

New study of Xia-Gibbs syndrome spurs development of patient registry

May 7 2018

Researchers at the Human Genome Sequencing Center at Baylor College of Medicine have conducted a study of 20 individuals with Xia-Gibbs syndrome, a rare condition that has symptoms of severe developmental delay, sleep apnea, delayed speech and generalized upper body weakness, and have established a registry to collect genetic and other clinical information from patients with the condition.

As discovered in 2014 by a team led by Dr. Richard Gibbs, professor and director of the Human Genome Sequencing Center at Baylor, Xia-Gibbs syndrome is the result of new changes within the AT-Hook DNA Binding Motif Containing 1 gene (AHDC1). The new study, which appears in the *American Journal of Medical Genetics*, shows that the gene is more susceptible to mutations than previously known.

Thorough study of the 20 individuals showed details of the clinical picture: scoliosis was a common feature, boys were less likely to develop speech and thinning of a region of the brain, known as the corpus callosum, often occurred.

"Understanding the correlations between the molecular studies and the patient phenotypes is an important gateway to fully understanding the disorder," said Dr. Michael Wangler, assistant professor of molecular and human genetics at Baylor and contributing author on the study.

The registry, which is private and HIPAA protected, is the next step in identifying and studying the full spectrum of the disorder. The current

group of 30 enrollees in the registry learned about the study through their doctors, social media, including a private Facebook page for families affected by Xia-Gibbs syndrome, or elsewhere on the internet.

"We know of about nearly 70 families worldwide, and roughly half of these are participating in the registry thus far," said Dr. Yunyun Jiang, postdoctoral associate in the Human Genome Sequencing Center and lead author on the study.

As more patients join the registry, accompanying laboratory work aims to understand the full range of underlying molecular changes that cause the symptoms of Xia-Gibbs syndrome.

"We recently encountered a patient who is more than 50 years old, which is remarkable because this condition is typically only identifiable in children," said contributing author Dr. David Murdock, assistant professor of molecular and human genetics at Baylor and assistant director of the clinical lab in the Human Genome Sequencing Center. "When we fully understand how this gene damage causes Xia-Gibbs syndrome, we can begin the path to a cure."

Current work in the Human Genome Sequencing Center is focused on whether mild mutations lead to a more mild form of the disorder, as well as the construction of animal models to help study the condition.

"Given the rarity of this disease, scientists and physicians are seldom able to gather all the information about a group like this. It is something that has really been lacking in the field," said Gibbs, corresponding author on the study and also the Wofford Cain Chair and Professor in Molecular and Human Genetics at Baylor. "With the growth of the registry and future mouse model studies, there is hope that we will get to the bottom of this one."

More information: Yunyun Jiang et al. The phenotypic spectrum of Xia-Gibbs syndrome, *American Journal of Medical Genetics Part A* (2018). [DOI: 10.1002/ajmg.a.38699](https://doi.org/10.1002/ajmg.a.38699)

Provided by Baylor College of Medicine

Citation: New study of Xia-Gibbs syndrome spurs development of patient registry (2018, May 7) retrieved 24 April 2024 from <https://medicalxpress.com/news/2018-05-xia-gibbs-syndrome-spurs-patient-registry.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.