

Certain mutations affect kidney disease risk and prognosis

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Certain gene mutations affect individuals' risk of developing a serious kidney condition, as well as their prognosis after being diagnosed with the disease, according to a study appearing in an upcoming issue of the *Journal of the American Society of Nephrology (JASN)*. The findings may help in diagnosing and treating patients with the disease, and in determining the risks that patients' relatives face for developing it as well.

Several abnormalities in genes that encode complement proteins—molecules of the immune system that mediate the first defense against pathogens—are thought to contribute to atypical [hemolytic uremic syndrome](#) (aHUS), a rare kidney condition that results from blood vessel injury, thrombosis, and the destruction of [red blood cells](#). The syndrome causes kidney failure in about half of cases. To see which genes or combinations of genes are important, Marina Noris, PhD, Elena Bresin, MD, Giuseppe Remuzzi, MD (Mario Negri Institute for Pharmacological Research, in Italy) and their colleagues in the European Working Party on Complement Genetics in Renal Diseases screened nearly 800 patients with aHUS and their relatives.

Among the major findings:

- A single mutation in the CFH, C3, or CFB complement genes conferred an increased aHUS risk.
- The concomitant presence of a mutation in the MCP or CFI

complement genes and mutations in other complement genes increased aHUS risk.

- In patients with aHUS, 50% of those with a combination of mutations in MCP and other complement genes developed [kidney failure](#) within three years of aHUS onset compared with 19% of patients with an isolated MCP mutation.
- [Plasma treatment](#) led to remission in patients with combined mutations and single mutations to a similar extent, but [disease recurrence](#) was more likely after a kidney transplant in those with a combined MCP mutation than in those with an isolated MCP mutation.

"The results presented in this paper underline the complexity of the genetics of aHUS and indicate that a complete screening for all susceptibility genes should be performed, in particular before making a conclusive decision about transplantation," said Dr. Noris.

The authors noted that an online database is being updated to provide information on the various mutations that, alone or in combination, are associated with aHUS (www.fh-hus.org).

More information: The article, entitled "Combined Complement Gene Mutations in Atypical HUS Influence Clinical Phenotype," will appear online on February 21, 2013, [doi: 10.1681/ASN.2012090884](https://doi.org/10.1681/ASN.2012090884)

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