

Hereditary pancreatitis rates 70 times higher among Australian Indigenous people

May 23 2022



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The estimated prevalence of hereditary pancreatitis in South Australia is

higher than in Europe, particularly among Indigenous young people, according to research published today by the *Medical Journal of Australia*.

South Australian researchers, led by Mr. Denghao Wu, a Ph.D. candidate at the University of Adelaide, working at the Royal Adelaide Hospital, analyzed data from 44 people from ten families who received molecular diagnoses of hereditary pancreatitis during 2006–21 (including 25 Indigenous people).

"The estimated prevalence of hereditary pancreatitis was 1.1 per 100,000 population for non-Indigenous and 71 per 100,000 population for Indigenous South Australians," Wu and colleagues reported.

"Among people with adult-onset [chronic pancreatitis](#) admitted to South Australian [public hospitals](#) during 2001–2019, the proportions of Indigenous people (12%) and women (38%) were smaller than we report for hereditary pancreatitis."

Hereditary pancreatitis encompasses a group of rare fibro-inflammatory conditions characterized by recurring bouts of severe abdominal pain that usually require opiate analgesia for relief; initial symptoms are typically reported before the age of ten years. It progresses to exocrine failure and diabetes, and increases the risk of adenocarcinoma of the pancreas.

"Mutations in the cationic trypsinogen (= serine protease 1) gene (PRSS1) on chromosome 7q were identified as major etiological factors in 1996," the authors wrote.

"More than 20 pathogenic PRSS1 mutations have since been described, accounting for 80% of all cases of hereditary pancreatitis in people of European ancestry. Other implicated [genes](#) include those that encode

serine protease inhibitor Kazal-type 1 (SPINK1, on chromosome 5q32), cystic fibrosis transmembrane conductance regulator (CFTR), chymotrypsin C (CTRC), and A-type carboxypeptidase (CPA1). Modifier genes, such as the calcium-sensing receptor gene (CASR), also influence the risk of hereditary pancreatitis."

In the current study, 36 people had mutations in PRSS1, five in SPINK1, and three in both PRSS1 and SPINK1.

"Thirty-eight people (86%) regularly used opioids for [pain relief](#), and 39 described the level of onset pain as moderate or high (89%). The physical impact of pancreatitis-related pain during the preceding four weeks was described as moderate or high by 35 people (79%) and the mental health impact as moderate or high by 18 people (41%). Thirty-two people (73%) reported five or more pancreatitis attacks during the preceding twelve months, and twelve (27%) more than ten attacks," Wu and colleagues wrote.

"Our estimated prevalence of hereditary pancreatitis in South Australia (2.5 per 100,000 population) is much higher than the rates reported by European studies. Further, the estimated prevalence among Indigenous people (71 per 100,000) was almost 70 times that for non-Indigenous South Australians (1.1 per 100,000).

"The high prevalence of hereditary pancreatitis, and particularly of PRSS1 mutations, among Indigenous Australians may have one or more explanations: for example, founder effects, leading to reduced [genetic variation](#); a population bottleneck caused by an abrupt decline in [population size](#), randomly increasing the relative frequency of some gene variants; or a different PRSS1 pseudogene architecture in Indigenous people that facilitates more frequent gene conversion events.

"We have identified the number of South Australians with hereditary

pancreatitis eligible for newer therapies, including total pancreatectomy with islet autotransplantation.

"Hereditary pancreatitis is an important but inadequately recognized cause of early onset pancreatitis in South Australia, particularly in young Indigenous people," Wu and colleagues concluded.

More information: Denghao Wu et al, The clinical and genetic features of hereditary pancreatitis in South Australia, *Medical Journal of Australia* (2022). [DOI: 10.5694/mja2.51517](https://doi.org/10.5694/mja2.51517)

Provided by Wiley

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