) [51–89]	68.4 (7.4) [52–84]	72.8 (6.4) [56–88]	71.4 (9.1) [52–90]	0.020*
Sex (M/F)	35/34	24/26	33/13	31/39	0.027**
Age at onset [years]	68.0 (8.6) [48–88]; 81%	61.4 (8.3) [44–78]	69.6 (8.8) [38–87]; 98%	-	0.267#
MMSE	24.1 (3.9) [12–30]	-	25.8 (4.0) [16–30]; 67%	-	0.016#
H&Y	-	2.3 (0.9) [1–5]; 72%	2.4 (0.9) [1–3]; 32%	-	0.461#
Disease Duration [years]	2.2 (1.2) [0.5–5]; 81%	7.0 (4.1) [0–15]	3.2 (3.3) [0–21]; 98%	-	<0.001%
Biobank	DDBB	BMDDB; CNS-lab	DDBB; BMDDB	BMDDB; CNS-lab	

\*: Welch ANOVA. \*\*: chi-squared test. #: Mann–Whitney test. %: Kruskal–wallis test. MMSE: Mini Mental State Examination (MMSE); H&Y: Hoehn & Yahr (7-scale); M: Male; F: Female; DDBB: Danish Dementia BioBank; BMDDB: Bispebjerg Movement Disorder Biobank; CNS-lab: Centre for Neuroscience and Stereology. \$: PD vs. DLB (collectively), *p* < 0.05

% of max binding

$$= \frac{(ECLIA_{\text{sample OD}} - ECLIA_{\text{OD at 1000 nM competitor (0\% binding)}})}{ECLIA_{\text{OD at 0 nM competitor (100\% binding)}}} \times 100$$

IgG, IgM and IgA anti- Syn/A /tau measurements

Total levels of anti- Syn/A /tau nAbs were measured by indirect ELISA as previously described [19, 20, 29] with few adjustments. In brief, 96-well polystyrene microtiter plates (Nunc MaxiSorp® at-bottom) were coated overnight with antigens ( Syn: 5 µg/mL (rPeptide, #S-1001-2); A<sub>1–42</sub>: 5 µg/mL (rPeptide, A-1002); tau: 0.5 µg/mL (rPeptide, T-1001)) in ice-cold 0.1 M carbonate buffer, pH 8.5 (Sigma Aldrich, #C3041). The plates were then emptied and blocked for 2 h at RT with PBS+BSA-3% (Sigma Aldrich, #05482)+Tergitol-0.1% (Sigma Aldrich #NP40S). Following a subsequent washing cycle of 5 times with PBS+0.05%-Tween-20 (Sigma Aldrich, #P7949), plasma samples were diluted (1:50 for anti- Syn/A /tau IgA and 1:100 for anti- Syn/A /tau IgG/IgM) in dilution buffer (PBS+0.1%BSA+0.05% + Tween-20) and incubated for 1 h at RT. After another washing cycle (5 times with PBS+0.05%-Tween-20), secondary HRP-conjugated anti-Ig antibodies (anti-IgG

**Table 2** Statistical comparison of nAb activities and levels for Syn, A and tau

Antigen	Model statistics			Groups		Sex	Age	DLB-AD	PD-AD	NC-AD	PD-DLB	NC-DLB	NC-PD
	p value	R <sup>2</sup>	F-stat	p value	p value	p value	p value	p value for multiple comparison*					
Syn	0.116	0.024	F(5,161) = 1.80	0.053	0.296	0.976							
	<b>0.033</b>	0.047	F(5,147) = 2.51	<b>0.016</b>	0.317	0.120							
	<b>1.9E-07</b>	0.150	F(5,211) = 8.62	<b>1.2E-07</b>	0.430	0.686							
	<b>6.1E-07</b>	0.139	F(5,213) = 8.01	<b>5.8E-07</b>	0.286	0.260							
	0.334	0.004	F(5,207) = 1.15	0.225	0.168	0.588							
A	p value	R <sup>2</sup>	F-stat	p value	p value	p value							
	0.033	0.040	F(5,177) = 2.49	<b>0.022</b>	0.450	0.225							
	0.089	0.025	F(5,177) = 1.95	0.068	0.337	0.915							
	<b>0.038</b>	0.030	F(5,224) = 2.41	<b>0.035</b>	0.318	0.167							
	<b>0.020</b>	0.037	F(5,222) = 2.76	<b>0.026</b>	0.174	0.456							
Tau	<b>0.049</b>	0.026	F(5,227) = 2.26	<b>0.015</b>	0.770	0.456							
	p value	R <sup>2</sup>	F-stat	p value	p value	p value							
	0.817	0.015	F(5,180) = 0.45	0.937	0.267	0.587							
	0.109	0.023	F(5,174) = 1.83	0.082	0.155	0.916							
	<b>0.033</b>	0.031	F(5,226) = 2.47	<b>0.010</b>	0.533	0.756							
IgM	<b>0.020</b>	0.036	F(5,227) = 2.74	<b>0.036</b>	0.057	0.575							
	0.557	0.005	F(5,225) = 0.79	0.871	0.078	0.891							

\*: Multiple comparison modulated for covariables (sex and age)

(1:20,000; Abcam, #ab98624), biotin-conjugated anti-IgM (1:5,000; Sigma Aldrich, #B1265), and anti-IgA (1:1,000 for Syn/A ; 1:2000 for tau; Cermo Fisher Scientific, #PA1-74395) were diluted in dilution buffer, added to the plates and incubated for 2 h at RT. An additional step was carried out for the biotin-conjugated IgM antibody, with streptavidin–peroxidase (1:10,000; Sigma Aldrich, #S5512) for 30 min at RT. Next, the plates were washed once again (5 times with PBS+0.05%-Tween-20), and tetramethylbenzidine (TMB) Liquid Peroxidase Substrate (Sigma Aldrich, #T8665) was added for 30 min in the dark at RT prior to reaction termination by the addition of 0.5 N sulfuric acid (Sigma Aldrich, #319570). Finally, the absorbance was measured at 450 nm and 620 nm on a MultiSkan™ FC Microplate Reader (Cermo Fisher Scientific, USA). All data were normalized to positive controls on each individual plate. Positive controls consisted of pooled plasma samples, from controls and patients, added to each plate to account for plate-to-plate variability.

#### Statistical analyses

For demographic group comparison, we applied Welch ANOVA followed by the Games-Howell test for multiple comparisons for age since the data was normally distributed but has difference in variances, the chi-squared test for sex, the Mann Whitney U test for age at onset, MMSE, Hoehn & Yahr and disease duration. Outliers were removed from analyses using ROUT with false discovery rate (FDR),  $Q=1\%$ . Normality was assessed using the D'Agostino Omnibus test. For group comparison, we applied multiple linear regression modeling including covariates age and sex, since small discrepancies between groups were observed, using ANOVA from the *car* package [30]. For multiple comparisons, the *glht* and *mcp* functions from the *multcomp* package [31] were applied using Tukey's range test. Correlations between measured outcomes and clinical data were assessed using Spearman's rank-order correlation. Spearman's correlation matrices were constructed using *corrplot* package [32]. Data were analyzed using R v. 3.5.2 [33] and GraphPad Prism 9.4.1 (GraphPad Software Inc., USA).

## Results

### Anti- Syn/A /tau high-affinity nAbs in AD, DLB and PD patients

To assess the functionality of nAbs and their capacity to form stable immunocomplexes across various diseases, we analyzed the binding affinity of anti- Syn, -A , and -tau IgG nAbs in patients with AD, DLB and PD, as well in control individuals. To perform these analyses, we utilized our well-characterized competition assay with minor adjustments [19, 20, 34]. Based on initial competition curves obtained from a subset of 10 randomly

selected age- and sex-matched patients and control individuals (Fig. 1A, D and G), we chose two different conditions to formally evaluate the high-affinity nAb repertoire.

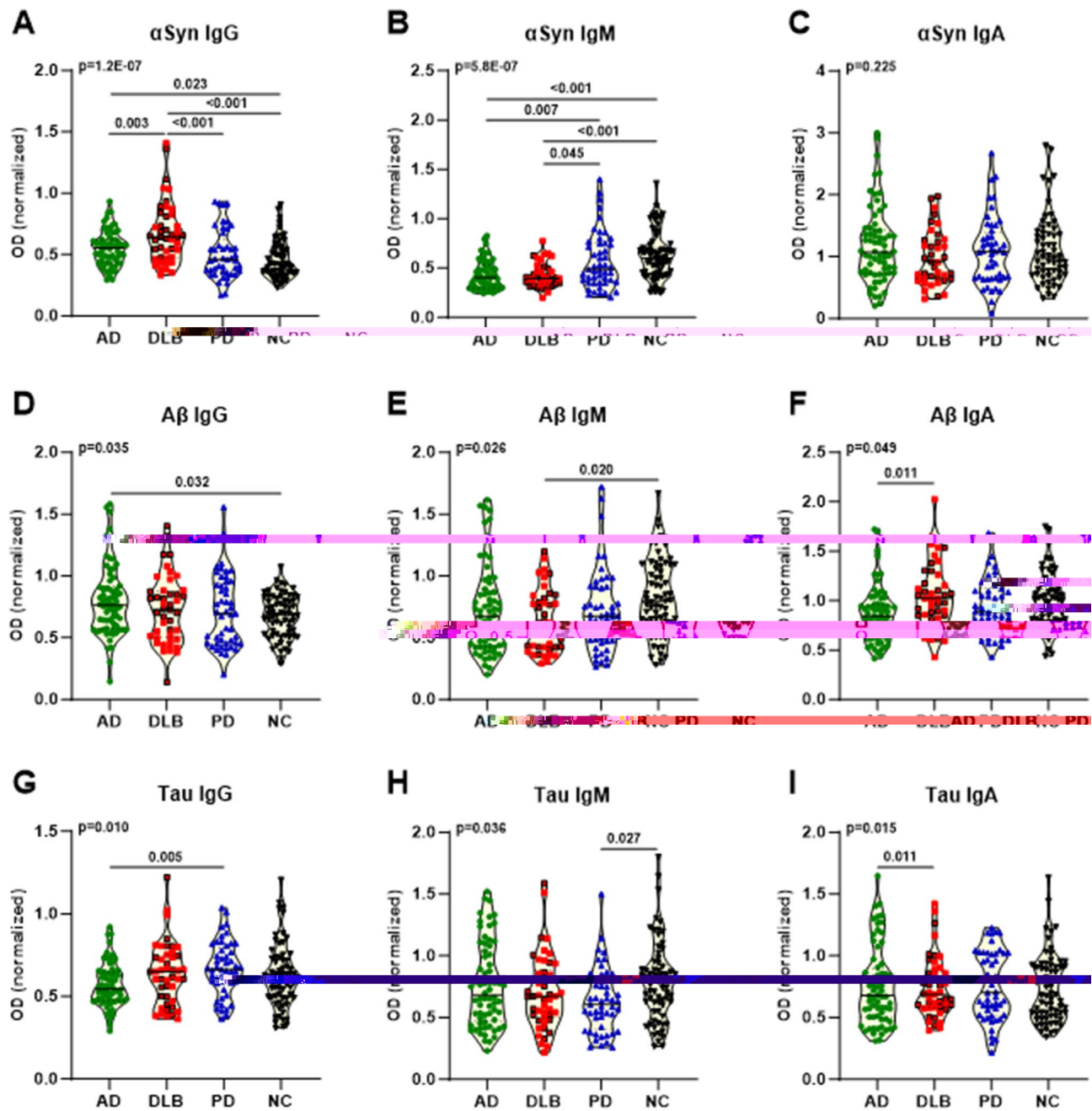
The analysis of individual samples revealed notable differences in the high-affinity repertoire of anti- Syn and anti-A IgG nAbs. When exposed to 0.2 nM free Syn, PD patients ( $p=0.045$ ) and AD patients ( $p=0.037$ ) (Fig. 1B; Table 3) exhibited a significantly reduced repertoire of high-affinity anti- Syn IgG compared to controls. Additionally, when exposed to 0.6 nM free A , AD patients only demonstrated significantly lower amounts of high-affinity anti-A IgG compared to controls. No differences in tau affinity reactivity were observed between groups.

### Anti- Syn/A /tau IgG, IgM and IgA nAbs in AD, DLB and PD patients

To explore the reactivity of different antibody classes in the immune system, namely, IgG, IgM, and IgA, toward Syn, A , and tau in patients with AD, DLB, PD, and controls, we conducted indirect ELISA analyses. Exploring the repertoires of anti- Syn IgG, IgM, and IgA nAbs, we observed that AD and DLB patients exhibited significantly higher levels of anti- Syn IgG than controls (AD:  $p=0.023$ ; DLB:  $p<0.001$ ) (Fig. 2A; Table 3). More significantly, DLB patients exhibited increased levels of anti- Syn IgG compared to both AD ( $p=0.003$ ) and PD patients ( $p<0.001$ ) (Fig. 2A; Table 3). In terms of anti- Syn IgM, both AD and DLB patients demonstrated reduced levels compared to PD (AD:  $p=0.007$ ; DLB:  $p=0.045$ ) and controls (AD:  $p$



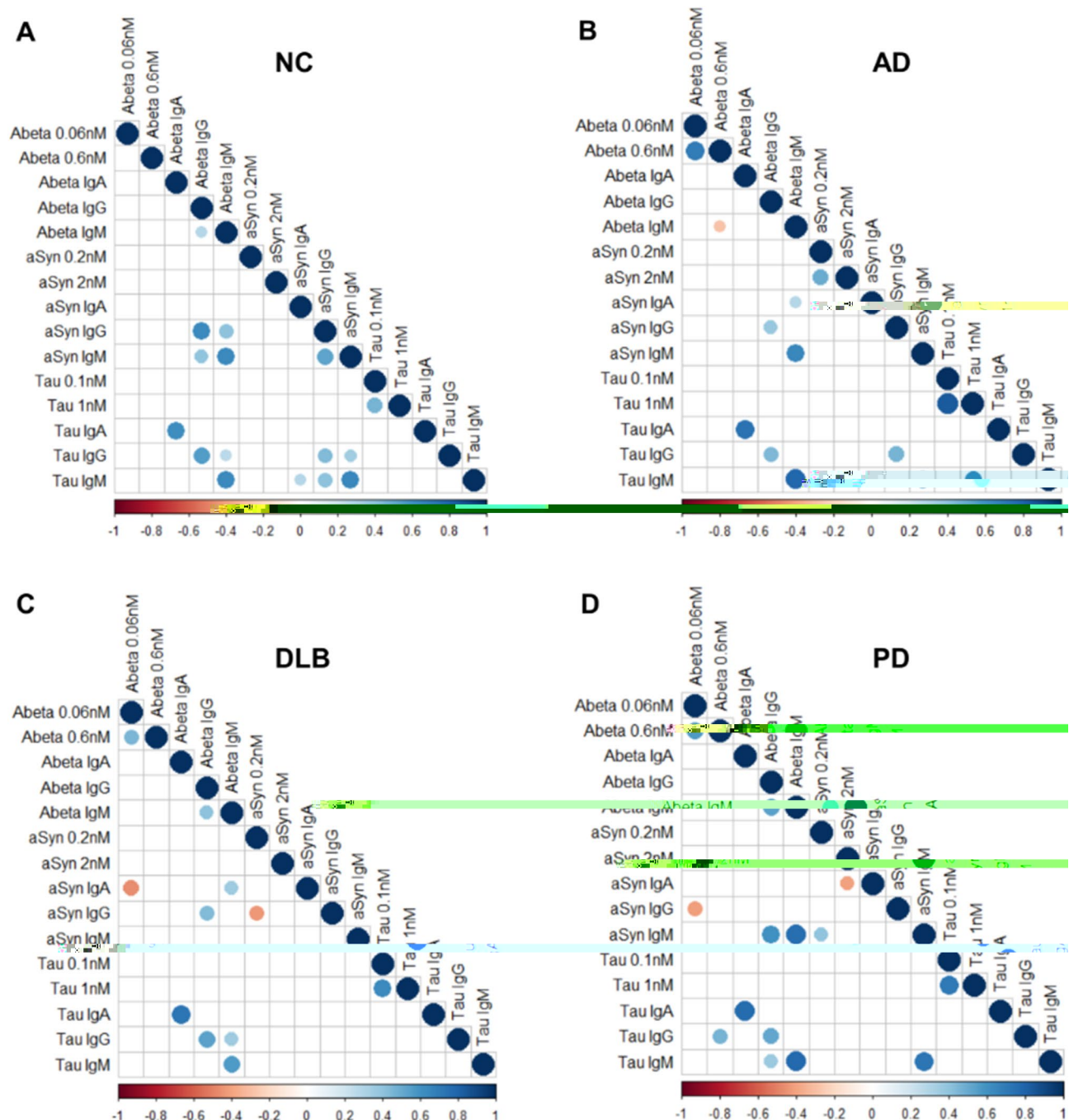




**Fig. 2** Relative anti- $\alpha$ -Syn (A-C),  $A\beta$  (D-F) and  $\tau$ -tau (G-I) IgG, IgM and IgA nAb levels in AD, DLB and PD patients as well as controls. Data are presented as normalized optical (normalized to positive controls on each individual plate) densities in truncated violin plots with median (horizontal line). Group comparisons were performed by applying multiple linear regression models including covariates age and sex and post hoc multiple comparison testing using Tukey's range test. Statistically significant p-values ( $<0.5$ ) are depicted

in Fig. 3A and Table S3. This was with the exception of the anti- $A\beta$  IgG versus anti- $\tau$  IgM relationship ( $r=0.230$ ,  $p=0.059$ ). Notably, strong correlations persisted across all four examined groups (AD, DLB, PD and controls) for the anti- $\tau$  IgA, IgG, IgM and the anti- $A\beta$  IgA, IgG, IgM (Fig. 3A-D, Table S1-4), respectively. Furthermore, a positive correlation was found between anti- $A\beta$  IgM and IgG ( $r=0.284$ ,  $p=0.019$ ), and similarly between anti- $\alpha$ -Syn IgM

and IgG ( $r=0.525$ ,  $p=1.2E-05$ ). A positive correlation was also observed between the activity of anti- $\tau$  IgG for two concentrations of free  $\tau$  ( $r=0.466$ ,  $p=4.3E-04$ ). Interestingly, in controls, no correlation was between the two activity measures for anti- $A\beta$  IgGs ( $r=-0.082$ ,  $p=0.601$ ), contrasting with the positive correlations observed in AD ( $r=0.693$ ,  $p=1.2E-08$ ), DLB ( $r=0.456$ ,  $p=0.009$ ) and PD ( $r=0.567$ ,  $p=1.7E-04$ ) patients.



**Fig. 3** Spearman's correlation matrix of the anti-Syn/A, tau, and IgG, IgM, and IgA levels. A: amyloid-beta, Ig: immunoglobulin, Syn: alpha-synuclein. Scalebar ranging from Spearman's  $r = -1$  (red, negative correlation) to  $+1$  (blue, positive correlation). Only significant correlations are shown. P-values  $< 0.05$  were considered significant

In AD patients, while many correlations remained (Fig. 3B, Table S4), there were a few exceptions, in addition to the previously described A $\beta$  high-a $\beta$  correlation. These exceptions included four positive correlations: between high-a $\beta$  anti-Syn 0.2 nM and anti-Syn 2 nM ( $r=0.492$ ,  $p=4.4E-04$ ), anti-Syn IgA and anti-A $\beta$  IgM ( $r=0.289$ ,  $p=0.024$ ), high a $\beta$  anti-tau 0.1 nM

and anti-tau 1 nM ( $r=0.835$ ,  $p=3.2E-13$ ) and anti-tau IgG versus anti-tau IgM ( $r=0.259$ ,  $p=0.034$ ). Additionally, a negative correlation was noted between anti-A $\beta$  IgM and high-a $\beta$  anti-A $\beta$  IgGs ( $r=-0.285$ ,  $p=0.047$ ).

In case of DLB patients compared to healthy controls, eight positive correlations associated with anti-Syn IgA, IgG, and IgM versus anti-A $\beta$ , and anti-tau IgG, and IgM



were eliminated (Fig. 3C, Table S5). Moreover, anti-Syn IgA and IgG showed negative correlations with high-affinity anti-A $\beta$  IgG ( $r=-0.462$ ,  $p=0.012$ ) and high-affinity anti-Syn IgG ( $r=-0.430$ ,  $p=0.018$ ), respectively. Additionally, the anti-Syn IgG and anti-Syn IgM interrelationship was ablated ( $r=-0.117$ ,  $p=0.497$ ).

In PD patients, compared to healthy controls, there was a notable impact on the correlations involving anti-Syn and anti-tau IgGs (Fig. 3D, Table S6). Specifically, anti-Syn IgG exhibited a negative correlation with high-affinity anti-A $\beta$  IgGs ( $r=-0.418$ ,  $p=0.013$ ), diverging from the previously observed positive correlations with anti-A $\beta$  IgG, anti-tau IgG and IgM in controls, which were no longer present. Furthermore, anti-Syn IgA was found to be negatively correlated with high-affinity anti-Syn IgGs ( $r=-0.403$ ,  $p=0.020$ ). Positive interrelationships were observed between anti-tau IgG and high-affinity anti-A $\beta$  IgGs ( $r=0.460$ ,  $p=0.004$ ), anti-tau IgM and anti-A $\beta$  IgGs ( $r=0.369$ ,  $p=0.008$ ), and between anti-Syn IgM and high-affinity anti-Syn IgGs ( $r=0.394$ ,  $p=0.023$ ).

## Discussion

In the explorations of neurodegenerative diseases such as Alzheimer's disease (AD) and Parkinson's disease (PD), the role of immune clearance has emerged as a topic of significant interest. This stems from the historical precedent set by the discovery of nAbs against the A $\beta$  protein in AD as early as 2001 [35]. However, even with decades of research, the exact function, and implications of these nAbs remain a subject of debate. Our study offers a fresh perspective by examining the binding affinity of nAbs to essential proteins associated with AD, DLB and PD patients.

By employing our well-characterized competition assay [18–20, 34], we analyzed the high-affinity repertoire of nAbs against Syn, A $\beta$ , and tau in AD, DLB and PD patients, as well as healthy control subjects. We extended our prior findings in PD to include AD, demonstrating significantly reduced high-affinity anti-Syn IgG nAbs compared to controls, and further demonstrating reduced high-affinity A $\beta$  nAbs in AD compared to controls. Although the common paradigm separates

Syn pathology into PD and A $\beta$  pathology into AD, the broader landscape of neurodegenerative disorders reveals that between 20 and 40% of all AD cases show pathological Syn accumulation in the brain [36, 37]. In addition to co-occurrence in pathology, several mechanisms have been proposed for the role of Syn involvement in AD.

Syn interacts with A $\beta$  and tau, promoting their aggregation and toxicity and contributing to the complexity and severity of neurodegenerative processes [38]. It is likely a key effector in neurotransmitter release and synaptic function, which have been shown to be compromised in AD [39]. Additionally, emerging evidence suggests that

Syn acts as a culprit in neuroinflammatory processes and contributes to activating microglia, as observed both in PD and potentially AD [40, 41]. Although these potential links are intriguing, we can only speculate which processes are present in this study's AD patients and whether they have Syn co-pathology, which could explain their reduced functionality of anti-Syn nAbs.

Earlier research has emphasized the ability of IgG nAbs to regulate inflammation and to facilitate the clearance of neurotoxic aggregates [42]. The presence of disease-specific IgG nAbs targeting pathological proteins such as Syn, A $\beta$  and tau suggests an intricate interplay between the immune system and the pathogenesis of neurodegenerative disorders [11, 43]. Although the intracellular location of Syn and tau aggregates makes it unlikely that nAbs penetrate the cell membrane and clear the aggregates intracellularly, they are more likely to scavenge the extracellular space, clearing material transmitting between cells and aiding in the degradation of cellular material after apoptosis. Considering that nAbs play a crucial role in regulating immune clearance mechanisms, any abnormalities in the nAb profile could potentially exacerbate the pathogenesis of neurological conditions. This observation was recently established by the success of two different passive immunization strategies in treating AD. The effectiveness of lecanemab and donanemab in AD patients [23, 24] manifests the importance of functional regulation of key pathological proteins. Passive immunization seems to have the potential to bolster compromised immune system functions in neurodegenerative diseases. To date, no conclusive study of passive immunization in PD patients has proven successful. However, promising secondary outcomes were recently achieved in the AMULET study, a Phase II passive immunization trial in MSA patients. This trial was based on the hypothesis that the treatment would not clear existing aggregates but would instead slow down or halt the spread and seeding of aSyn to other cells [44]. These results, taken together with the lecanemab and donanemab trials in AD, further imply that intravenously administered antibodies can partially reach the brain, consistent with previous findings showing differences in nAbs in CSF samples from PD and AD, which also correlate with plasma levels [19, 35, 45–48]. Several factors, including the absence of precise antibody candidates, defined pathological hallmarks, and challenges in enrolling patients with varying disease durations or early in the disease onset into the trials, could contribute to this lack of success. Furthermore, the fact that the main pathological processes are different would be the most obvious reason. This contrast may suggest distinct nAb-associated disease mechanisms or pathological responses between AD and PD. Confirming this, recent studies offer seemingly paradoxical perspectives on the role of

B cells in disease pathogenesis. For instance, Scott et al. [49], posited that B cells play a protective role in a PD model, whereas a 2021 study hinted at their pathological role in an AD model [50].

Plasma IgM levels were altered in AD, DLB and PD patients. Interestingly, we observed that AD and DLB patients had reduced anti-Syn IgM titers compared to PD patients and controls. DLB patients also presented reduced anti-A  $\beta$  IgM titers. IgM nAbs bind to disease-specific proteins and influence the aggregation of these proteins [42, 51]. The pentameric structure of IgM endows it with multivalency, allowing it to bind to multiple copies of proteins, such as Syn, A  $\beta$  and tau [52, 53].

This property could facilitate more efficient clearance. IgM's relatively unspecific yet rapid immune response makes it an essential component of the innate immune system. In neurodegenerative diseases, this could mean that IgM acts as an early responder to neural inflammation or protein aggregation, is secreted quickly, and plays a role in complement activation. Harvesting these properties has been proposed as a passive immunization strategy, with the scFv-Fc format allowing for multimerization into pentameric structures, improving the binding and functionality of the antibodies [54]. On the other hand, reduced antigen-specific IgM nAbs have been observed leading to increased IgG nAbs towards self-antigens [55], possibly explaining the subsequent increased anti-aSyn IgG nAbs in AD and DLB. Further understanding the multifaceted roles of IgM in AD, DLB and PD could offer novel insights into neurodegenerative disease pathogenesis and explore its potential as a therapeutic approach.

Finally, this study is the first to explore the relevance of pathology-related IgA compartments in neurodegenerative diseases. Our data suggest a positive correlation between the Hoehn and Yahr (H & Y) scale and anti-Syn IgA nAbs in individual plasma samples from PD patients (Fig. S1D). IgAs are primarily known for their role in mucosal immunity, such as lining the gastrointestinal tract, as well as other openings inside the nose and mouth [56]. In addition to their localization at mucosal sites, IgAs are also found in the circulatory system [56]. One of their critical functions is to maintain harmonious homeostasis between the microbiota and the host's immune response [57]. The role of the gut-brain axis, especially in PD, has attracted increasing interest. One intriguing observation is the identification of Syn pathology at the gut's mucosal lining in PD patients [58, 59]. Furthermore, specific infections such as *Helicobacter pylori* have been implicated in PD [60], and urinary tract infections have been associated with the atypical parkinsonian disorder, MSA [61], possibly triggering Syn misfolding, which can spread to the brain via the vagus nerve [62, 63]. Specifically, related to IgA in the context of the gut-brain axis, recent studies found that the IgA to IgM/

IgD ratio was nearly 2-fold increased in PD patients [64]. Moreover, IgA-producing plasma cells are not only present in the meningeal venous sinuses but also associated with increases during aging and after an intestinal barrier breach [65]. B-cell receptor sequencing further identified these cells as originating from the intestine [65]. In the realm of AD, recent research has reported elevated IgA levels in the plasma and brain tissue of APOE-4 non-carriers, establishing intracerebral transfer of IgAs [17].

These and our discoveries indicate a possible connection between gut-specific IgA responses, healthy aging and the onset or progression of PD and related neurodegenerative conditions.

As neurodegenerative pathologies progress and redistribute, distinct changes in nAb function and concentration emerge, with variability across diseases. In AD, A  $\beta$  pathology accumulates prior to tau pathology [66], and both occur before clinical symptoms manifest. This sequence implies that nAb-response dynamics may differ between A  $\beta$  and tau. Similarly, in PD and DLB, Syn accumulation is an early event, possibly starting many years prior to disease onset [67, 68], which may explain the heightened anti-Syn response observed in early and prodromal PD stages [20, 45] and in idiopathic REM sleep behavior disorder (iRBD) patients [49].

The precise interactions among nAb affinity, disease pathology, and immune regulation remain complex and incompletely understood. However, differential responses in nAbs targeting key neurodegenerative proteins suggest intricate interplay between nAb affinity and disease processes. One of the most striking observations is the lack of correlation between the affinity of anti-A  $\beta$  IgGs in controls (Fig. 3A) and the strong positive correlations in AD and PD patients (Fig. 3B and D). One hypothesis posits that nAb-producing B cells are pre-existing and slight antigenic pushes, drives generation of IgGs and IgAs by differentiating into plasma cells. This process enables hypermutation and class-switching, as suggested by Reynefeld et al. 2020 [69]. The further persistence of IgG and IgM correlations, particular in controls, suggest that these nAbs may serve protective and regulatory roles, which become disrupted in disease. The reduction in these correlations in patients implies an overall breakdown in the immune system's ability to coordinate the recognition and removal of misfolded proteins. On the other hand, the disruptions could merely be driven by the chronic presence of A  $\beta$  plaques and increased A  $\beta$  1–42 levels in the brains of AD patients and Syn Lewy body aggregates in PD patients, or both in DLB patients and that these high-affinity nAbs are sequestered in the brain, which could contribute to their lower levels in peripheral circulation. However, the non-changes in relation to tau and the absence of correlation between affinity and disease duration, cognitive impairment and motor

disability talks against it. Only in DLB patient, a significant decrease was observed in relation to anti-Syn IgG affinity and levels, suggesting that both affinity and levels are decreased during disease progression (Table S2), suggesting a link between development of Syn accumulation and anti-Syn nAbs. This needs to be explored further, in brain and body, and although, we can only speculate at this stage, the breakdown of interrelationships between different nAbs across disease groups support the theory that chronic neuroinflammation and immune dysregulation are shared features across neurodegenerative diseases.

The present study, while insightful, has several limitations. First, its cross-sectional design captures antibody dynamics at a single time point, limiting understanding of their progression over time. Longitudinal studies are needed to clarify these changes, potentially in the prodromal stages. Second, peripheral blood measurements may not fully reflect central nervous system (CNS) pathology, as blood-brain barrier integrity and antibody sequestration in the brain were not directly assessed. The study also focuses on A $\beta$ , tau, and Syn autoantibodies, potentially overlooking other disease-relevant proteins.

The mechanisms behind the observed antibody variations, particularly the role of nAbs in disease progression, remain speculative. Finally, the study did not explore other immune pathways or antibody subclasses that may play critical roles in neurodegeneration. Future research addressing these limitations is necessary for deeper mechanistic understanding.

Conclusions

The multifaceted nature of neurodegenerative diseases is reflected in the aberrant levels of nAbs and their classes. While the utility of nAbs as diagnostic biomarkers remains a subject of ongoing debate, we argue that their inherent variability among groups and individuals limits their effectiveness in this role. However, their importance in elucidating disease mechanisms should not be underestimated, and they may prove valuable in identifying subgroups within the disease spectrum. Our study provides evidence of a dysfunctional immune system in neurodegenerative diseases, suspected to impair the endogenous clearing mechanism of pathological proteins, namely A $\beta$ , Syn and tau. This suggests a relationship between disease-specific immunoglobulins and pathogenesis, although the specific nature of this relationship has yet to be clearly defined.

Abbreviations

AD	Alzheimer's disease
A $\beta$	Amyloid-beta
PD	Parkinson's disease
aSyn	Alpha-synuclein
DLB	Dementia with Lewy bodies
nAbs	Naturally occurring autoantibodies

Ig	Immunoglobulin
MSA	Multiple system atrophy
CSF	Cerebrospinal fluid
PDD	Parkinson's disease dementia
VaD	Vascular dementia
FTD	Frontotemporal dementia
MSA	Multiple system atrophy
PSP	Progressive supranuclear palsy
LRKK2	Leucine-rich repeat kinase 2
bvFTD	Behavior-variant FTD
SCA	Spinocerebellar ataxia
sAD	suspected AD
ADRD	AD-related dementia
NDs	Neurodegenerative disorders
NPH	Normal pressure hydrocephalus
RBD	REM-sleep behavior disorder
ApoE	Apolipoprotein E
E4	Epsilon 4
MCI	Mild cognitive impairment
OND	Other neurological diseases
IC	Inflammatory controls
ALS	Amyotrophic lateral sclerosis
CJD	Creutzfeld-Jakob disease
CBD	Corticobasal degeneration
WE	Wernicke encephalopathy
BMDB	Bispebjerg Movement Disorder Biobank
DDBB	Danish Dementia BioBank
CNS-lab	Centre for Neuroscience and Stereology
ECLIA	Electrochemiluminescence immunoassay
MSD	Mesoscale Discovery
PBS	Phosphate Buffered Saline
BSA	Bovine Serum Albumin
ELISA	Enzyme-linked Immunosorbent assay
TMB	3,3', 5,5' tetramethylbenzidine dihydrochloride
FDR	False Discovery Rater
nM	Nanomolar
H&Y	Hoehn & Yahr staging (7-stage)
NC	Non-neurological controls

Supplementary Information

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Supplementary Material 1

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Author contributions

LK and KD contributed equally to this work and conducted the majority of experiments and analyzed data and authored the manuscript. AHS, KW, LS, SB, A-MH, AL, SGH, GW contributed to patient material and clinical assessments. FP, JAR, RD and SA: contributed by allocating resources, and interpreting data. TB: contributed to the design, conception and execution of experiments and interpreted data. JF: secured funding, contributed to conception, design and execution of experiments, interpreted data and authored the manuscript. All authors reviewed and revised the manuscript.

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Data availability

No datasets were generated or analysed during the current study.

## Declarations

### Ethics approval

This project was approved by the Danish National Committee on Health Research Ethics, Copenhagen Regional Area (j.no.: H-15016232) and the Danish Data Protection Agency (j.no.: P-2020-937). All participants gave written consent for inclusion in the biobanks according to the World Medical Association Declaration of Helsinki.

### Competing interests

The authors declare no competing interests.

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