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PHOSPHORYLATED TAU: CANDIDATE BIOMARKER FOR AMYOTROPHIC LATERAL SCLEROSIS

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Abstract

IMPORTANCE—An increasingly varied clinical spectrum of cases with amyotrophic lateral sclerosis (ALS) has been identified, and objective criteria for clinical trial eligibility is necessary.

OBJECTIVE—We sought to develop a cerebrospinal fluid (CSF) biomarker sensitive and specific for the diagnosis of ALS.

DESIGN—Case-control study.

SETTING—Academic medical center.

PARTICIPANTS—51 individuals with ALS and 23 individuals with a disorder associated with a four-repeat tauopathy (4R-tau).

MAIN OUTCOME MEASURE—CSF level of tau phosphorylated at threonine 181 (ptau), and ratio of ptau to total tau (ttau).

RESULTS—Using a cross-validation prediction procedure, we found significantly reduced CSF levels of ptau and ptau:ttau in ALS relative to 4R-tau and to controls. In the validation cohort, the receiver operating characteristic area under the curve for the ptau:ttau ratio was 0.916, and the comparison of ALS to 4R-tau showed sensitivity=92% and specificity=91.7%. Correct classification based on low CSF ptau:ttau was confirmed in 18 (85.7%) of 21 cases with autopsy-proven or genetically-determined disease. In patients with available measures, ptau:ttau in ALS correlated with clinical measures of disease severity such as Mini Mental State Exam (n=51) and ALS Functional Rating Scale-Revised (n=42), and regression analyses related ptau:ttau to MRI

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(n=10) evidence of disease in the corticospinal tract and white matter projections involving prefrontal cortex.

CONCLUSIONS AND RELEVANCE—CSF ptau:ttau may be a candidate biomarker to provide objective support for the diagnosis of ALS.

Keywords

amyotrophic lateral sclerosis; cerebrospinal fluid; phosphorylated tau; biomarker

INTRODUCTION

Amyotrophic lateral sclerosis (ALS) is a progressive neurodegenerative condition with upper motor neuron (UMN) and lower motor neuron (LMN) motor deficits. ALS patients experience a rapid rate of decline over 3–5 years¹. Diagnostic evaluation of ALS is aimed typically at exclusion of other disorders. Phenotypic variability has resulted in controversy about clinical stratification strategies²: Patients may show strictly LMN or UMN disease, may have disease restricted to a particular segment (e.g., bulbar) or region (e.g., flail arm), and may be strongly lateralized^{3,4}. ALS may exhibit non-pyramidal motor system involvement, including cognitive difficulty in 33%–50%^{5,6} that extends to frontotemporal degeneration (FTD)⁷. Clinically presymptomatic ALS may exist in carriers of genetic mutations such as *TARDBP*⁸ and *C9orf72* hexanucleotide repeat expansion^{9,10}. Given these challenges in an era of disease-modifying therapies, it is critical to identify objective biomarkers of ALS during life.

ALS is considered part of the frontotemporal lobar degeneration (FTLD) spectrum of disorders. Approximately 95% of individuals with ALS have transactive DNA binding protein of ~43 kDa (TDP-43) at autopsy, and TDP-43 is also the histopathologic feature in half of FTLD¹¹. Most of the remaining FTLD patients have hyperphosphorylated tau¹². Perhaps the most common tauopathy is associated with four-repeat tau (4R-tau) in progressive supranuclear palsy (PSP). Deposition of pathologic tau is negligible in ALS, except for individuals with Guam ALS/parkinsonism which is predominantly a tauopathy¹³.

Tau can be assayed in cerebrospinal fluid (CSF). Studies of CSF total tau (ttau) levels in FTLD have been mixed^{14–22}. Since CSF ttau may be elevated following any neuronal injury, assays for tau phosphorylated at threonine 181 (ptau) attempt to improve specificity. Elevated CSF ptau is found in several conditions involving tau pathology, including AD²³, pathologically-confirmed FTLD-tau such as PSP^{24–26}, and FTLD due to a genetically-determined tauopathy²⁷. By comparison, patients with FTLD-TDP pathology as in ALS may have low CSF ptau because tau pathology is rare in these patients²⁴.

The present study evaluated the possibility that CSF ptau is reduced in ALS, while PSP individuals likely to have tau pathology were expected to have higher CSF ptau levels. We used a cross-validation prediction procedure to assess CSF ptau as a candidate ALS biomarker. We also assessed the relationship between CSF ptau and clinical markers of disease burden such as the Functional Rating Scale-revised²⁸. White matter (WM) neuroimaging may be valuable diagnostically in ALS^{29–31} and PSP^{32–34}. To evaluate the

extent of WM disease and relate this to CSF ptau, we obtained fractional anisotropy (FA) measures of WM disease in ALS.

METHODS

Clinical Evaluation

Participants—We studied 51 patients with ALS and 23 patients likely to have 4R-Tau pathology recruited from the ALS Center and the Penn FTD Center at the University of Pennsylvania. Experienced neurologists diagnosed ALS according to El Escorial-revised criteria³⁵, with initial evaluation showing definite=7, probable=18, possible=18, and suspected=5. Three additional individuals had ALS-FTD, with co-occurring FTD diagnosed according to published clinical criteria³⁶. Onset was bulbar=10, cervical=15, thoracic=1, and lumbosacral=22 (onset was unknown in those with FTD). Five patients had autopsy confirmation, 5 had a *C9orf72* expansion, and 1 had a pathogenic *TARDBP* mutation (p.N390S), consistent with TDP-43 pathology. The 4R-tau cohort was comprised of patients clinically diagnosed with PSP (n=15), which is highly associated with 4R-tau pathology at autopsy³⁷, autopsy-confirmed four-repeat tauopathy (CBD=3, PSP=2), and pathogenic mutations consistent with 4R-tau (*MAPT* E10+16=2 with one autopsy-confirmed, and *MAPT* p301.L=1). A subset of 43 ALS and PSP patients participated in another CSF study³⁸. We excluded patients with a three-repeat tauopathy to define a homogeneous contrast group. CSF was also available in 23 healthy seniors screened for dementia using a mini-mental state exam (MMSE)³⁹ score >28/30, were screened for AD pathology using an autopsy-validated t-tau to beta-amyloid ratio (<0.34)⁴⁰ and no neurological or psychiatric history. Table 1 summarizes demographic features (all p>0.05). Another cohort of 28 demographically-matched healthy seniors (age, education and gender p-values>0.1) with no neurological or psychiatric history were recruited as neuroimaging controls.

All individuals participated in a written informed consent procedure with their caregivers, when appropriate, that was approved by the Institutional Review Board at the University of Pennsylvania.

Functional measures—As summarized in Table 1, ALS patients, 4R-tau patients, and Seniors were evaluated clinically with the MMSE (n=78). A subset of ALS patients (n=42) were additionally evaluated on the Functional Rating Scale (ALSFRS-R)²⁸.

Lumbar Puncture Procedure and Analysis

Lumbar puncture (LP)—CSF samples were obtained during routine diagnostic LP, as described⁴⁰. Briefly, LP was performed at the L3/L4 lumbar space using a 20-gauge needle to collect 15 ml of CSF in polypropylene tubes (Corning Life Sciences, Lowell, MA). Samples were aliquotted and immediately stored at –80°C until analysis. CSF sample collection, storage, and analysis were performed according to published standard operating procedures⁴¹.

CSF analysis—Samples were analyzed using a Luminex xMAP platform (INNO-BIA AlzBio3™, Innogenetics, Ghent, Belgium) (n=52) or an ELISA assay (INNOTEST®,

Innogenetics, Ghent, Belgium) (n=13), as described⁴¹. Briefly, the xMAP platform utilized capture monoclonal antibodies (MAbs) 4D7A3 (A β ₁₋₄₂), AT120 (ttau), and AT270 (ptau) bound to color-specific beads. We used an assay sensitive to phosphorylation at threonine-181 since this is the Alzheimer's Disease Neuroimaging Initiative standard for which the highly reliable Luminex method is available⁴¹. Biomarker analytes were detected using reporting MAbs 3D6 (A β ₁₋₄₂) and HT7 (ttau, ptau). Some older samples were analyzed with an ELISA method, where MAbs for capturing and reporting ttau and ptau were AT120/HT7 and BT2, HT7/AT270, respectively. As described previously²⁰, ELISA values for A β also were measured using an "in house" method with the capturing MAb BAN-50, and the reporting MAb BC-05. Using an autopsy-validated formula⁴⁰, a linear regression model converted natural-log transformed raw CSF values from ELISA to an xMAP equivalent.

Statistical analysis—We evaluated overall group-level differences for raw ptau, ttau, and ptau:ttau ratio with non-parametric Kruskal-Wallis and *post hoc* Mann-Whitney U tests for descriptive purposes. We also confirmed that a potential covariate for ALS progression rate, defined using previously published criteria (48 - ALSFRS-R)/Disease Duration in Months)²⁸ did not contribute to group differences. We used a cross-validation procedure to evaluate ptau, ttau, and ptau:ttau ratio as candidate biomarkers for individual patient screening. We randomly divided ALS and 4R-tau cohorts into comparably sized training (ALS n=26; 4R-Tau n=11) and validation (ALS n=25; 4R-Tau n=12) cohorts. eTable 1 summarizes the training and validation cohorts. Since ALS and 4R-tau differ in age and disease duration and these factors may influence CSF analyte levels⁴², we performed a logistic regression for each CSF analyte that included age and disease duration nuisance covariates. These logistic regressions were completed in the training cohort to generate a probabilistic likelihood of ALS, and then these probabilities were entered into receiver operating characteristic (ROC) curves. We defined the optimal cutoff to assess sensitivity and specificity at a probability 0.703, equivalent to the proportion of ALS patients in the training cohort (26 out of 37) and then applied this logistic regression model to the independent validation cohort. We report screening accuracy using a chi-squared (χ^2) test: Patients in the validation cohort whose ALS probability exceeded the 0.703 threshold were predicted as having ALS and otherwise assigned to the predicted 4R-tau group. We performed Pearson's correlations between each analyte (ptau:ttau ratio, ptau, and ttau) and functional measures, summarized in Table 1. For each correlation, we used the predicted probability of ALS as an age- and disease duration-adjusted proxy for each CSF analyte.

Imaging Procedure and Analysis

Acquisition—Diffusion-weighted MRIs were available for 10 ALS patients (1 with ALS-FTD) from a SIEMENS 3.0T Trio scanner using an 8-channel coil. Diffusion-weighted images (DWI) were acquired with a 30-directional sequence involving single-shot, spin-echo, diffusion-weighted echo planar imaging (FOV=245mm; matrix size=128 \times 128; number of slices=57; voxel size=2.2mm isotropic; TR=6700msec; TE=85msec; fat saturation). We acquired 30 volumes with diffusion weighting (b=1000 s/mm²) along 30 non-collinear directions per subject, and either one (n=2) or four (n=17) without diffusion weighting (b=0 s/mm²). When four volumes were collected without diffusion weighting,

these volumes were averaged to increase signal-to-noise ratio. Reasons for exclusion included health and safety (e.g., difficulty breathing while supine, metallic implants, shrapnel, claustrophobia) and lack of interest in an imaging study. DWI were also available for 9 4R-tau patients. T1-weighted MRI volumes were also acquired in the same scanning session with MPRAGE acquisition parameters: repetition time=1620msec; echo time=3msec; slice thickness=0.9mm; flip angle=15°; matrix=192×256, and in-plane resolution=0.9×0.9mm.

Preprocessing—Whole-brain MRI volumes were preprocessed using PipeDream (<https://sourceforge.net/projects/neuropipedream/>) and Advanced Normalization Tools (ANTs, <http://www.picsl.upenn.edu/ANTs/>), as described⁴³. Briefly, PipeDream deformed each individual dataset into local template space in a canonical stereotactic coordinate system. Each participant's T1 image was warped to the template via the symmetric diffeomorphic procedure in ANTs. For DWI, motion and distortion artifacts were removed by affine co-registration of each image with diffusion weighting to the unweighted (b=0) image. Diffusion tensors (DTs) were computed using a linear least-CSF biomarker for ALS Grossman et al 8 squares algorithm⁴⁴ implemented in Camino⁴⁵, and tensors were reoriented using the preservation-of-principal-directions algorithm⁴⁶. Fractional anisotropy (FA) was computed from the DT image for each subject. Distortion between T1 and DT images was corrected by registering the FA image to the T1 image. The DT image was then warped to template space by applying both the FA-to-T1 and T1-to-template warps for each subject. FA images were smoothed using a 4mm full-width half-maximum isotropic Gaussian kernel.

Statistical Analysis—Analyses of FA were performed in SPM8 using the two-samples t-test module. FA volumes were analyzed using an explicit mask (FA > 0.25) to constrain comparisons to WM regions. Comparisons of ALS patients to healthy seniors used a height threshold of $q < 0.05$ with false discovery rate (FDR) correction for multiple comparisons, and comparisons of 4R-tau to seniors used a height threshold of $q < 0.005$ with FDR correction. Both comparisons used an extent threshold of 200 voxels. Regression analyses related FA to the adjusted ptau:tttau ratio at $p < 0.05$ (uncorrected) with a 50-voxel extent. Regression analyses were constrained to WM fibers with reduced FA using explicit masks generated from the results of the direct comparisons with healthy seniors; different thresholds were used for group comparisons to create disease masks of comparable size. Using a deterministic tractography procedure in Camino, WM fibers were tracked in a healthy elderly template using the DTI sequence described above. Fiber tracts that passed through voxels of reduced FA were retained to define the masks for regression analyses. This was done to limit the interpretation of a correlation between WM and CSF to WM fibers with disease.

RESULTS

Median raw CSF analyte values for ALS, 4R-tau, and Seniors are illustrated in Figure 1. Kruskal-Wallis tests revealed group differences for ptau:tttau ratio [$\chi^2=30.55$; $p < 0.001$] and for ptau ng/ml [$\chi^2=22.80$; $p < 0.001$]. Planned *post hoc* Mann-Whitney tests revealed that, relative to 4R-tau, ALS has reduced ptau:tttau ratio [$Z=3.74$; $p < 0.001$] and reduced ptau

levels [$Z=2.82$; $p=0.005$]. ALS also had reduced ptau:ttau ratio [$Z=4.92$; $p<0.001$] and ptau levels [$Z=4.36$; $p<0.001$] relative to Seniors. There was no group effect for ttau ng/ml [$\chi^2=1.73$; $p>0.1$]. By comparison, 4R-tau had only marginally reduced ptau [$Z=2.27$; $p=0.05$] relative to Seniors.

ROC analyses illustrated in Figure 2 showed an area under the curve (AUC) for ptau:ttau ratio of 0.916 ($p<0.001$). In the training cohort, the probabilistic-ALS cutoff achieved 80.8% sensitivity and 90.9% specificity. A cross-validation analysis using the same cutoff in the validation cohort revealed 92% sensitivity and 91.7% specificity [$\chi^2=24.90$; $p<0.001$]. An analysis of ptau alone also was robust (AUC=0.923; $p<0.001$). We found 80.8% sensitivity and 81.8% specificity in the training cohort, but the validation cohort achieved high sensitivity (88%) with only modest specificity (75%) [$\chi^2=17.42$; $p<0.001$]. The ttau analyte also achieved a significant AUC (AUC=0.885; $p<0.001$), with 84.6% sensitivity and 81.8% specificity in the training cohort, and high sensitivity (92%) with modest specificity (75%) in the validation cohort [$\chi^2=14.69$; $p<0.001$].

Follow-up analyses of individuals with autopsy confirmation or a genetic mutation ($n=21$) showed correct classification in 18 (85.7%) of 21 patients using the most robust analyte, ptau:ttau ratio. The three misclassified cases included one *C9orf72* expansion patient, one *MAPT* (E.10+16 C>T) mutation patient, and one autopsy-confirmed ALS patient.

Correlation analyses in ALS using age- and disease-duration-adjusted CSF levels revealed that MMSE is related to pttau:ttau ratio ($r=0.342$; $p<0.05$), ptau ($r=0.354$; $p<0.05$), but not to ttau; ALSFRS-R is related to ptau ($r=0.448$; $p<0.005$) and ttau ($r=0.406$; $p<0.01$), but less to pttau:ttau ratio ($r=0.263$). CSF levels were not related to Progression Rate ($r<0.25$).

Figure 3 Panel A illustrates reduced WM FA in ALS that extends throughout frontal WM, the corpus callosum and the anterior limb of the internal capsule. Specific anatomic loci and WM tracts are summarized in eTable 2. Regression analysis related reduced ptau:ttau ratio to reduced WM FA in the corticospinal tract subjacent to primary motor cortex, prefrontal WM projections, and the corpus callosum (not shown). Figure 3 Panel B shows areas of reduced FA in PSP. Peak foci of reduced FA, summarized in eTable 2, were found in frontal, parietal, corpus callosum, internal capsule and brainstem regions. Regression analysis related reduced ptau:ttau ratio to reduced FA in the midbrain and uncinate fasciculus (not shown).

DISCUSSION

CSF levels of phosphorylated tau were very low in ALS. A cross-validation analysis revealed that ptau and ptau:ttau ratio appear to distinguish individuals with ALS from 4R-tau and from Seniors. This was confirmed in the subgroup of patients with known histopathology. Lower ptau and ptau:ttau ratio correlated with clinical measures of disease, and with MRI measures of reduced WM FA in the corticospinal tract and prefrontal cortex in ALS subgroups.

The histopathologic abnormality in sporadic ALS is TDP-43, and ALS patients (except for those with ALS/parkinsonism who are Chamorro from Guam) have negligible brain

hyperphosphorylated tau at autopsy²⁴. Thus, we predicted low CSF ptau levels in ALS. The ptau:ttau ratio was consistently sensitive and specific, generalizing from training to validation cohorts, and thus is a candidate biomarker for screening ALS. Another study of TDP-43 proteinopathies and 4R-tauopathies reported similar findings³⁸. Additionally, almost all cases with known TDP-43 pathology had a low ptau:ttau ratio. Two of the 3 incorrectly classified cases had genetically-determined disease, and we cannot rule out that these patients may have additional pathology due to another condition⁴⁷.

Some previous work described elevated CSF ttau in ALS^{48,49}, while others reported normal ttau levels⁵⁰. Interpretation of inconsistent results should be performed cautiously because of the substantial variability associated with the ELISA method used in those studies⁴¹. The present study used a more reliable Luminex method to assess most CSF analyte levels. Other studies reported significantly reduced CSF amyloid precursor protein levels in ALS, and elevated CSF A β levels were related to shorter survival^{51,52}, possibly reflecting the small number of ALS patients who have concurrent Alzheimer's pathology⁵³. Some studies described abnormal axonal markers that were related to survival⁵⁴ and abnormal glial markers that were related to progression rate⁴⁹, although our observations of reduced ptau were unrelated to survival and progression rate. Two reports described CSF TDP-43 levels in ALS^{55,56}, although the variability of results, including substantial overlap with control values, suggests that TDP-43 assays may be premature.

Lower CSF ptau and ptau:ttau in ALS correlated with clinical measures, and although there are many measures of clinical functioning, this suggests that ptau:ttau ratio may be a sensitive marker of disease. ALS is associated with cognitive difficulty in many individuals^{5,6}, and we found that ptau and ptau:ttau ratio correlates with cognitive functioning. ALS-FRS-R is commonly used to reflect disease severity in ALS and this correlated with ptau. Additional converging evidence suggesting that ptau:ttau ratio may be biologically meaningful comes from WM neuroimaging in anatomic regions known to be compromised pathologically in ALS⁵⁷. Since CSF ptau:ttau ratio appears to be related to both clinical and imaging measures, ptau:ttau ratio may be a candidate marker to assess eligibility in clinical trials for disease-modifying treatments of ALS.

Our findings also suggest that a low ptau:ttau ratio may be specific for ALS. We demonstrated this by contrasting ALS with individuals highly likely to have 4R-tau histopathology. Others also have reported comparative studies to demonstrate specificity^{52,54}. While not unreasonable to expect elevated CSF ptau levels of in this cohort because tau is hyperphosphorylated in autopsy assessments of these conditions, some reports have described elevated levels¹⁴⁻¹⁷, some normal levels^{18,25,58}, and some reduced levels¹⁹ relative to healthy controls. This variability may be due in part to mixed etiology in clinically-diagnosed groups and the less reliable ELISA method used in most prior studies. Regardless of the basis for previous findings, our observations suggest a reliable difference between individuals with ALS and those highly likely to have 4R-Tau pathology.

Several caveats should be kept in mind when evaluating our findings. Our cohort was relatively small. Tau is phosphorylated at several sites, and assaying other phosphorylation sites may be informative. Most participants were assessed soon after the onset of typical

ALS, and it would be important for future work to assess ALS patients with other phenotypic presentations and lengths of disease, and to evaluate CSF ptau levels in these different presentations. The contrast group consisted of patients four-repeat forms of tau because of limited CSF available from individuals with three-repeat tau pathology and our desire to have a relatively homogeneous contrast group, and it would be important to evaluate CSF ptau levels in individuals with three-repeat tau pathology. Limited MRI assessments were available because of patient limitations, and verification of FTLT-DTP pathology was possible in only a subset of cases. With these caveats in mind, our cross-validation prediction design suggests that individual patients with ALS highly likely to be due to FTLT-DTP pathology are characterized by a low CSF ptau:ttau ratio relative to individuals highly likely to have FTLT-Tau pathology, and low CSF ptau:ttau is associated with several clinical and imaging measures of ALS.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Data analysis was conducted by Murray Grossman, Corey T. McMillan, and John Powers.

The corresponding author had full access to all the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

Dr. Hu declares that he has filed a provisional patent on phosphorylated-tau: total-tau ratio as a biomarker for FTLT-DTP and ALS through Emory University.

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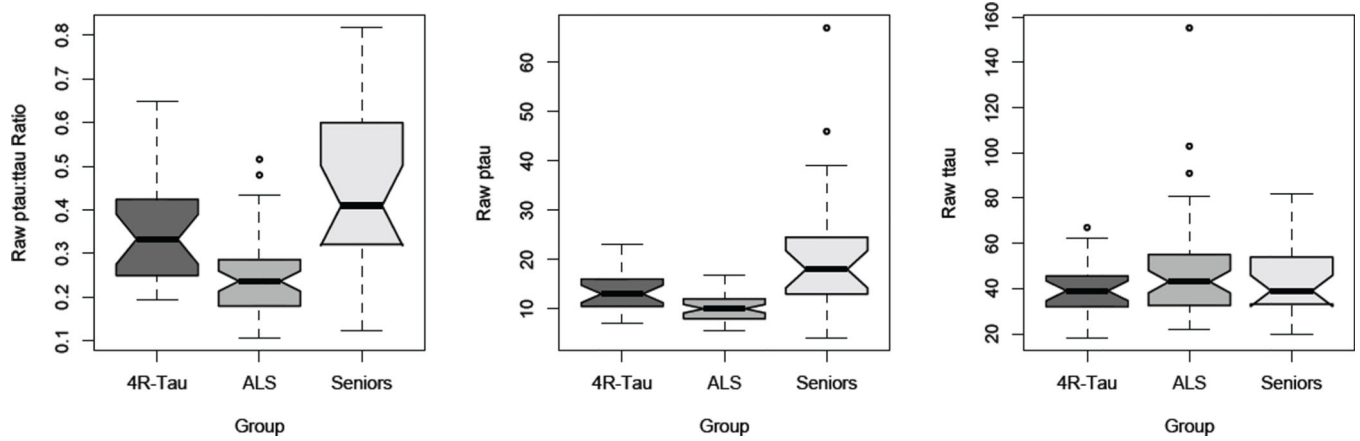


FIGURE 1. BOXPLOTS OF CSF ANALYTES PTAU NG/ML, TTAU NG/ML, AND PTAU:TTAU RATIO IN AMYOTROPHIC LATERAL SCLEROSIS, FOUR-REPEAT TAUOPATHY, AND HEALTHY SENIORS. NOTE. Dark lines in boxplots illustrate median CSF value, notches illustrate interquartile range (non-overlapping notches are significantly different), and error bars represent full range of data.

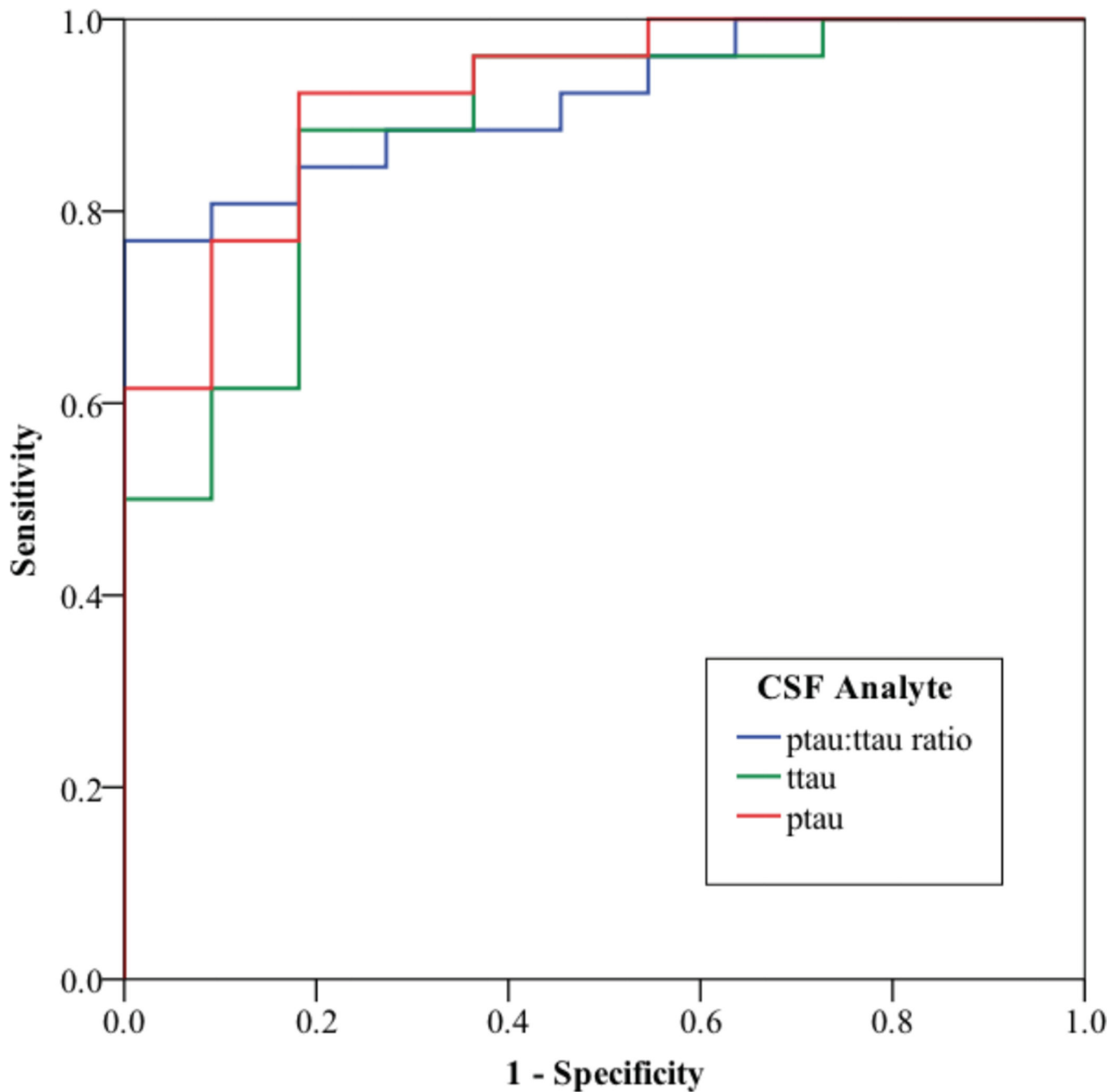


FIGURE 2. RECEIVER OPERATING CHARACTERISTIC CURVE ILLUSTRATING THE SENSITIVITY AND SPECIFICITY OF CSF PTAU, TTAU, AND PTAU:TTAU RATIO IN AMYOTROPHIC LATERAL SCLEROSIS RELATIVE TO FOUR-REPEAT TAUOPATHIES.

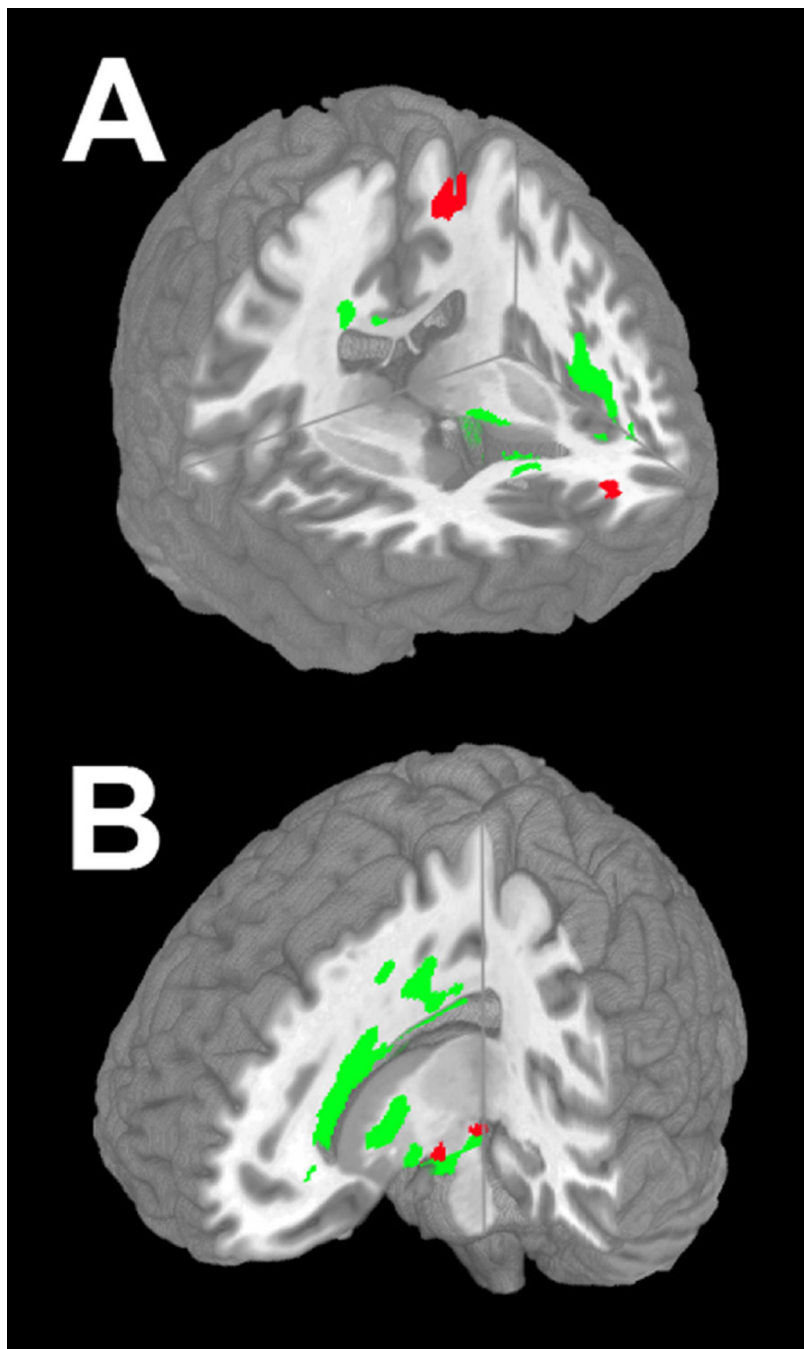


FIGURE 3. REDUCED WHITE MATTER FRACTIONAL ANISOTROPY IN AMYOTROPHIC LATERAL SCLEROSIS AND 4R-TAU, AND REGRESSIONS RELATING ADJUSTED CSF PTAU:TTAU RATIO TO FRACTIONAL ANISOTROPY. NOTE. Panel A: Right anterior view of anatomic distribution of reduced fractional anisotropy in ALS ($q < 0.05$, FDR-corrected; green). Red areas indicate anatomic distribution of regressions relating adjusted ptau:ttau ratio to fractional anisotropy in corticospinal tract, prefrontal centrum semiovale, and body of corpus callosum (not illustrated). Panel B: Left anterior view of

anatomic distribution of reduced fractional anisotropy in 4R-tau ($q < 0.005$, FDR-corrected; green). Red areas indicate anatomic distribution of regressions relating adjusted ptau:ttau ratio to fractional anisotropy in midbrain, right uncinate (not illustrated).

TABLE 1

MEAN (\pm S.D.) DEMOGRAPHIC & FUNCTIONAL MEASURES FOR AMYOTROPHIC LATERAL SCLEROSIS (ALS), FOUR-REPEAT TAUOPATHIES (4R-TAU), AND HEALTHY SENIORS

FUNCTIONAL MEASURE	ALS (N=51)	4R-TAU (N=23)	HEALTHY SENIORS (N=23)
Age	54.88 (10.6)	65.4 (9.4) *	59.9 (6.0)
Education	14.86 (3.9)	14.55 (3.4)	15.9 (4.3)
Gender (M / F)	35 / 16	14 / 9	12 / 11
MMSE (adjusted)	27.53 (3.3)	24.45 (5.13) ^{^#}	29.39 (0.7)
Disease Duration (months)	24.72 (14.1)	54.87 (23.4) *	--
ALSFRS-R (n=42)	37.74 (7.9)	--	--
Progression Rate (n=42)	0.46 (0.4)	--	--

NOTE:

* 4R-TAU differs significantly from ALS at $p < 0.001$;

[^] 4R-TAU differs significantly from ALS at $p < 0.01$;

[#] 4R-TAU differs significantly from healthy seniors at $p < 0.001$.