

QRICH1

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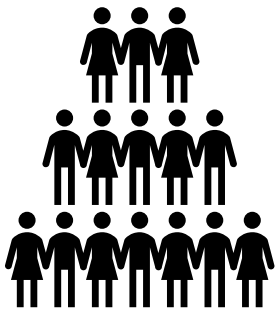
QRICH1-associated developmental delay and intellectual disability in 36 unrelated individuals

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Targeted gene(s)/phenotype under study : *QRICH1*

Abstract : *De novo* pathogenic variants of *QRICH1* (Glutamine-rich protein 1, OMIM #617387) has recently been described in **five patients** (Ververi et al. 2018; Lui et al. 2019). The variants have been associated with developmental delay and intellectual disability, mild facial dysmorphism and chondrodysplasia in some cases. Through Gene Matcher (Genematcher.org) we have now identified **19 further patients** with *QRICH1*-variants. Currently, we are in the process of defining the phenotype-genotype spectrum of *QRICH1*-related disorders in the patient cohort (**a total of 24** including the published cases) and preparing a manuscript. We welcome further cases to this study to reach a better understanding of this rare disorder. We aim to close inclusion of further cases 15th July 2020.

Coordinating clinician/researcher: Zeynep Tümer



CURRENT STATUS

- The current number of unpublished individuals are now 27 – two of the patients are through ITHACA collaboration (Prof. Favier's group)
- 4 new patients reported meanwhile (Föhrenbach et al. 2020).
- **TOTAL** number of individuals will be 36 (including 9 published)

QRICH1 has 11 isoforms and all encode the same 776 amino acid protein.

- glutamine (Q) rich region
- **CARD domain** (Caspase activation and recruitment domain): uncharacterized
- **DUF3504 domain** (domain of unknown function): a protein-protein interaction domain found in several proteins involved in apoptosis, inflammation and immune responses.

**Protein Truncating
Variants: n=23**

p.(Arg16*)
p.(Gln46Serfs*200)
p.(Gln46_Gln47delinsHis*)
c.310-2A>C; p.?

p.(Gln181*)
p.(Gln275*)
p.(Ser278Leufs*25)
p.(Gly306Argfs*48)
p.(Asp321Thrfs*47)
p.(His329Thrfs*39)
p.(Gln420*)
p.(Pro432Thrfs*23)
p.(Gln460*)
p.(Arg511*)
p.(Arg536*)
p.(Tyr543Metfs*23)
p.(Phe552Serfs*14)
c.1787-2A>G; p.?
p.(Glu605Glyfs*25)
c.1896-2A>G; p.?
p.(Arg652Alafs*9)
p.(Arg652*)
p.(Trp739*)



p.(Met252Ile)
p.(Pro284Leu)

p.(His394Asp)
p.(Gln435Arg)
p.(Gly527Arg)
p.(Tyr550Cys)
p.(Tyr574Asp)
p.(Val603Leu)
p.(Phe628Leu)
p.(Ser736Asn)

**Missense Variants:
n=10**

28 *de novo*, 3 paternally inherited (c.961del, c.1655del, c.310-2A>C), 2 being investigated.

Collaborators from 27 different locations *(author list not complete yet and not in the final order)*

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- Anya Revah-Politi, Kwame Anyane-Yeboah - Columbia University
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- Natasha Brown, Smitha Kumble - Australia
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Article is in preparation and not submitted yet