

A Study among the Genotype, Functional Alternations, and Phenotype of 9 *SCN1A* Mutations in Epilepsy Patients

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Supporting information

Supplementary Table 1 Summary of the studied SCN1A variants and used mutagenesis primers.

Variant	Longer transcript ENST00000303395.8		Shorter transcript ENST00000637988.1		Region	Mutagenesis primers (5' - 3')
	cDNA position	Protein change	cDNA position	Protein change		
D249E	c.747T>G	p.Asp249Glu	c.747T>G	p.Asp249Glu	DIS4-S5	GAAGCTCTCAGAGGTAATGATCCT TTCACAGACTGGATCAGGGCT
E788K	c.2362G>A	p.Glu788Lys	c.2329G>A	p.Glu777Lys	DIIS1-S2	ATGGCCATGGAGCACTATCCA GAAAAGAGTATTTAAGACAATACAGATGGT
W384*	c.1151G>A	p.Trp384*	c.1151G>A	p.Trp384*	DIS5-S6	GACTCAGGACTTCTAGGAAAATCTTTATC ATTAGTCGAAACAAGGACAAAAAAGC
E78D	c.234G>T	p.Glu78Asp	c.234G>T	p.Glu78Asp	N-terminal	CAGAGCCCTGGATGACCTGGACC ACACCATCTCTGGAGGAATGT
T1934I	c.5801C>T	p.Thr1934Ile	c.5768C>T	p.Thr1923Ile	C-terminal	TTTAAAGCGAATTGTAAAAACAAGC AGGTGGCGTCTGTAAGCAC
R1596C	c.4786C>T	p.Arg1596Cys	c.4753C>T	p.Arg1585Cys	DIVS2-S3	AACTCATCTCTATGCCATTATTATTT TCAGTACACACTCTCCAGTAAATAGC
M909K	c.2726T>A	p.Met909Lys	c.2693T>A	p.Met898Lys	DIIS5	CCGTGGTCGGCAAGCAGCTCTTTGGT CAAAAATGAAGACGATGATGGCCAAG
E78*	c.232G>T	p.Glu78*	c.232G>T	p.Glu78*	N-terminal	CAGAGCCCTGTAGGACCTGGAC ACACCATCTCTGGAGGAATGT
E1587K	c.4759G>A	p.Glu1587Lys	c.4726G>A	p.Glu1576Lys	DIVS2	GCTATTTACTGGAAAGTGTGTACTGAA ACAATGAACACCAGATTGATGCGT