

Brugada syndrome: etiological, forms and genetic basis- 2008

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1) Sporadic forms without genetic mutation: » 70% to 82% of the cases.

2) With genetic mutation: » 18% to 30% of cases

(2-1) Brugada syndrome type 1 or B1: OMIN number: 601144, Locus: 3p21; Gene: mutation in alpha subunit of the sodium channel (SCN5A Nav1.5) gene¹.

(2-2) Brugada syndrome type 2 or B2: Locus: 3p24, gene loss-of-function mutations in gene encoding the cardiac L-type calcium channel: Mutation in glycerol-3 phosphate dehydrogenase-1 like (GPD1L ^{2; 3}). The entity is associated with progressive conduction disease, a low sensitivity to procainamide, and a relatively good prognosis⁴.

(2-3) Brugada syndrome type 3 or B3: Locus: 12.p 13.3 Gene: mutation in CACNA1C.Ca_v1.2 gene⁵.

(2-4) Brugada syndrome type 4 or B4: Locus: 10p12.33, Gene: loss-of-function mutations in gene encoding the cardiac L-type calcium channel: The CACNB2b.CAvb2b gene. The entity consists of an ST-segment elevation in the right precordial ECG leads, a shorter-than-normal QT interval (≤ 370 ms), and a familial history of sudden cardiac death⁵.

Observation: Types 3 and 4 are very rare, and probably have and overlap genotype (shorter-than-normal QT interval (≤ 370 ms) interval + Brugada syndrome ECG phenotypes)

(2-5) Brugada syndrome type 5 or B5: OMIN number: 604433, *KCNE3* which coassembles with *KCND2* beta subunit act in

the modulation of I_{to} potassium outward current in the human heart. Mutations in *KCNE3* can underlie the development of the Brugada syndrome⁶

(2-6) Brugada syndrome type 6 or B6: OMIN number: 600235, SCN1B beta 2 subunit of the sodium channel SCN5A. Watanabe et al⁷ investigated SCN1B, which encodes the function-modifying sodium channel beta1 subunit, in 282 probands with Brugada syndrome and in 44 patients with conduction disease, none of whom had SCN5A mutations. The authors identified 3 mutations segregating with arrhythmia in 3 kindreds. Two of these mutations were located in a newly described alternately processed transcript, beta1B. Both the canonical and alternately processed transcripts were expressed in the human heart and were expressed to a greater degree in Purkinje fibers than in heart muscle, consistent with the clinical presentation of conduction disease. Sodium current was lower when NaV1.5 was coexpressed with mutant beta1 or beta1B subunits than when it was coexpressed with WT subunits. These findings implicate SCN1B as a disease gene for human arrhythmia susceptibility.

3) Atypical Brugada syndrome or idiopathic ventricular fibrillation with J waves in inferior, lateral or inferior lateral leads

Until to day, two identified mutations

(3-1) SCN5A⁹

(3-2) KCNJ8_{KATP}^{10; 11} The ECGs have an elevation of the QRS-ST junction of at least 0.1 mV from baseline in the inferior or lateral lead, manifested as QRS slurring or notching. Among patients with a history of idiopathic ventricular fibrillation, there is an increased prevalence of early repolarization.

4) “Acquired” or secondary forms¹²

(4-1) Drug-induced like congenital LQTS

(4-2) Electrolytes abnormalities-induced

(4-3) Acute ischemia-induced

(4-4) Fever-induced

(4-5) Hypothermic-induced

(4-6) Mechanical compression of right ventricle outflow tract - induced

(4a) without genetic background

(4b) with genetic background

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