

Computer model of spread of dementia can predict future disease patterns years before they occur

March 21 2012

Researchers at Weill Cornell Medical College have developed a computer program that has tracked the manner in which different forms of dementia spread within a human brain. They say their mathematic model can be used to predict where and approximately when an individual patient's brain will suffer from the spread, neuron to neuron, of "prion-like" toxic proteins -- a process they say underlies all forms of dementia.

Their findings, published in the March 22 issue of *Neuron*, could help patients and their families confirm a diagnosis of dementia and prepare in advance for future cognitive declines over time. In the future -- in an era where targeted drugs against dementia exist -- the program might also help physicians identify suitable brain targets for <u>therapeutic intervention</u>, says the study's lead researcher, Ashish Raj, Ph.D., an assistant professor of computer science in radiology at Weill Cornell Medical College.

"Think of it as a weather radar system, which shows you a video of <u>weather patterns</u> in your area over the next 48 hours," says Dr. Raj. "Our model, when applied to the baseline <u>magnetic resonance imaging</u> scan of an individual brain, can similarly produce a future map of degeneration in that person over the next few years or decades.

"This could allow <u>neurologists</u> to predict what the patient's



neuroanatomic and associated <u>cognitive state</u> will be at any given point in the future. They could tell whether and when the patient will develop speech impediments, <u>memory loss</u>, behavioral peculiarities, and so on," he says. "Knowledge of what the future holds will allow patients to make informed choices regarding their lifestyle and therapeutic interventions.

"At some point we will gain the ability to target and improve the health of specific <u>brain regions</u> and <u>nerve fiber</u> tracts," Dr. Raj says. "At that point, a good prediction of a subject's future anatomic state can help identify promising target regions for this intervention. Early detection will be key to preventing and managing dementia."

Tracking the Flow of Proteins

The computational model, which Dr. Raj developed, is the latest, and one of the most significant, validations of the idea that dementia is caused by proteins that spread through the brain along networks of neurons. It extends findings that were widely reported in February that Alzheimer's disease starts in a particular brain region, but spreads further via misfolded, toxic "tau" proteins. Those studies, by researchers at Columbia University Medical Center and Massachusetts General Hospital, were conducted in mouse models and focused only on Alzheimer's disease.

In this study, Dr. Raj details how he developed the mathematical model of the flow of toxic proteins, and then demonstrates that it correctly predicted the patterns of degeneration that results in a number of different forms of dementia.

He says his model is predicated on the recent understanding that all known forms of dementia are accompanied by, and likely caused by, abnormal or "misfolded" proteins. Proteins have a defined shape, depending on their specific function -- but proteins that become



misshapen can produce unwanted toxic effects. One example is tau, which is found in a misfolded state in the brains of both Alzheimer's patients and patients with frontal temporal dementia (FTD). Other proteins, such as TDP43 and ubiquitin, are also found in FTD, and alpha synuclein is found in Parkinson's disease.

These proteins are called "prion-like" because misfolded, or diseased, proteins induce the misfolding of other proteins they touch down a specific neuronal pathway. Prion diseases (such as mad cow disease) that involve transmission of misfolded proteins are thought to be infectious between people. "There is no evidence that Alzheimer's or other dementias are contagious in that way, which is why their transmission is called prion-like."

Simple Explanation for Clinically Observed Patterns of Dementia

Dr. Raj calls his model of trans-neuronal spread of misfolded proteins "very simple." It models the same process by which any gas diffuses in air, except that in the case of dementias the diffusion process occurs along connected neural fiber tracts in the brain.

"This is a common process by which any disease-causing protein can result in a variety of dementias," he says.

The model identifies the neural sub-networks in the brain into which misfolded proteins will collect before moving on to other brain areas that are connected by networks of <u>neurons</u>. In the process the proteins alter normal functioning of all brain areas they visit.

"What is new and really quite remarkable is the network diffusion model itself, which acts on the normal brain connectivity network and manages



to reproduce many known aspects of whole brain disease patterns in dementias," Dr. Raj says. "This provides a very simple explanation for why different dementias appear to target specific areas of the brain."

In the study, he was able to match patterns from the diffusion model, which traced <u>protein</u> disbursal in a healthy brain, to the patterns of brain atrophy observed in patients with either Alzheimer's disease or FTD. This degeneration was measured using MRI and other tools that could quantify the amount of brain volume loss experienced in each region of the patient's brain. Co-author Amy Kuceyeski, Ph.D., a postdoctoral fellow who works with Dr. Raj, helped analyze brain volume measurements in the diseased brains.

"Our study demonstrates that such a spreading mechanism leads directly to the observed patterns of atrophy one sees in various dementias," Dr. Raj says. "While the classic patterns of dementia are well known, this is the first model to relate <u>brain</u> network properties to the patterns and explain them in a deterministic and predictive manner."

More information: Related study: <u>medicalxpress.com/news/2012-03</u> ... scans-alzheimer.html

Provided by New York- Presbyterian Hospital

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