

Biomarker-Based Corticobasal Syndrome Classification: The Added Value of Deep Phenotyping and Population Diversity

Jacy Bezerra Parmera, MD, PhD,^{1*} and Blas Couto, MD, PhD²

We read with great interest the article by Palleis and colleagues,¹ which prospectively stratified 50 patients with corticobasal syndrome (CBS) using cerebrospinal fluid (CSF)/amyloid-positron emission tomography (PET) (A β), tau-PET, and CSF α -synuclein (α Syn) seed amplification assays (SAA), relating these biomarker profiles to both cross-sectional and longitudinal clinical features. The authors are to be commended for their comprehensive, multimodal biomarker approach and for elucidating the molecular heterogeneity underlying CBS.

As the authors note, the presence of biomarkers in older individuals does not necessarily indicate that they are the primary drivers of the CBS phenotype. We were, however, intrigued by the finding that A β + patients were significantly older than A β - patients. Particularly, CBS with underlying Alzheimer's disease (AD) pathology typically manifests in younger-onset AD and affects younger patients than CBS due to corticobasal degeneration and CBS-predominant variant of progressive supranuclear palsy (PSP).² We inquire whether A β positivity in this cohort primarily reflects age-related co-pathology rather than primary AD neuropathological changes. This finding aligns with α Syn-SAA positivity, which also correlates with AD biomarkers. Additionally, in the Palleis et al. sample, tau-predominant patients were the younger subgroup (ie, less likely to have age-related co-pathology), and regardless of the biomarker used for patient stratification, those with one or more positive biomarkers were older than those with none.

Beyond its focus on biomarkers, this study would also help identify key clinical features that remain highly informative for assessing the likelihood of an underlying 4R-tauopathy versus AD or Lewy-type synucleinopathy (LTS). Those include the presence of oculomotor dysfunction,³ hyposmia/anosmia, rapid eye

movement sleep behavioral disorder, and visual hallucinations.^{4,5} The authors reported the subscores of the PSP-Rating Scale data stratified by biomarker positivity in Figure S4, although no inferential statistics were used for hypothesis-driven analysis. In Figures S4 to S6, we can see that the ocular motor dysfunction subscore was higher in the tau+ subgroup than in the tau- subgroup, whereas visual hallucinations were twice as frequent in patients stratified as LTS + AD than in those stratified as AD only. We consider that adding these clinical aspects is meaningful for interpreting those cases' biomarker profiles, as it would illustrate both the utility and the challenges of future application of this stratification in clinical practice. This would allow not only leveraging the deep clinical phenotyping of prospective cohorts such as the Munich cohort¹ but also integrating it with initiatives across the world spanning diverse PSP populations.^{6,7}

Relatedly, when uncertainty exists regarding primary versus co-pathology, FDG (fluorodeoxyglucose)-PET patterns can provide complementary information on neurodegenerative topography and may be particularly useful alongside amyloid biomarkers, as previously demonstrated for differentiating AD-related CBS from non-AD CBS.⁵ We therefore support the integration of combined biomarker strategies (A β biomarkers, FDG-PET, and SAA) in ambiguous cases.

Overall, the study provides compelling evidence that biomarker profiles correspond to clinically relevant differences, including greater cognitive impairment in A β -positive cases and milder motor progression in α Syn-positive patients. We believe that incorporating deep phenotyping data for each subgroup would further enhance the interpretability and translational applicability of this critical biomarker-based CBS framework. ■

¹Department of Neurology, Hospital das Clínicas, Faculdade de Medicina da Universidade de São Paulo (HC-FMUSP), São Paulo, Brazil; ²Instituto de Neurociencia Cognitiva y Traslacional (INCYT, CONICET-INECO-Universidad Favaloro), Buenos Aires, Argentina

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*Correspondence to: Dr. Jacy Bezerra Parmera, Department of Neurology, University of São Paulo School of Medicine, Rua Doutor Eneas de

Carvalho Aguiar, 255, Cerqueira Cesar, 05403-911 São Paulo, SP, Brazil; E-mail: jacy.parmera@hc.fm.usp.br

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