For some of us, they carry the bright blue of our grandfather’s eyes. For others they result in the characteristic cleft chin or the familial tendency toward color blindness. But in some families, the genetic mutations handed down from generation to generation aren’t as benign. And for one family in particular, the mutation results in early-onset Alzheimer’s disease.

“We heard about a family in Girardota a number of years ago,” said UC Santa Barbara neuroscientist Kenneth S. Kosik, who studies Alzheimer’s disease and other tau protein pathologies and works closely with research partners in Colombia, where this family is located.

The subject of a new paper published in the journal Alzheimer’s & Dementia, the family was discovered over the course of Kosik’s work in the Medellin region. While studying the genetics of a different, roughly 6,000-member family in the Colombian province of Antioquia with a now well-documented history of early-onset Alzheimer’s disease due to a mutation in the PSEN1 (presenilin) gene, the researchers came across this other, smaller family presenting the same symptoms.

“We analyzed their genes for mutations,” said Kosik, UC Santa Barbara’s Harriman Professor of Neuroscience and co-director of the campus’s Neuroscience Research Institute. “And, remarkably, we found another mutation in the very same gene. And they are unrelated to the first family.”
Though located on the same gene as the first family’s, genetic analysis revealed that this genetic mutation was in a different location, ruling out the possibility that this smaller group of people was a subset of the larger family. “Having it arise twice in two families that are in pretty close proximity is a little surprising,” Kosik said. “But improbable things happen.”

Though the two Colombian families live down the road from each other today, further genetic investigation revealed that their mutations originated on two separate continents, in two different populations. By sequencing their genomes, the researchers were able to follow the mutations backward, complementing their genetic work with an examination of historical and demographic records.

Using identity-by-descent analysis, the first, larger family’s mutation was traced back to the Spanish Empire, and is thought to have been brought to South America by a single founder who might have been a conquistador during the colonization of Colombia in the early 16th century. The second, smaller family’s PSEN1 mutation, meanwhile, has been traced to Africa.

“We pinned it down pretty clearly to show that in the region around the mutation the sequences around there were of African origin,” Kosik said. These findings were further bolstered by Colombia’s history: In the 16th century, West Africans were imported into the country as enslaved people, and, like the conquistadores of the same era, mixed and mingled with the local population. In fact, the concentration of Afro-Colombians led to the modern-day establishment of the department of Chocó, an enclave on the western coast of the country inhabited predominantly by descendants of the enslaved Africans.

Such genetic mutations are not uncommon, Kosik said.

“All of us carry different mutations,” he explained. Often, these errors in our code don’t survive. They don’t beat the odds to become part of the next generation, or they die with their last, childless carriers; in some cases they make it impossible to reproduce. But in Colombia, through millennia of population dynamics — the violence and introduced diseases brought on by conquest, the ebb and flow of populations, the mingling and also the isolation — this PSEN1 mutation persisted.

“It was never selected against,” Kosik said. While we think of 40 as a ‘young’ age to have genetic early-onset Alzheimer’s, it’s late enough in a person’s life for them to have passed on their genes to the next generation before it manifests. And the idea
of a genetic component to this condition wouldn’t come until the 20th century.

**A Double-edged Sword**

When a genetic link to the first Colombian family’s early onset form of Alzheimer’s was established in 1997, it provided a valuable opportunity. With the family’s consent, the research collaboration led by Kosik and neuroscientist Francisco J. Lopera of the Grupo de Neurociencias de Antioquia at the Universidad de Antioquia, was able to document the progression of symptoms from its earliest stages and later, with additional investigators, to conduct clinical trials for a drug with potential to arrest the disease.

“If you really understand what are the first things that change, what might be the effects that trigger these changes, you might find a way to interfere with that pathway,” said lead author Juliana Acosta-Uribe, a graduate student researcher in the Kosik Group who joined the team as she was completing her medical training at Universidad de Antioquia. This second family could provide insight into, among other things, whether or not the progression of genetic early-onset Alzheimer’s is similar for all people with a PSEN1 mutation or if it is family-specific, she said.

For carriers of PSEN1 mutations, this knowledge could be something of a double-edged sword. An explanation for the condition that strikes with frightening regularity members of their family as soon as they reach their fourth decade could ease the uncertainty and point to a potential cure. But it also could impart to these families the dilemma that may force them to choose between having children with a significant chance of inheriting a gene mutation that all but guarantees dementia at age 50, or selecting against the mutation by electing not to have children.

“These families are traditionally large families, where you can have eight, 10, 13 kids,” Acosta-Uribe said. “The patients we’re seeing now, they have already had their own children. So it’s their kids that are now really understanding what’s happening and asking questions.”

“We’re now very actively discussing these questions of disclosure,” Kosik said of a separate, but no less significant research endeavor. “We’re really very involved in finding out which people want to know if they have this mutation, and what they would do differently if they knew or didn’t know.”

Research on this study was also conducted by Laura Ramirez Aguilar (lead author), Margarita M. Giraldo, Sonia Moreno, Ana Baena, Rosario Costumal, David Aguillón,
Lucía Madrigal, Amanda Saldarriaga, Alexander Navarro, Gloria P. Garcia and Daniel C. Aguirre-Acevedo of Universidad de Antioquia. Ethan Geier and Jennifer S. Yokoyama of UC San Francisco; Nicholas Cochran and Richard M. Myers of HudsonAlpha Institute for Biotechnology; and Yakeel Quiroz of Harvard Medical School also contributed research to this study.

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**About UC Santa Barbara**

The University of California, Santa Barbara is a leading research institution that also provides a comprehensive liberal arts learning experience. Our academic community of faculty, students, and staff is characterized by a culture of interdisciplinary collaboration that is responsive to the needs of our multicultural and global society. All of this takes place within a living and learning environment like no other, as we draw inspiration from the beauty and resources of our extraordinary location at the edge of the Pacific Ocean.