



Clinicopathologic complexities – the multi-proteinopathy problem

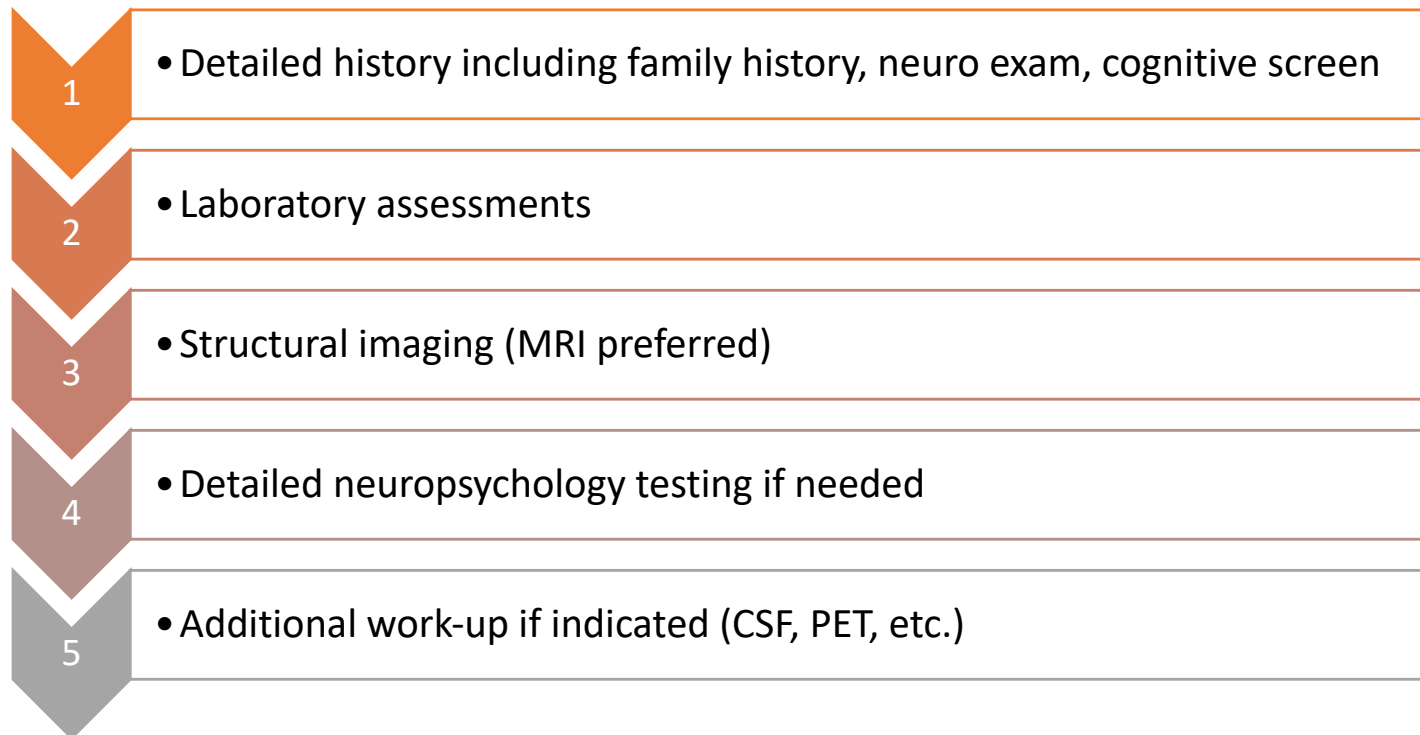
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Barbara and Peer Baekgaard Professor of Alzheimer's Disease Research
Clinical Core Leader, Indiana Alzheimer's Disease Center

Department of Neurology

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The Diagnostic Process



HPI + Cognitive Data + Neurologic Exam

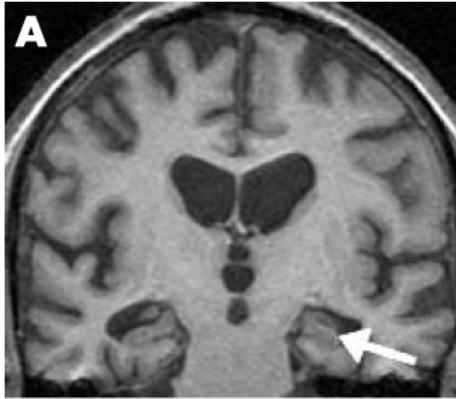
- Formulate the differential diagnosis on the basis of published criteria
 - NIA-AA criteria for MCI and dementia / Dubois criteria
 - AD variants – Crutch criteria for PCA; Gorno -Tempini IvPPA criteria
 - FTD – Lund-Manchester or Neary criteria
 - McKeith DLB criteria
 - Etc.



Rule Out Etiologies

- Depression
- Hypothyroidism
- Vit. B12 deficiency
- End organ failure
- NPH
- Subdural hematoma
- Toxic encephalopathy
- HIV/Syphilis as appropriate

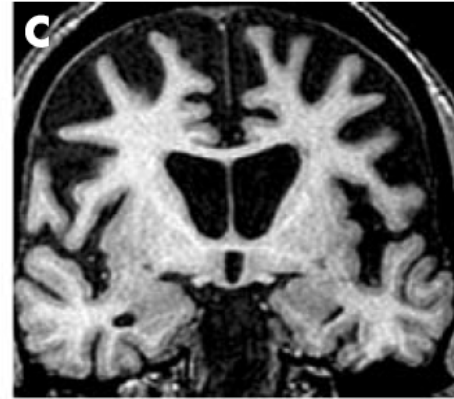




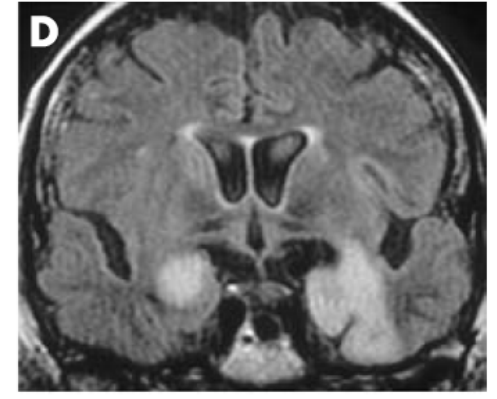
Alzheimer's disease



Semantic dementia



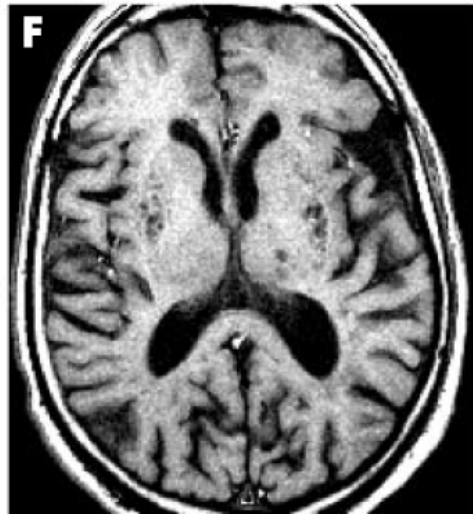
Frontal dementia



Limbic encephalitis



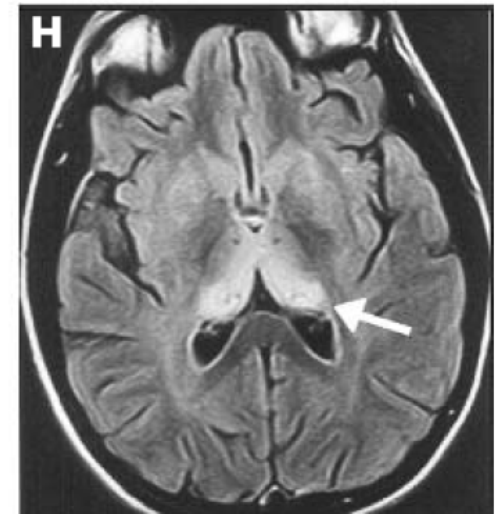
Tumour



Lacunar state

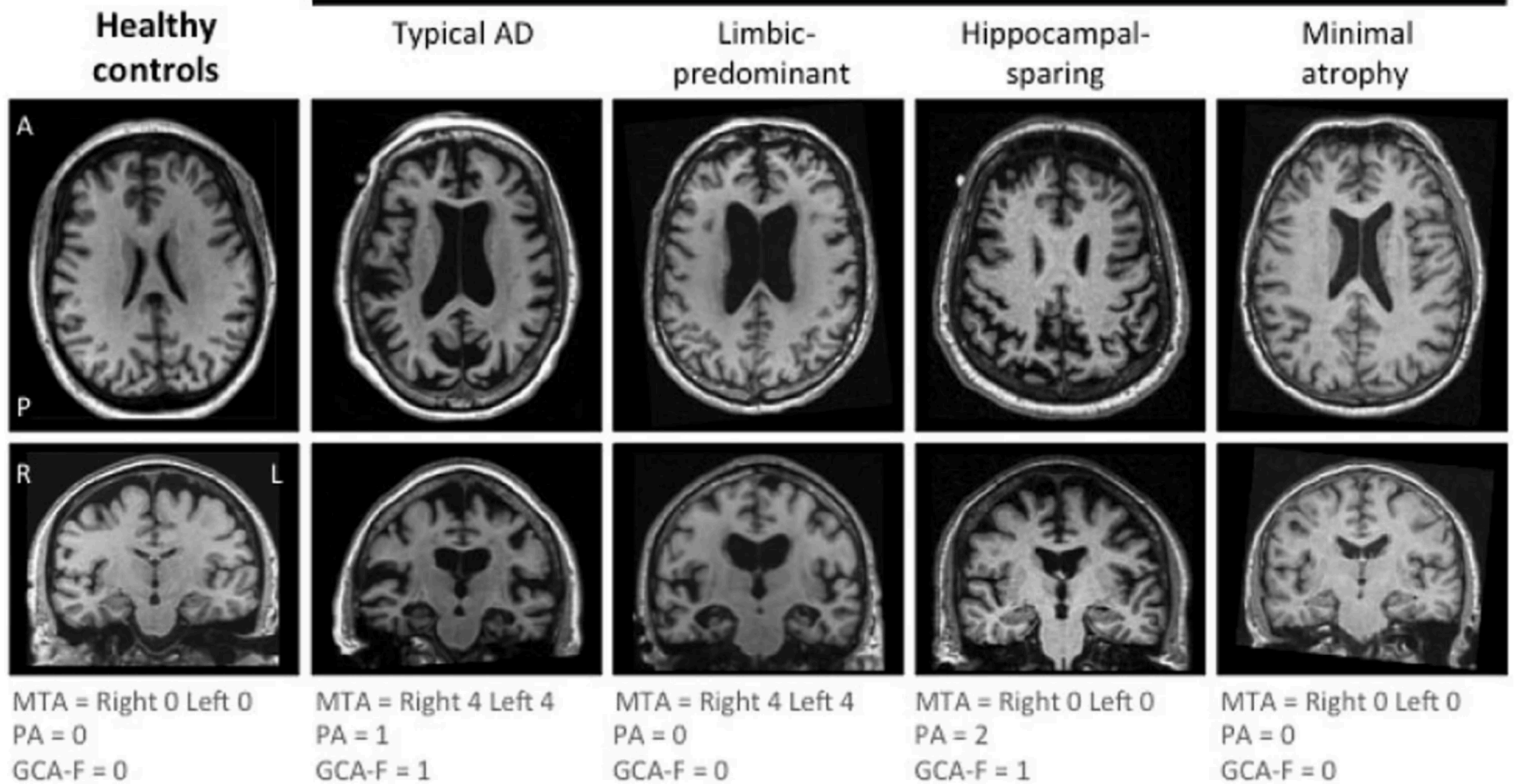


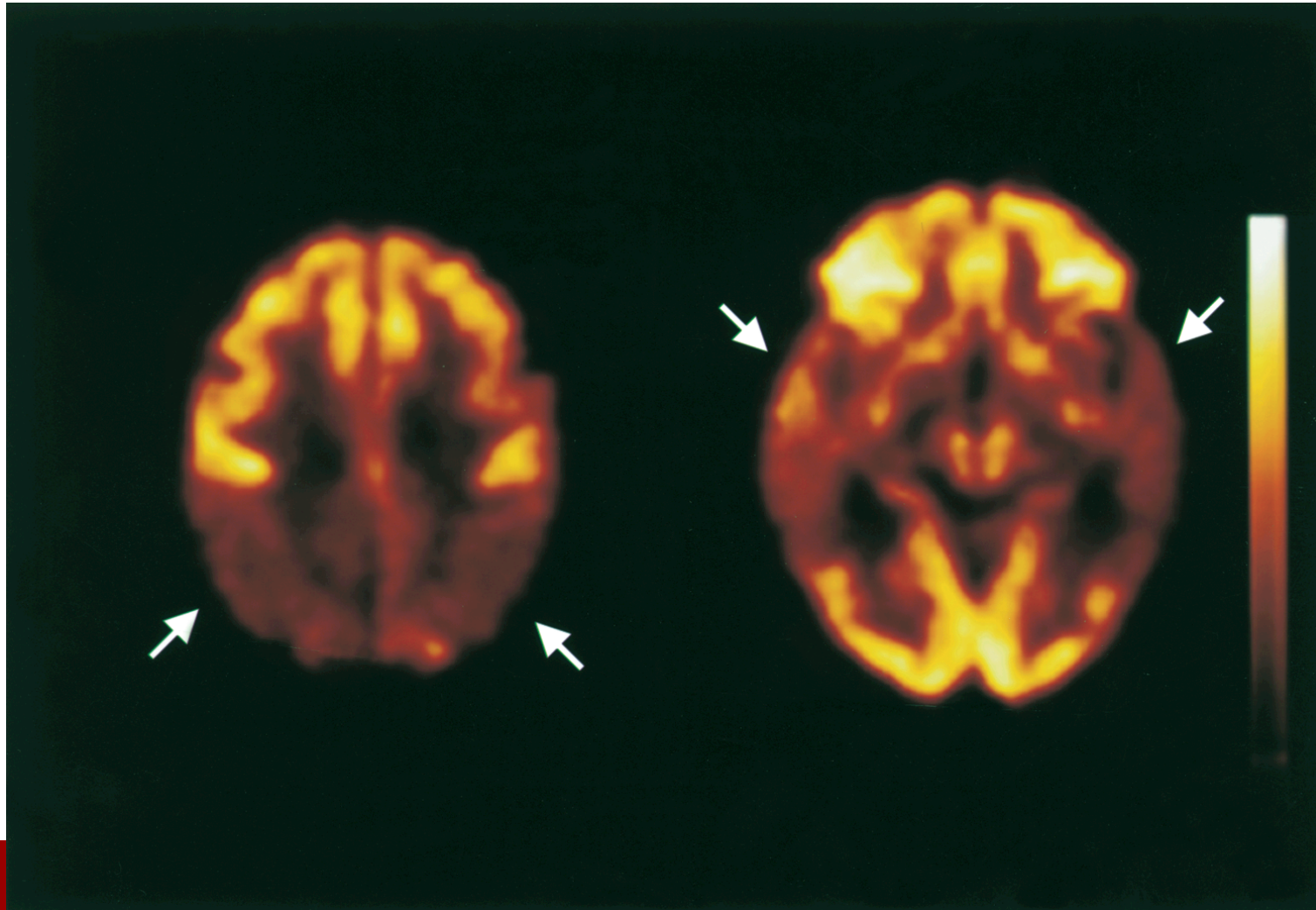
CADASIL



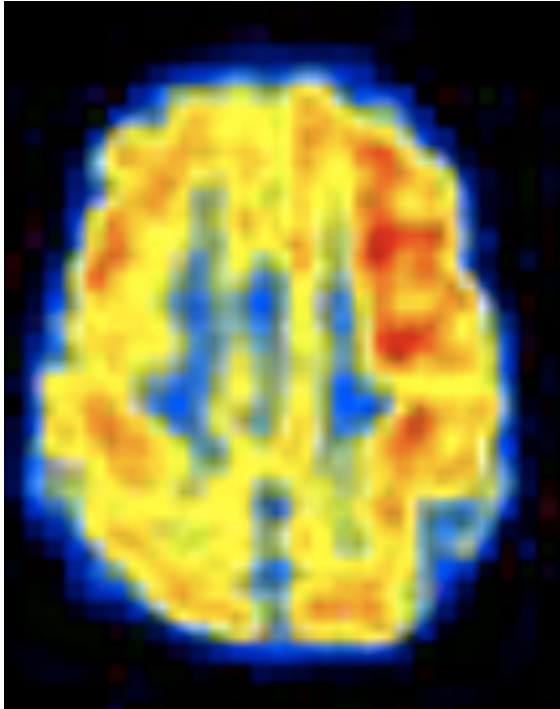
nvCJD

Mild Cognitive Impairment

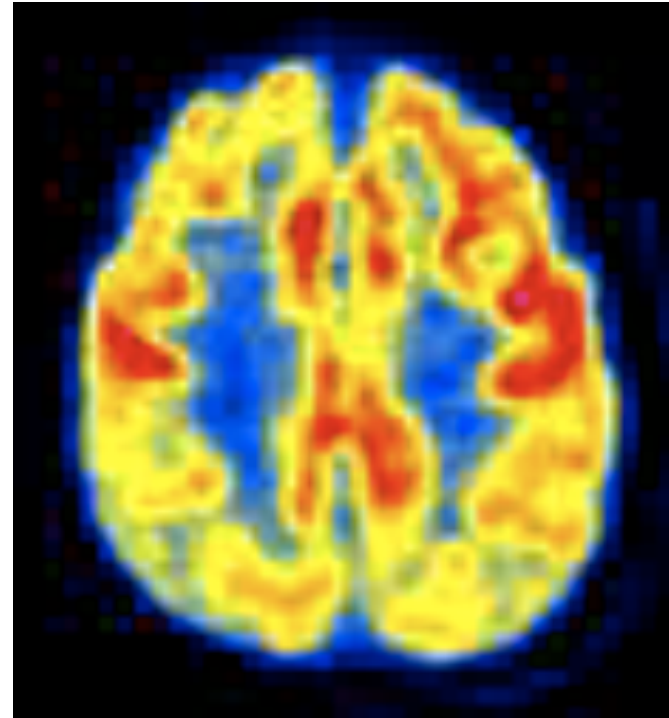




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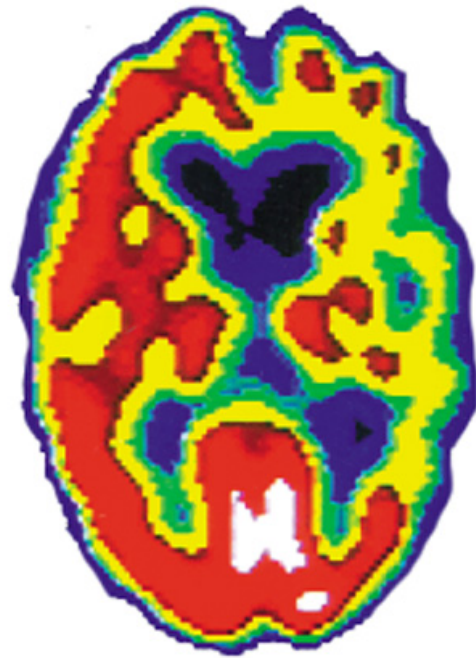
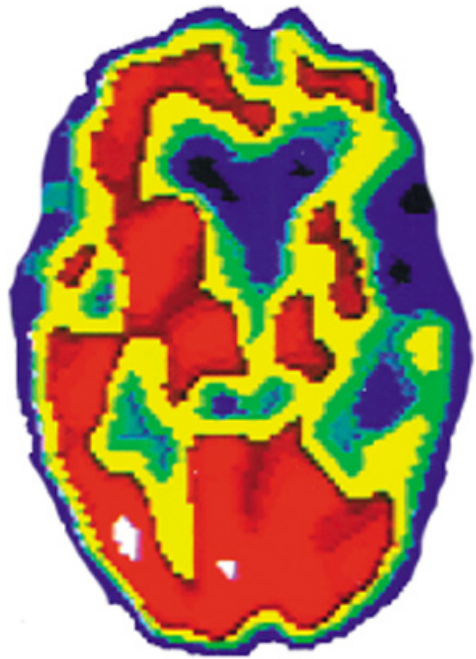


CBD



DLB

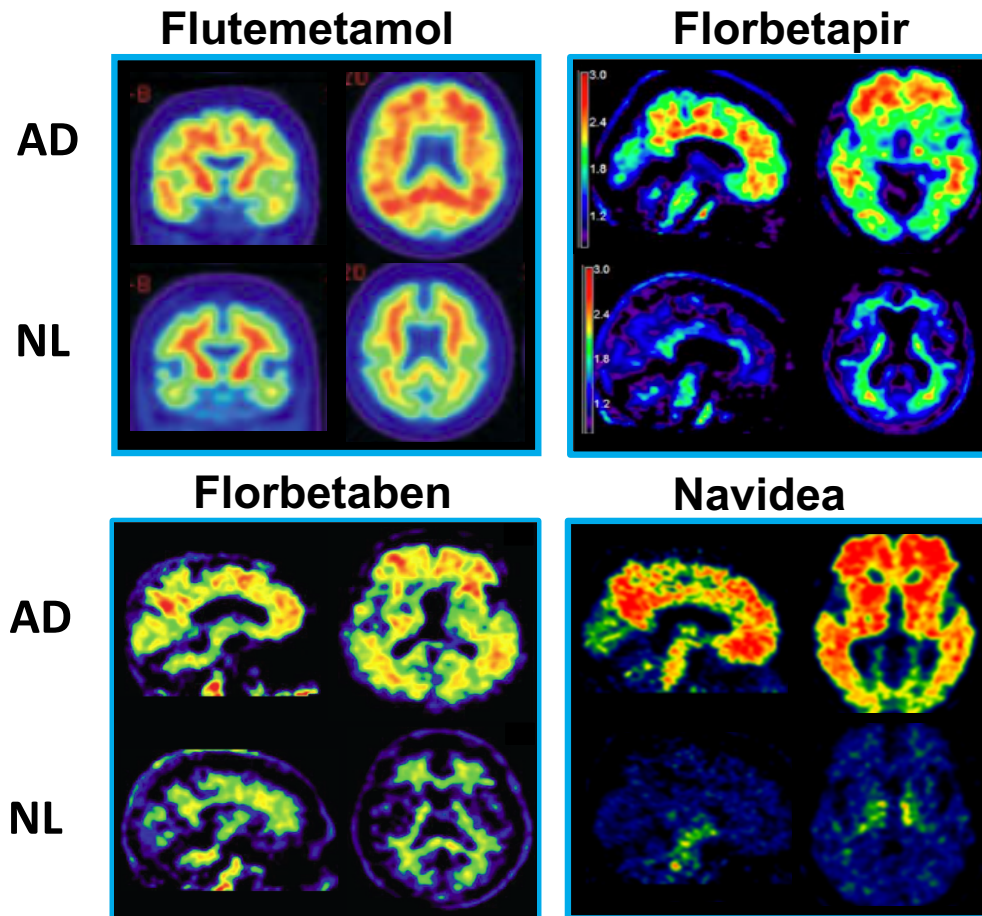




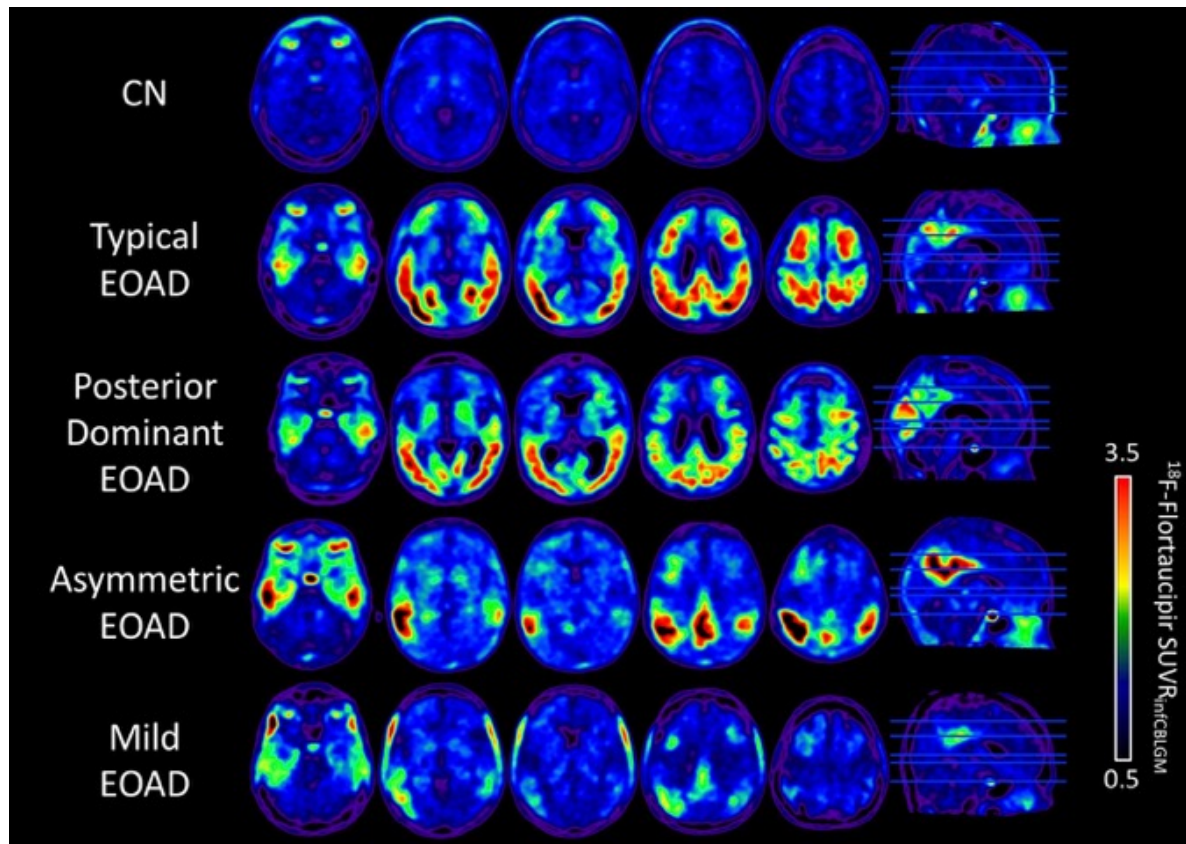
PNFA



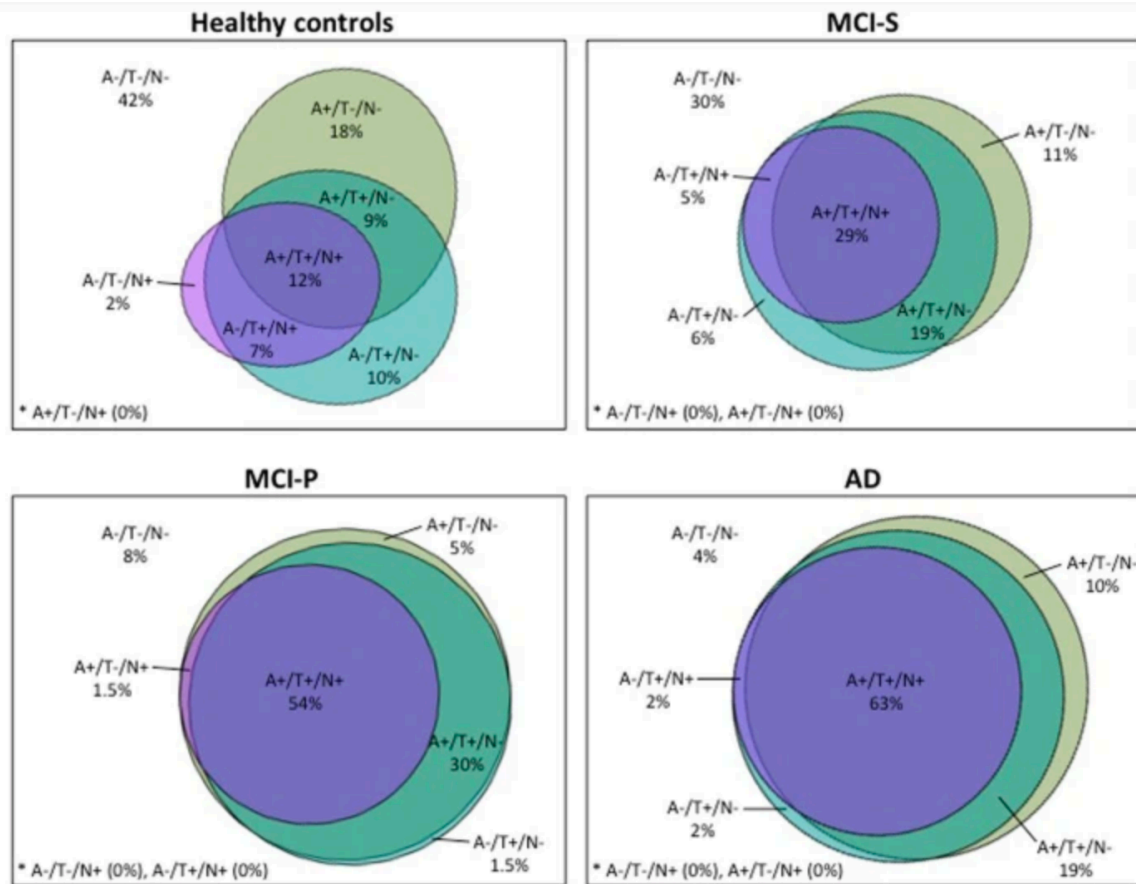
Amyloid PET



Tau PET – LEADS data



ROS MAP

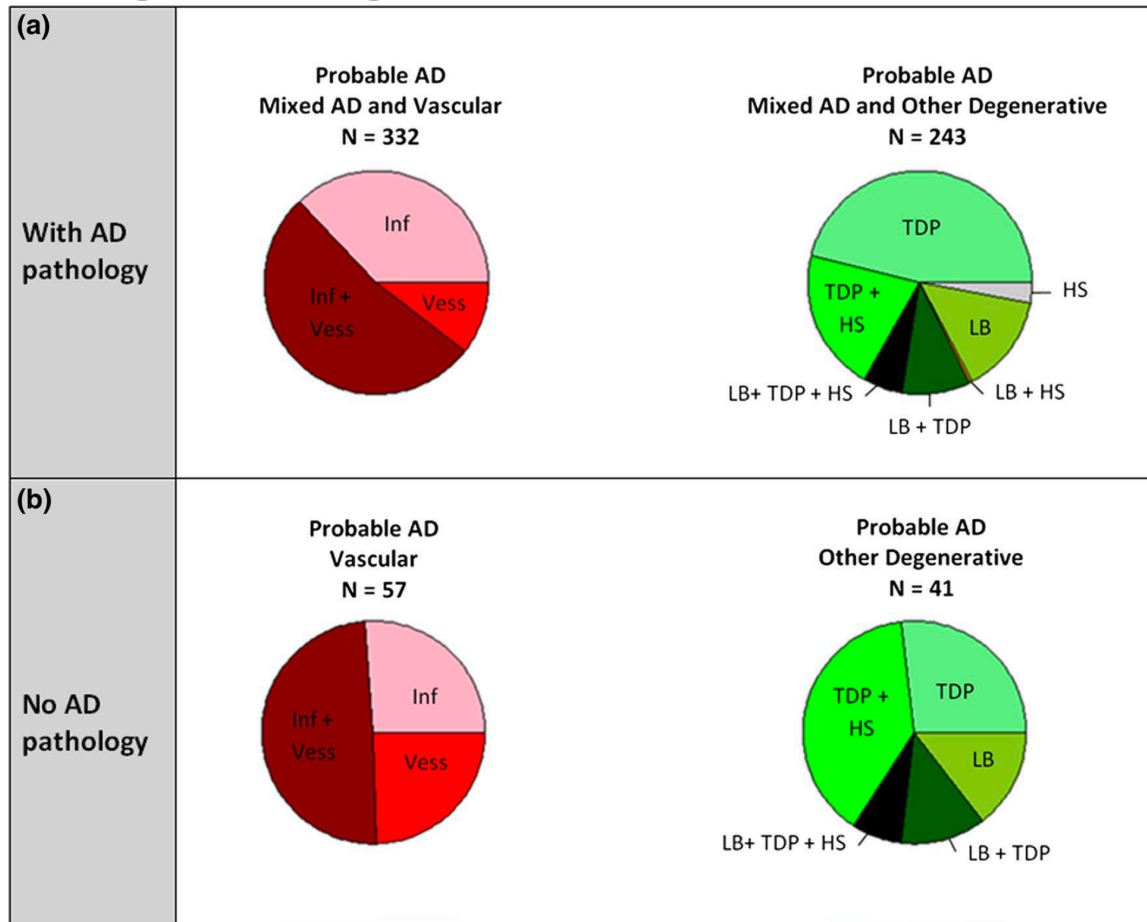


Prevalence of each A/T/N group Percentages of participants in each ATN group for healthy controls, MCI-S, MCI-P, and AD. HC = healthy controls. MCI-S = MCI participants that are clinically stable across time. MCI-P = MCI participants that progress to AD. AD = Alzheimer's disease. MCI = mild cognitive impairment. CSF = cerebrospinal fluid. A - = CSF A β normal. A + = CSF A β abnormal. T - = CSF p-tau normal. T + = CSF p-tau abnormal. N- = CSF t-tau normal. N + = t-tau abnormal.

Kapasi et al, Acta Neuropathol 2017

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And it is getting more complicated...



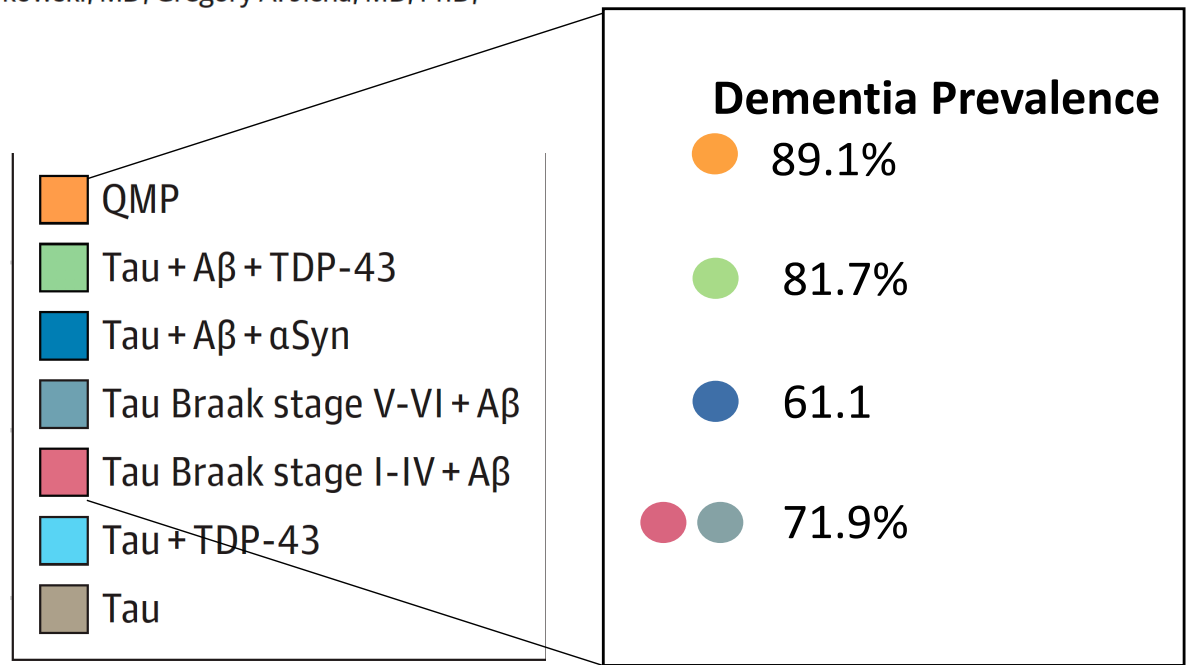
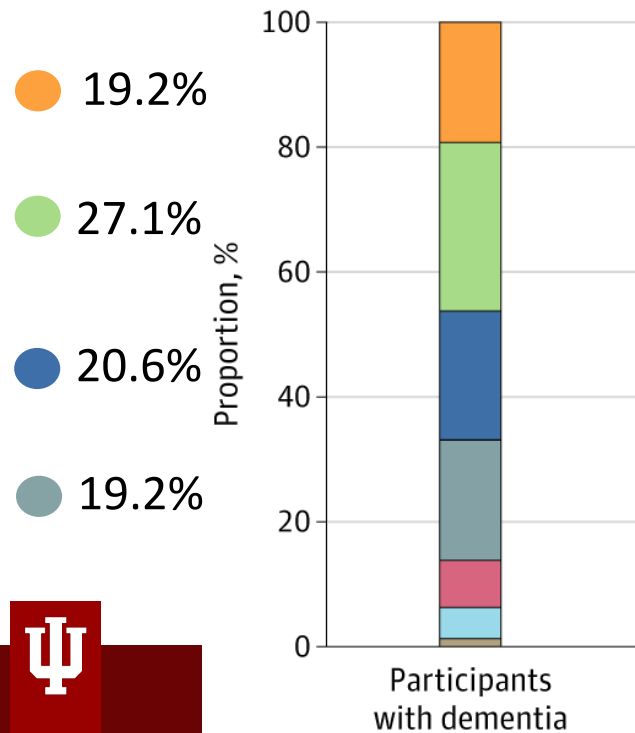
ROS MAP

Kapasi, Acta Neuropath 2017



Prevalence and Clinical Phenotype of Quadruple Misfolded Proteins in Older Adults

Shama Karanth, MDS; Peter T. Nelson, MD, PhD; Yuriko Katsumata, PhD; Richard J. Kryscio, PhD; Frederick A. Schmitt, PhD; David W. Fardo, PhD; Matthew D. Cykowski, MD; Gregory A. Jicha, MD, PhD; Linda J. Van Eldik, PhD; Erin L. Abner, PhD, MPH



JAMA Neurology | **Original Investigation**

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- 14/83 (16.9%) of cognitively normal participants had quadruple proteinopathy
- MCI with quadruple proteinopathy showed faster conversion to dementia (more aggressive disease)
- PURE AD was rare (19.2%)



LATE (Nelson, Brain 2019)

- LATE is 100x more common than FTD due to TDP-43
- Clinically: older patients (LATE-20% in 80+ years old), more gradual cognitive decline and more agitation than pure AD but the combination – accelerated cognitive decline > pure AD
- Imaging: asymmetric, more severe hippocampal atrophy (anterior hippocampus), hypometabolism/atrophy in the medial temporal lobe and inferior frontal gyrus



ARTAG (Nolan et al. J Neuropathol Exp Neurol 2019)

- 64% of autopsy cases (N=83) showed ARTAG
 - 67% of amnesic predominant
 - 43% of CBD
 - 57% of lvPPA
 - 57% of very mild dementia
 - 50% of PCA
 - All CN (N=3)



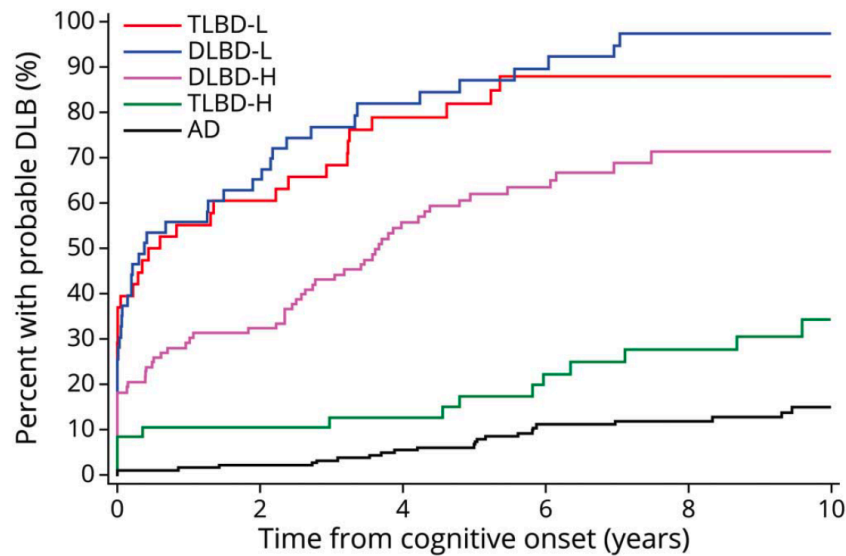
LBD spectrum (Ferman et al, Neurology 2020)

- In the presence of **tau NFTs** the diagnostic accuracy of clinical Dx of probable DLB (2+ of VH, fluctuations, parkinsonism, RBD) and DLB syndrome (1 sx) declined
 - w/o NFT - 96% sensitivity for prob DLB with diffuse LB
 - w NFT – 70% sensitivity for prob DLB with diffuse LB (+ onset of DLB clin syndrome later in the disease course)
- Among those with pure AD pathology 16% had 2+ and 22% 1 feature c/w clinical Dx of probable DLB or DLB syndrome



LBD + NFT affect both the clinical presentation and prognosis

Figure 1 Cumulative incidence model for the latency of probable dementia with Lewy bodies (DLB) across pathologic subgroups



L - NFT-
H - NFT+



Causative Pathology

- Can clinicians accurately predict all concomitant proteinopathies in the absence of biomarkers?

WE CAN'T

- **In cases with multiple proteinopathies, a formal clinicopathologic consensus conference might be warranted** to determine the primary etiology via interpretation of the pathology data in the context of the observed clinical syndrome

