

Identification of a Mutation in *CNNM4* by Whole Exome Sequencing in an Amish Family and Functional Link between *CNNM4* and *IQCB1*

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Abstract

We investigated an Amish family in which three siblings presented with an early-onset childhood retinal dystrophy inherited in an autosomal recessive fashion. Whole exome sequencing was performed and identified a homozygous nonsense mutation (c.C1813T, p.R605X) in the cyclin and CBS domain divalent metal cation transport mediator 4 (*CNNM4*). Here we show that *CNNM4* interacts with *IQCB1*, which causes Leber Congenital Amaurosis (LCA) when mutated. A truncated *CNNM4* protein starting at R605 significantly increased the rate of apoptosis, and significantly increased the interaction between *CNNM4* and *IQCB1*. Mutation p.R605X may cause Jalili syndrome by a nonsense-mediated decay mechanism, affecting the function of *IQCB1* and apoptosis, or both. Our data, for the first time, functionally link Jalili syndrome gene *CNNM4* to LCA gene *IQCB1*, providing important insights into the molecular pathogenic mechanism of retinal dystrophy in Jalili syndrome.

Results

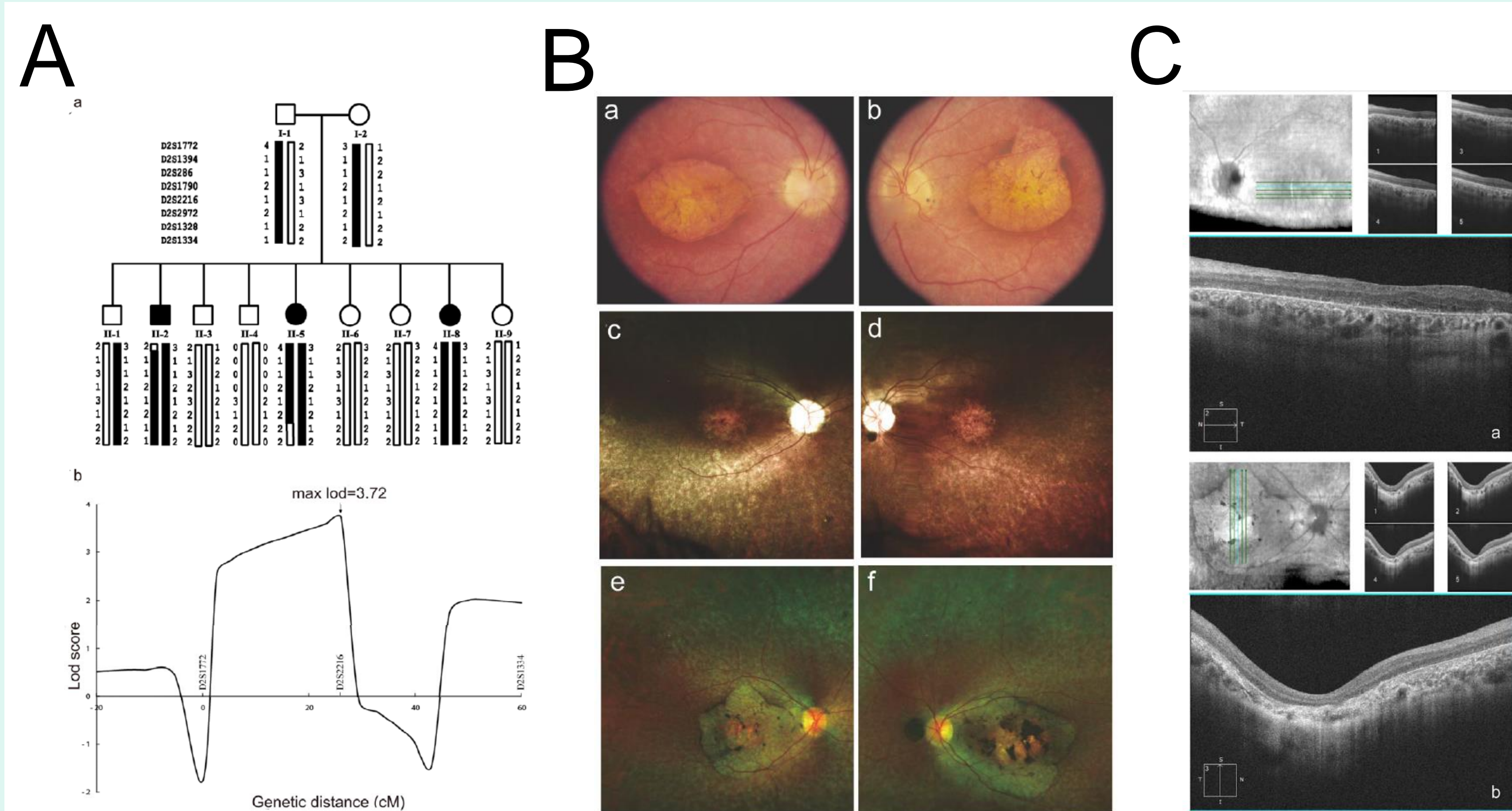


Figure 1 (A) Linkage analysis. (B) Fundus photos of three siblings with Jalili syndrome. (C) Representative OCT.

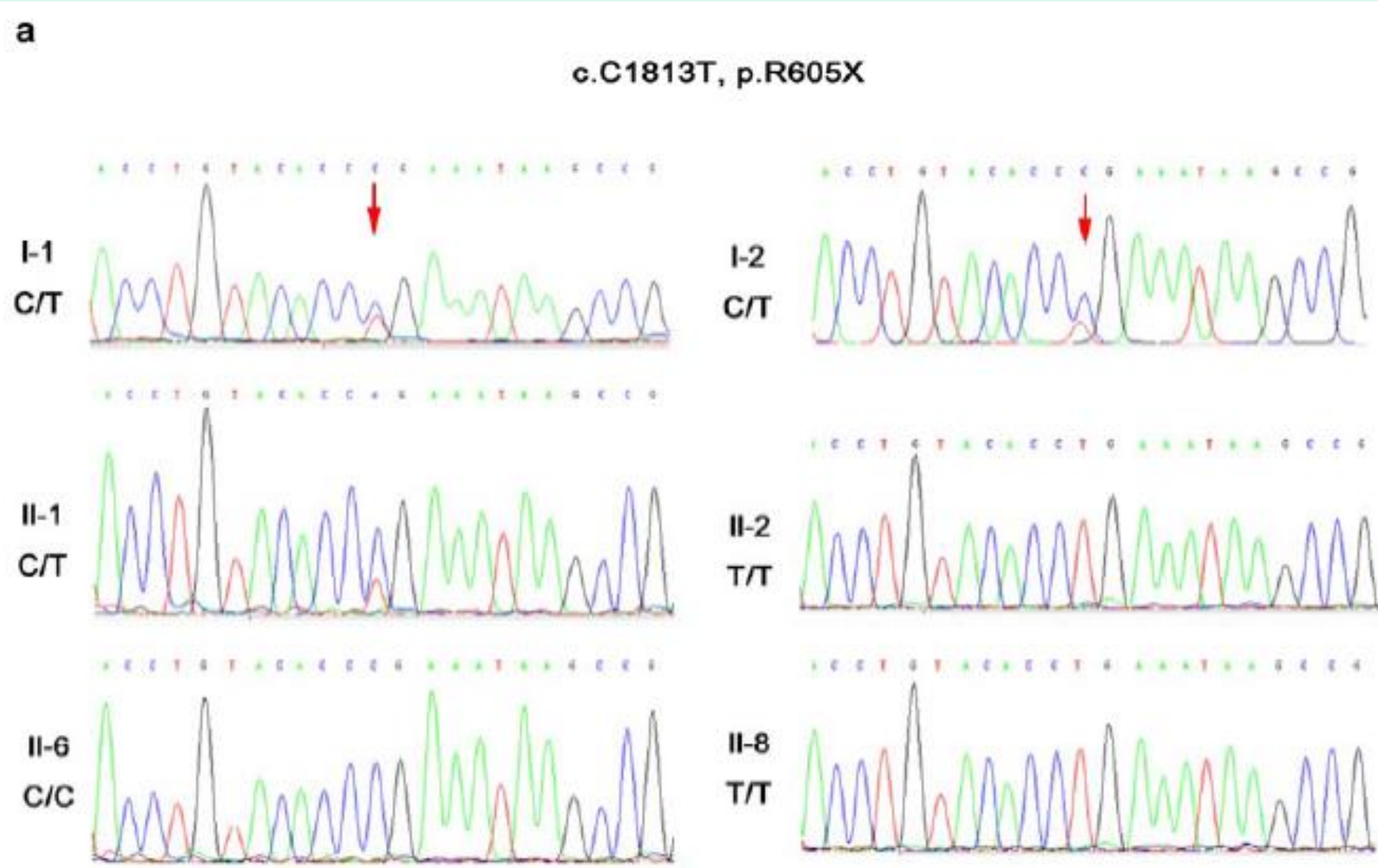


Figure 2 Identification of a *CNNM4* mutation co-segregating with the disease in the Amish family.

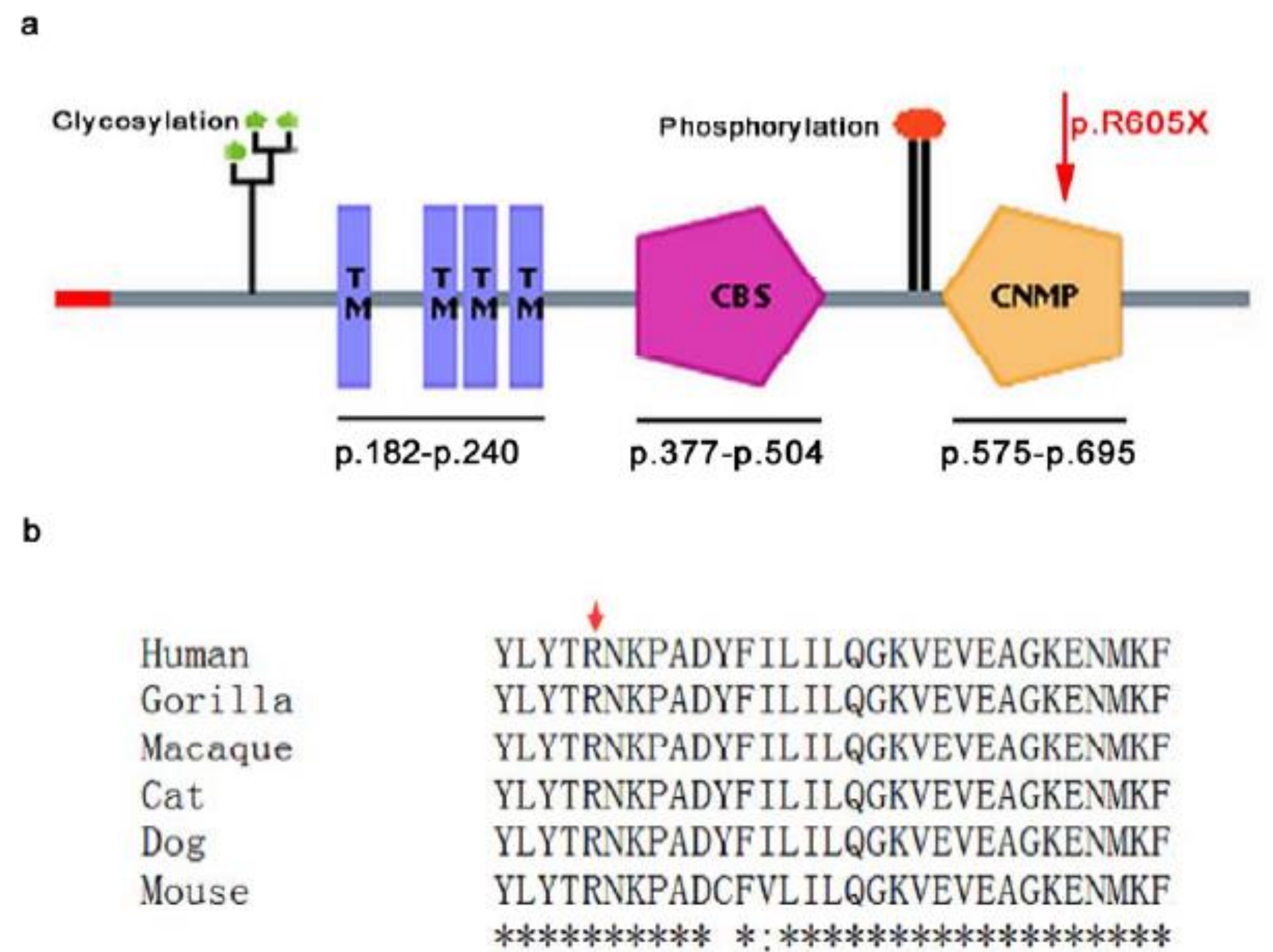


Figure 3 Schematic drawing and multiple sequence alignment of the *CNNM4* protein.

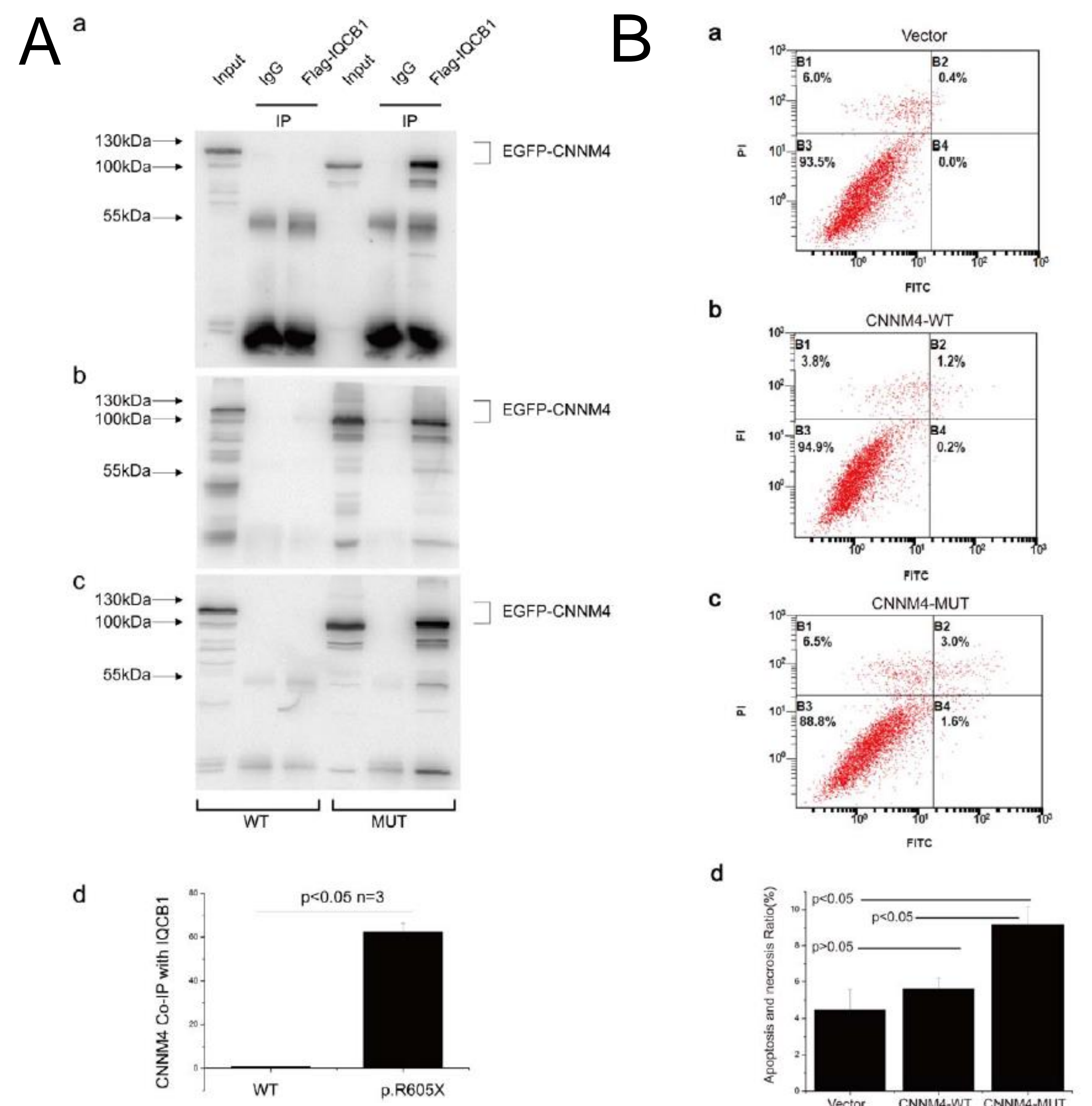


Figure 4 (A) The *CNNM4* protein interacts with *IQCB1* encoded by an LCA-causing gene. (B) The *CNNM4* protein interacts with *IQCB1* encoded by an LCA-causing gene.

Acknowledgements

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