

Title: *KCNQ3*-Related Disorders *GeneReview* Table 3

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Note: The following information is provided by the authors listed above and has not been reviewed by *GeneReviews* staff.

Table 3. Overview of the Available Genetic, Clinical, and Functional Data from Families Carrying *KCNQ3* Mutations: BFNE and BFIE

DNA Nucleotide Change	Amino Acid Change	Localization	BFNE	BFIE	Additional Clinical Data	Functional Effects	Reference
c.895G>A	p.Glu299Lys	S5-S6 linker region; Pore	+	-	This variant segregates with the BFNE phenotype in 4/5 sibs of a three-generation family; possibly contributes to the Rolandic epilepsy occurring in this family	Reduction in current amplitude	Neubauer et al [2008]; Hahn & Neubauer [2009]
c.914A>G	p.Asp305Gly		+	-		Reduced current amplitude of heteromeric channels	Singh et al [2003]
c.925T>C	p.Trp309Arg		+	-		Homomeric channels lacked potassium current and heteromeric channels displayed a dramatic reduction of current	Hirose et al [2000]; Uehara et al [2008]; Sugiura et al [2009]
c.929G>T	p.Gly310Val		+	-		---	Charlier et al [1998]
c.950T>C	p.Ile317Thr		+	-	Moderate psychomotor delay	Reduction in current amplitude	Soldovieri et al [2014]
c.988C>T	p.Arg330Cys		+	-	The same variant occurred in two families	---	Li et al [2006]; Li et al [2008]; Fister et al 2013]

DNA Nucleotide Change	Amino Acid Change	Localization	BFNE	BFIE	Additional Clinical Data	Functional Effects	Reference
c.1403A>G	p.Asn468Ser	C-terminus	-	+	Three siblings affected	No effect; possibly a non pathogenic variant	Singh et al [2003]
c.1720C>T	p.Pro574Ser		+	-	Found in a typical BFNE family also carrying a <i>KCNQ2</i> missense mutation likely pathogenic. Also detected in other complex phenotypes (see Table 2); in fact, it has been detected in four individuals with rolandic epilepsy and in one with rolandic epilepsy and moderate psychomotor delay, in 8/455 patients with idiopathic generalized epilepsies (IGE; absent in 454 controls), and in three patients with autism spectrum disorders, with no additional neurologic features. In the latter study, this variant has been detected at low frequency (0.2-0.4%) also in control individuals.	No difference versus wt; possibly a variant of unknown significance; reduced current amplitude only when co-expressed with Kv7.5	Neubauer et al [2008]; Hahn & Neubauer, [2009]; Miceli et al [2009]; Lemke et al [2012]; Gilling et al [2013]
c.2338C>T	p.Arg780Cys		-	+		---	Zara et al [2013]
c.2462A>G	p.Asn821Ser		+	-	Found in a typical BFNE family also carrying a <i>KCNQ2</i> deletion/insertion likely pathogenic; the <i>KCNQ3</i> variant does not co-segregate with the disease	No difference versus wt; possibly a variant of unknown significance	Bassi et al [2005]

BFNE: benign familial neonatal epilepsy

BFIE: benign familial infantile epilepsy

FS: febrile seizures

Reference sequences: NM_004519.3 (*KCNQ3*), cDNA numbering begins with +1 as the A of ATG initiation codon.

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