Relevance of plasma biomarkers to pathologies in Alzheimer’s disease, Parkinson’s disease and frontotemporal dementia

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Amyloid plaques and tau tangles are pathological hallmarks of Alzheimer’s disease (AD). Parkinson’s disease (PD) results from the accumulation of α-synuclein. TAR DNA-binding protein (TDP-43) and total tau protein (T-Tau) play roles in FTD pathology. All of the pathological evidence was found in the biopsy. However, it is impossible to perform Stein examinations in clinical practice. Assays of biomarkers in plasma would be convenient. It would be better to investigate the combinations of various biomarkers in AD, PD and FTD. Ninety-one subjects without neurodegenerative diseases, 76 patients with amnesic mild cognitive impairment (aMCI) or AD dementia, combined as AD family, were enrolled. One hundred and nine PD patients with normal cognition (PD-NC) or dementia (PDD), combined as PD family, were enrolled. Twenty-five FTD patients were enrolled for assays of plasma amyloid β 1–40 (Aβ1–40), Aβ1–42, T-Tau, α-synuclein and TDP-43 using immunomagnetic reduction (IMR). The results show that Aβs and T-Tau are major domains in AD family. α-synuclein is highly dominant in PD family. FTD is closely associated with TDP-43 and T-Tau. The dominant plasma biomarkers in AD family, PD family and FTD are consistent with pathology. This implies that plasma biomarkers are promising for precise and differential assessments of AD, PD and FTD in clinical practice.

In addition to cognitive and behavioral inspections, the demand for biological examinations is strongly increasing for neurodegenerative diseases in clinics1–3. From the perspective of pathogenesis, neurodegenerative diseases result from the misfolding of specific proteins in the brain4–6. Pathological studies on animal biopsy or human autopsy demonstrate that the accumulation of these misfolded proteins causes neuronal damage, cognitive impairment and behavioral disorders2–11. For example, amyloid plaques and tau-protein tangles are pathological hallmarks of Alzheimer’s disease (AD)12–14. Lewy bodies composed of α-synuclein were found in the brain steins of patients with Parkinson’s disease (PD)15,16. TAR DNA-binding protein (TDP-43) and total tau protein (T-Tau) play roles in the pathology of patients with frontotemporal dementia (FTD)17,18. To observe the pathological evidence of neurodegenerative diseases, positron emission tomography (PET) scans with the aid of tracers to specifically label the accumulated proteins in brains have been developed or are under development19–22. Unfortunately, to date, only tracers for amyloid PET are available in clinical practice. It is impossible to use PET scans to clarify the pathologies for PD or FTD in clinics. Even for amyloid PET scans, the cost issue makes it difficult to perform in routine practice. Assays of these biomarkers in cerebrospinal fluid (CSF) are alternative inspections for neurodegenerative diseases23–26. Several research results show high correlations between CSF biomarkers

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and pathology or clinical diagnosis. However, due to side effects, lumbar puncture is not popularly used in clinical practice. Thus, biological examinations with biomarkers face severe bottlenecks.

With the successful development of ultrasensitive assay technologies in the early 2000s, the precise detection of biomarkers in human plasma instead of CSF has become feasible. Many researchers and neurologists have been interested in studies on plasma Aβ, T-Tau, TDP-43 and other biomarkers. Most studies have concentrated on the discrimination between individual neurodegenerative diseases and normal controls using plasma biomarkers. It was reported that plasma Aβ1–42 and T-Tau are able to differentiate AD from normal controls. PD patients show significantly higher levels of plasma α-synuclein than normal controls. Relatively higher levels of plasma TDP-43 in FTD were reported. However, comprehensive studies on the effects of Aβs, T-Tau, α-synuclein and TDP-43 on AD, PD and FTD are rare.

In this work, normal controls and patients with either AD, PD or FTD were enrolled. Plasma Aβ1–40, Aβ1–42, T-Tau, α-synuclein and TDP-43 were assayed for each participant using immunomagnetic reduction (IMR). Since 2011, the results of assaying plasma Aβ1–40, Aβ1–42, T-Tau, α-synuclein and TDP-43 have been reported by independent groups over the world. In addition to ultra-high sensitivity and specificity, the published results evidenced the high consistencies in measured plasma biomarkers with CSF biomarkers, plasma Aβ1–42/ Aβ1–40 with amyloid PET, plasma T-Tau with magnetic resonance MRI, plasma Aβ1–42 x T-Tau, α-synuclein, and TDP-43 with clinical diagnosis of AD, PD and FTD, respectively. With these clinical validations, IMR kits have been registered with CE IVD and approved as an in-vitro medical devices by Taiwan Food and Drug Administration. Hence, IMR is reliable to be used to explore the plasma Aβ1–40, Aβ1–42, T-Tau, α-synuclein and TDP-43. The contributions of each plasma biomarker to AD, PD and FTD were analyzed to clarify the role of each biomarker in these neurodegenerative diseases.

Methods
Enrollment of subjects. All subjects were enrolled at hospitals in Taiwan with approvals by the ethics committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA committees of every hospital. Each subject was identified as normal control (NC), Alzheimer’s disease (AD), Parkinson’s disease (PD) or frontotemporal dementia (FTD) according to the diagnostic guidelines of NIA-AA

Preparation of plasma samples. A 9 ml or 6 ml K3 EDTA lavender-top tube was used for blood draw with each enrolled subject, followed by centrifugation at 1500–2500 g at room temperature for 15 min. Plasma was collected and aliquoted into cryotubes (0.5 ml aliquots) and stored at −20 °C. The freezing of plasma should be performed within 4 h after blood draw. Each frozen plasma sample was placed in wet ice and then positioned at room temperature for IMR measurement.

Assays of biomarkers. IMR reagents (MF-AB0-0060, MF-AB2-0060, MF-TAU-0060, MF-ASC-0060, MF-TDP-0060, MagQu) were used to assay Aβ1–40, Aβ1–42, T-Tau, α-synuclein and TDP-43 in plasma with the aid of an IMR analyzer (XacPro-S, MagQu). There are two stages of quality controls for each batch of IMR measurement. The first stage is to calibrate the reading of the sensor of the analyzer. 120-μl, 10-mg-Fe/ml magnetic fluid is used as a calibrator for each detecting channel. The output signals of blank test (without calibrator) and calibrator test (with calibrator) should be within the acceptable range. In addition, the signal ratio of calibrator test to blank test should be higher than 20. The second stage is as follows. For each batch of IMR measurement for a given biomarker, control samples (several tens of pg/ml) with known concentrations are assayed together with tested samples. The deviations of measured concentrations of control samples should be less than 15%. Duplicated measurements were performed for each biomarker of a plasma sample. The averaged value of the duplicated measurements was reported.

Statistical method. The software, GraphPad Prism 6.01, was used to perform the data analysis. Ages, CDR, MMSE scores, H–Y stage, concentrations of biomarkers were presented as the means ± standard deviations for each enrolled group. MMSE scores and concentrations of biomarkers were compared between two enrolled groups, e.g. AD family versus NC, using a t test, and p values were determined. The value of 0.05 for p was a criterion to determine the significance.
Significantly higher levels of plasma T-Tau were found in the AD family (34.51 ± 9.91 pg/ml; \( p < 0.001 \)) and FTD (41.53 ± 20.51 pg/ml; \( p < 0.001 \)) than in the NC group (17.46 ± 9.28 pg/ml; \( p < 0.01 \)). T-Tau showed a significantly positive correlation with the dementia-rank severity (CDR) and the cognitive impairment (MMSE) for all groups. There was no significant difference in the plasma levels of T-Tau between the AD family and FTD. The effect sizes of T-Tau for all groups are listed in Table 2, except for FTD. The effect size of T-Tau for the AD family was calculated as 34.51/17.46 = 1.98.

No significant difference was found in the plasma concentrations of TDP-43 among the NC group, AD family, and PD family (0.165 ± 0.062 pg/ml, 0.154 ± 0.106 pg/ml, and 0.116 ± 0.075 pg/ml, respectively). However, FTD showed relatively high levels of TDP-43 (0.356 ± 0.202 pg/ml; \( p < 0.001 \)) compared to the other groups. The effect size of TDP-43 for the FTD group was calculated as 0.356/0.116 = 3.06.

Table 1. Demographic information and measured concentrations of plasma biomarkers in enrolled subjects. NC: normal control, AD family: patients with amnestic cognitive impairment or Alzheimer's disease, PD family: Parkinson's disease patients with or without dementia, FTD: frontotemporal dementia. CDR: clinical dementia ranking, MMSE: mini-mental state examination, H–Y stage: Hoehn and Yahr stage. \( * p < 0.05 \) using NC as reference. ** \( p < 0.01 \) using NC as reference. *** \( p < 0.001 \) using NC as reference. **** \( p < 0.0001 \) using NC as reference.

### Ethical standards.

The study was started and conducted after approval of the study protocol by the regional ethical committee in all the joined hospitals, including Chang Bin Show Chwan Memorial Hospital, Tri-Service General Hospital, National Taiwan University Hospital, and Kaohsiung Chang Gung Memorial Hospital. The study was carried out in accordance with relevant guidelines and regulations, including the World Medical Association (WMA) Declaration of Helsinki. A written informed consent was obtained from all participants and/or their legal guardians.

### Results

The demographic information of the enrolled subjects is listed in Table 1. Ninety-one normal controls aged 64.1 ± 6.8 years were enrolled and referred to as the NC group. The female percentage was 65.9%. The clinical dementia ranking (CDR) of every participant in the NC group was zero. The score of the mini-mental state examination (MMSE) was 28.6 ± 1.6. The AD family had forty-one patients with amnesic cognitive impairment and thirty-five patients with AD dementia. The female percentage was 72.4%. The age of the AD family was 74.0 ± 9.8 years. The CDR of the AD family was 0.78 ± 0.52, and the MMSE score was 23.1 ± 5.3, which were significantly lower than those of the NC group (\( p < 0.0001 \)). The PD family had 47 PD patients with normal cognition and 62 patients with PD dementia. The female percentage was 67.9%. The age of the PD family was 65.8 ± 10.4 years. The CDR was 0.40 ± 0.47. The MMSE score was 22.4 ± 5.7, which resulted in a significantly lower score (\( p < 0.0001 \)) than that of the NC group. For example, the effect size of T-Tau for the AD family was calculated as 34.51/17.46 = 1.98.

No significant difference was found in the plasma concentrations of TDP-43 among the NC group, AD family, and PD family (0.165 ± 0.062 pg/ml, 0.154 ± 0.106 pg/ml, and 0.116 ± 0.075 pg/ml, respectively). However, FTD showed relatively high levels of TDP-43 (0.356 ± 0.202 pg/ml; \( p < 0.001 \)) compared to the other groups. The effect size of TDP-43 for the FTD group was calculated as 0.356/0.116 = 3.06.

The measured levels of plasma biomarkers in the NC, AD family, PD family, and FTD groups are listed in Table 1. According to published papers\(^{49,53,54}\), the concentrations of plasma biomarkers assayed with immuno-magnetic reduction (IMR) in normal controls are independent of age. Hence, it is not necessary to have correction to age in the analysis. For plasma α-synuclein, the NC group showed significantly higher levels (58.68 ± 13.28 pg/ml) than the AD family (49.70 ± 15.58 pg/ml; \( p < 0.0001 \)), PD family (44.96 ± 10.88 pg/ml; \( p < 0.0001 \)) and FTD (40.34 ± 4.69 pg/ml; \( p < 0.0001 \)). However, NC showed significantly lower levels of plasma TDP-43 (15.72 ± 2.48 pg/ml) than the AD family (19.61 ± 9.28 pg/ml; \( p < 0.0001 \)) and FTD (17.08 ± 3.37 pg/ml; \( p < 0.0001 \)) than in the NC group (17.46 ± 9.28 pg/ml; \( p < 0.01 \)). Relatively higher levels of plasma TDP-43 were found in the FTD (18.42 ± 3.02 pg/ml; \( p < 0.0001 \)) than in the NC group (0.165 ± 0.062 pg/ml; \( p < 0.0001 \)) and AD family (0.154 ± 0.106 pg/ml and PD family (0.116 ± 0.075 pg/ml). There was no significant difference in plasma TDP-43 levels among the NC group, AD family and PD family.

The effect sizes of each biomarker in the AD family, PD family and FTD group were investigated. For a biomarker, the effect size is calculated via the ratio of the mean concentration in a given neurodegenerative disease to that in the NC group. For example, the effect size of T-Tau for the AD family was calculated as 34.51/17.46 = 1.98. The levels of plasma α-synuclein in the AD family, PD family and FTD group decreased compared to the NC group. Instead of Aβ1–40, the inversion of Aβ1–40, i.e., 1/Aβ1–40, was used to calculate the effect size.

Table 1. Demographic information and measured concentrations of plasma biomarkers in enrolled subjects. NC: normal control, AD family: patients with amnestic cognitive impairment or Alzheimer's disease, PD family: Parkinson's disease patients with or without dementia, FTD: frontotemporal dementia. CDR: clinical dementia ranking, MMSE: mini-mental state examination, H–Y stage: Hoehn and Yahr stage. \( * p < 0.05 \) using NC as reference. ** \( p < 0.01 \) using NC as reference. *** \( p < 0.001 \) using NC as reference. **** \( p < 0.0001 \) using NC as reference.

| Group       | NC       | AD family | PD family | FTD      |
|-------------|----------|-----------|-----------|----------|
| n (female%) | 91 (65.9%) | 76 (72.4%) | 109 (67.9%) | 25 (60.2%) |
| Age (years) | 64.1 ± 6.8 | 74.0 ± 9.8 | 65.8 ± 10.4 | 75.2 ± 11.6 |
| CDR         | 0        | 0.78 ± 0.52 | 0.40 ± 0.47 | –        |
| MMSE        | 28.6 ± 1.6 | 23.1 ± 5.3**** | 22.4 ± 5.7**** | –        |
| H–Y stage   | –        | –         | 2.20 ± 1.1 | –        |
| Aβ1–40 (pg/ml) | 58.68 ± 13.28 | 49.70 ± 15.58**** | 44.96 ± 10.88**** | 40.34 ± 4.69**** |
| Aβ1–42 (pg/ml) | 15.72 ± 2.48 | 19.61 ± 5.21**** | 17.08 ± 3.37**** | 18.42 ± 3.02**** |
| T-Tau (pg/ml) | 17.46 ± 9.28 | 34.51 ± 9.91**** | 26.64 ± 10.36**** | 41.53 ± 20.51**** |
| α-synuclein (fg/ml) | 96.7 ± 1179.8 | 4148 ± 1345* | 3648 ± 9065** | 543 ± 59.7 |
| TPD-43 (pg/ml) | 0.165 ± 0.062 | 0.154 ± 0.106 | 0.116 ± 0.075 | 0.356 ± 0.202**** |

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the effect size. This result demonstrates the role of amyloid β in AD. The FTD group showed the highest value of effect size in plasma T-Tau (\(= 2.38\)). In addition to T-Tau, the FTD group showed a relatively higher value (\(= 2.15\)) for the effect size in plasma TDP-43. The results revealed that the average levels of plasma T-Tau and TDP-43 in the FTD group were more than double those in the NC group. The average level of plasma T-Tau in the AD family was almost double that in the NC group. The PD family showed a relatively high value of effect size in plasma α-synuclein (\(= 37.7\)).

Notably, there are two reasons for this issue. Firstly, the levels of other biomarkers such as Aβ\(_{1–42}\), T-Tau, α-synuclein and TDP-43 elevate in AD, PD or FTD as compared to NC. The effect sizes of these biomarkers are higher than 1. However, the level of Aβ\(_{1–40}\) decreases in dementia groups as compared to NC. The effect size should be lower than 1, which is not consistent with other biomarkers. In order to have common expressions to show the increases in biomarker levels in dementia, the inversion of Aβ\(_{1–40}\), i.e., \(1/\text{Aβ}_{1–40}\), is used to calculate the effect size.

Secondly, if Aβ\(_{1–40}\) is used for calculating the effect size, the more changes in Aβ\(_{1–40}\) in AD, PD or FTD result in the lower values of the effect size. However, for other biomarkers, the more changes result in higher values of the effect size. This would under-estimate the dominance (or normalized effect size) of Aβ\(_{1–40}\) among biomarkers for AD, PD and FTD, as listed in Table 3. Hence, the inversion of Aβ\(_{1–40}\) i.e., \(1/\text{Aβ}_{1–40}\), is used to calculate the effect size to avoid this paralogism.

**Discussion**

As listed in Table 2, specific biomarkers show effect sizes of relatively higher values in AD, PD or FTD. It is worth investigating the dominance of biomarkers in neurodegenerative diseases (NDDs). In Table 2, the ranges of effect sizes among biomarkers differ greatly from each other. For example, the effect size of plasma α-synuclein ranges from 0.65 to 37.7, whereas it ranges from 1.09 to 1.25 for plasma Aβ\(_{1–42}\). To eliminate the difference in the ranges of effect size among biomarkers, the effect size of a given biomarker is normalized to the sum of effect sizes over NDDs. For instance, the normalized effect size of plasma Aβ\(_{1–42}\) in the AD family was calculated as 1.25/(1.25 + 1.09 + 1.17) = 0.356. The normalized effect sizes of every biomarker in the AD family, PD family and FTD group are listed in Table 3.

The dominance of a biomarker in an NDD is evaluated by calculating the percentage of the normalized effect size of the biomarkers in that of all biomarkers. According to Table 3, the dominance of plasma Aβ\(_{1–42}\) in the AD family is 0.356/(0.300 + 0.356 + 0.336 + 0.101 + 0.246) × 100% = 26.6%. The combinations of plasma Aβ\(_{1–40}\), Aβ\(_{1–42}\), T-Tau, α-synuclein and TDP-43 in the AD family, PD family and FTD are shown in Fig. 1.

For the AD family, the major domains were Aβ\(_{1–42}\) (26.6%), T-Tau (25.1%) and Aβ\(_{1–40}\) (22.4%). The three biomarkers are almost equally dominant in the AD family. This evidence highlights the associations between amyloidosis and tauopathy with AD from the plasma biomarker point of view.

It is well known that Aβs are peptides of amino acids 36–43 that result from cleavage of amyloid precursor protein (APP) by β and γ secretases. Certain misfolded Aβ, such as pyroglutamate Aβ, induce the accumulation of Aβ and the formation of insoluble Aβ plaques in the brain. The toxic Aβ plaques damage neurons, particularly those surrounding the hippocampus, resulting in memory disorders and cognitive decline in AD patients.
In addition, the hyperphosphorylation of tau proteins, which are abundant in brain neurons and stabilize the microtubules of neurons, causes the death of neurons in the brain\textsuperscript{59,60}. Due to neuron death, tau proteins are expressed by neurons, and neurofibrillary tangles can be observed in the biopsy. Meanwhile, regional atrophy of the brain occurs\textsuperscript{46}. Therefore, brain atrophy, Aβ plaques and neurofibrillary tangles in the brain are clinical features of AD.

The results in Fig. 1a showing the major dominance of Aβ\textsubscript{1–42}, T-Tau and Aβ\textsubscript{1–40} in the AD family is consistent with AD neuropathology. Some published papers point out that the plasma Aβ\textsubscript{1–42}-to-Aβ\textsubscript{1–40} ratio is significantly correlated with the density of Aβ plaques in the brain in AD\textsuperscript{45,61,62}. The elevations in plasma T-Tau levels due to regional atrophy of the brain in AD were demonstrated\textsuperscript{46}. All the results demonstrate the feasibility of using plasma Aβs and T-Tau to assess neuropathology or brain volumetry in AD.

Remarkably, as shown in Fig. 1a, TDP-43 is a minor biomarker in AD family (18.4%). Some papers have reported that TDP-43 pathology is frequently found in AD, especially in severe AD\textsuperscript{63–65}. AD patients with TDP-43 pathology have a rapid decline in cognition\textsuperscript{66}. Furthermore, TDP-43 species can coexist with Tau tangles in AD\textsuperscript{67}. All the findings provide strong evidence that TPD-43 could play a role in AD, as observed in Fig. 1a.

For the PD family shown in Fig. 1b, the most important dominant biomarker is α-synuclein, whose dominance (44.9%) is more than double that of individual Aβ\textsubscript{1–40} (16.8%), Aβ\textsubscript{1–42} (15.7%) and T-Tau (13.2%). According to the pathogenesis associated with PD, as α-synuclein molecules are phosphorylated, the β-sheets and oligomerization or fibrils of α-synuclein are easily formed, followed by the formation of Lewy bodies in dopaminergic neurons\textsuperscript{68–70}. Dopaminergic neurons with Lewy bodies become degenerative and unable to express dopamine. With the lack of dopamine, neurons in the motor cortex of a brain are damaged, which results in movement disorders in PD patients\textsuperscript{71–73}. The formation of Lewy bodies mainly consisting of α-synuclein in PD patients was demonstrated in autopsy steins\textsuperscript{74}. Therefore, α-synuclein is the most recognized biomarker for PD. The current results of plasma biomarkers also support the key role of α-synuclein in PD.
| Group                        | Aβ₁–40 | Aβ₁–42 | T-Tau | α-Synuclein | TDP-43 | Group                        | Aβ₁–40 | Aβ₁–42 | T-Tau | α-Synuclein | TDP-43 |
|------------------------------|--------|--------|-------|-------------|--------|------------------------------|--------|--------|-------|-------------|--------|
| aMCI (n = 41)                | 24.9%  | 27.3%  | 25.7% | 4.0%        | 18.1%  | AD family (n = 76)           | 22.4%  | 26.6%  | 25.1% | 7.5%        | 18.4%  |
| AD dementia (n = 35)         | 19.6%  | 24.2%  | 24.6% | 8.6%        | 23.0%  | PD family (n = 109)          | 16.8%  | 15.7%  | 13.2% | 44.9%       | 9.4%   |
| PD-NC (n = 47)               | 24.6%  | 22.1%  | 20.7% | 24.0%       | 8.6%   | PD dementia (n = 102)        | 21.9%  | 19.8%  | 23.9% | 0.8%        | 33.6%  |
| FTD (n = 25)                 | 21.0%  | 18.6%  | 23.3% | 0.7%        | 36.4%  | FTD (n = 25)                 | 21.9%  | 19.8%  | 23.9% | 0.8%        | 33.6%  |

Table 4. Dominance of plasma Aβ₁–40, Aβ₁–42, T-Tau, α-synuclein and TDP-43 in aMCI, ADD, PD-NC, PDD and FTD. aMCI amnesic mild cognitive impairment, AD Alzheimer’s disease, PD Parkinson’s disease, PD-NC Parkinson’s disease with normal cognition, FTD frontotemporal dementia.

Using IMR assays for plasma α-synuclein, research groups have validated the high discrimination of the PD family from NC (sensitivity > 80%, specificity > 80%)²⁰⁻²¹,4⁹. Furthermore, the levels of plasma α-synuclein positively correlated with the severity of cognitive impairment⁷⁵. The finding of the association of plasma α-synuclein levels with the thinning of the limbic cortex could support the positive correlation between plasma α-synuclein levels and cognitive decline⁷⁵. These results reveal the importance of assaying plasma α-synuclein to assess PD in clinical practice.

In Fig. 1c, it is clear that TDP-43 is the crucial biomarker for FTD (33.6%). Since the discovery of TDP-43 in 2006, it has been found that almost 50% of FTD and amyotrophic lateral sclerosis (ALS) have TDP-43 pathology.²⁶,²⁷ TDP-43 not only plays a role in nuclear transcription in relation to alternative splicing or exon skipping but also a role in RNA transport granules and in regulating local translation at distal locations.²⁸⁻²⁹. The aggregation of TDP-43 in the cytoplasm in FTD and ALS mainly results from hyperphosphorylation, ubiquitination or C-terminal truncation of TDP-43.³⁰⁻³². Thus, TDP-43 is recognized as a biomarker for FTD and ALS. The results of plasma biomarkers in Fig. 1c present the role of TDP-43 in FTD compared to amyloid, T-Tau and α-synuclein. Notably, the results in Fig. 1c show that T-Tau is the second major biomarker in FTD.

According to reported studies, in addition to TDP-43, tauopathy is a common characteristic pathological hallmark in FTD patients.³³⁻³⁵. The positive tauopathy in FTD patients was found to be due to mutations in the gene encoding tau (MAPT) on chromosome 17.³⁶. The mutations disrupt the normal binding of tau protein to tubulin, resulting in neuronal damage and pathological deposits of tau in the brain in FTD.

Other studies on plasma biomarkers in FTD reported increases in the levels of both TDP-43 and T-Tau compared to NC.³³,³⁵. Especially for TDP-43, FTD patients show significantly higher levels of plasma TDP-43 than AD and PD patients.³⁷. These results imply that the assay of plasma TDP-43 is promising for assessing FTD in the clinic. Both TDP-43 and T-Tau should be taken into account for developing drug therapy.

In Fig. 1, amyloid β and total tau protein are equally dominant in the AD family. In the PD family, only α-synuclein is dominant. TDP-43 is the first major biomarker, while T-Tau is the second major biomarker in FTD. The results clearly reveal that more than one biomarker should be considered in AD family and FTD. Combined therapies against amyloid β and total tau protein (or phosphorylated tau protein) in AD family and TDP-43 and T-Tau (or phosphorylated tau protein) in FTD should be investigated.

As AD family is split to aMCI and AD dementia, PD family is split to PD with normal cognition (PD-NC) and PD dementia. The dominance of plasma Aβ₁–40, Aβ₁–42, T-Tau, α-synuclein and TDP-43 in aMCI, AD dementia, PD-NC, PD dementia and FTD is analyzed, as listed in Table 4. The results are compared to Fig. 1 for AD family, PD family and FTD.

As AD family is split to aMCI and AD dementia, Aβ₁–40, Aβ₁–42 and T-Tau are dominant in aMCI, which is consistent with that in AD family. In AD dementia, the role of Aβ₁–40 is suppressed, whereas TDP-43 becomes one of the dominant biomarkers. As reported,²⁶⁻³⁰,³⁸⁻³⁹, TDP-43 pathology is frequently found in AD dementia, especially in severe AD. These findings provide strong evidence that TDP-43 could play a role in AD dementia.

As PD family is split to PD-NC and PD dementia, α-synuclein remains the definitely dominant biomarker in PD dementia. However, in PD-NC, in addition to α-synuclein, other biomarkers such as Aβ₁–40, Aβ₁–42 and T-Tau are equally dominant. The results reveal the involvements of amyloidosis and tauopathy in PD patients with normal cognition.

Some postmortem studies show the significant existence of amyloid fibrils and neurofibrillary tangles (NFTs) in brain of PD patients.³⁸⁻⁹⁰. It was suggested that aggregation of NFTs, the abnormal hyperphosphorylation of tau protein, the interaction between T-Tau and α-synuclein may all result in poor axonal transport or the cell death observed in PD patients.⁹⁰. Furthermore, the amyloidosis could contribute to the rapid progression of dementia in PD patients.³¹,³². However, the uptake of [¹¹C] Pittsburgh compound-B was found to be normal for the PD dementia.³³. All the results imply that amyloidosis and Tauopathy have roles in early-stage PD.

**Conclusion**

By using IMR assays for plasma Aβ₁–40, Aβ₁–42, T-Tau, α-synuclein and TDP-43 in NC, the AD family, the PD family and FTD, the dominant roles of these biomarkers in these neurodegenerative diseases were clarified. Consistent with neuropathological hallmarks, the current results show that plasma Aβs and T-Tau are major biomarkers in the AD family, while plasma TDP-43 could play a role in AD dementia. Plasma α-synuclein is dominant in the PD family compared to other biomarkers. Worth noting, amyloidosis and tauopathy also significantly contribute in early-stage PD. Plasma TDP-43 is very specific to FTD and is also involved in tauopathy. Thus, plasma biomarkers assayed with IMR clearly reflect the pathogenesis of neurodegenerative diseases. The results demonstrate the feasibility of precise and differential assessments of AD, PD and FTD using plasma...
biomarkers. It is strongly suggested that combined treatments against various biomarkers would be necessary for the therapy of neurodegenerative diseases.

**Data availability**

The dataset generated and analyzed in the current study is available from the corresponding author on reasonable request.

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**Author contributions**

This study was designed by S.Y.Y. and P.Y.C. and run by P.Y.C., F.C.Y., M.J.C., W.C.L., and C.H.L. S.Y.Y. prepared the manuscript. All authors approved the final manuscript.

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**Competing interests**

S.Y. Yang is an employee and shareholder of MagQu Co., Ltd. The other authors do not have conflicts of interest.

**Additional information**

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