

# iNPH—the mystery resolving

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**Idiopathic normal pressure hydrocephalus (iNPH) is characterized clinically by degradation of gait, cognition, and urinary continence. INPH is progressive (Andrén et al, 2014), still probably underdiagnosed (Williams et al, 2019) but potentially treatable by CSF diversion (Kazui et al, 2015). Familial aggregation is a strong indicator of genetic regulation in the disease process iNPH (Fig 1). Enlargement of brain ventricles is associated with failed cerebrospinal (CSF) homeostasis by so far mostly unknown mechanisms. A mutation of the cilia gene *CFAP43* in iNPH family, confirmed by a knocked-out mouse model (Morimoto et al, 2019), allelic variation of *NME8* (Huovinen et al, 2017), a segmental copy number loss in *SFMBT1* in selected iNPH patients (Sato et al, 2016), and current results by Yang et al (2021) indicate that cilia dysfunction is one of the key mechanisms behind iNPH.**

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See also: HW Yang et al (March 2021)

In this issue of *EMBO Molecular Medicine*, Yang et al (2021) report novel genetic and functional findings related to two loss of function deletions in *CWH43* gene in

patients with iNPH. This is one of the most important studies in the field of iNPH research so far, emphasizing the fact that eventually iNPH is not anymore considered as “idiopathic”. Furthermore, these results strongly support the existence of iNPH as an independent disease entity.

The discovery cohort used in the study of Yang et al is relatively small and thus, genetic studies in the larger patient cohorts are expected to identify new potential genetic loci related to iNPH. As with numerous other multifactorial diseases, international collaboration with large multicenter cohorts are needed to uncover the expected rare variants with large effect and more common variants with small effect.

Three out of the eight patients with *CWH43* alteration had potential iNPH-related family history, indicating that this variant can either be *de novo* mutation or that the familiar inheritance is underestimated due to the late onset of the clinical symptoms. Diagnostic criteria of iNPH need to be updated to include also prodromal iNPH, such as asymptomatic ventriculomegaly with features of iNPH on magnetic resonance imaging (AVIM, Kimihira et al, 2020). Identification of genetic variants similar to that now observed with *CWH43* will open a window to decipher the underlying

pathophysiological mechanisms further, paving the way to better treatments and novel prevention strategies in the future.

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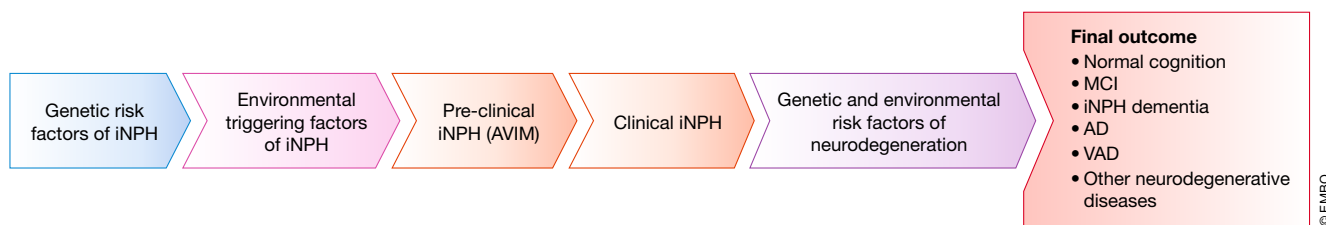


Figure 1. Hypothetical disease progress of iNPH.

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