

Somatic mutations in adrenals from patients with primary aldosteronism not cured after adrenalectomy suggest common pathogenic mechanisms between unilateral and bilateral disease

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Abstract

Objective: Primary aldosteronism (PA) is the most common form of secondary and curable hypertension. Different germline and somatic mutations are found in aldosterone producing adenoma (APA) and familial forms of the disease, while the causes of bilateral adrenal hyperplasia (BAH) remain largely unknown. Adrenalectomy is the recommended treatment for patients with APA; however, 6% of patients are not cured and show persistent PA after surgery suggesting BAH. The objective of this study was to analyze clinical data of patients with APA without biochemical success after adrenalectomy as well as the histological and genetic characteristics of their adrenal glands. **Design and Methods:** Clinical data of 12 patients without biochemical cure were compared to those from 39 PA patients with hormonal cure after surgery. Histological, morphological and genetic characterization of the adrenals was carried out by CYP11B2 and CYP11B1 immunostaining and by CYP11B2-guided NGS. **Results:** Patients with absent hormonal cure displayed longer duration of arterial hypertension and lower lateralization index of aldosterone production. In 10 patients, APA expressing CYP11B2 were identified. No difference in histological and morphological characteristics was observed between patients with or without hormonal cure. Somatic mutations in APA driver genes were identified in all CYP11B2 positive APA; *CACNA1D* mutations were the most frequent genetic abnormality. **Conclusions:** Patients with absent biochemical cure were diagnosed later and exhibited lower lateralization index of aldosterone production, suggesting asymmetric aldosterone production in the context of BAH. Somatic mutations in adrenal glands from those patients indicate common mechanisms underlying BAH and APA.

1 **Introduction**

2 Primary aldosteronism (PA) is the most frequent cause of endocrine hypertension, with
3 a prevalence of $\approx 5\%$ of hypertensive patients in primary care and 10% to 20% in reference
4 centers (1-3). PA patients exhibit hypertension associated with high levels of plasma
5 aldosterone, low levels of plasma renin and in some cases hypokalemia. The excessive
6 aldosterone production is attributable in the majority of cases to an unilateral aldosterone
7 producing adenoma (APA) or to a bilateral adrenal hyperplasia (BAH) (4).

8 In the last decade, different studies identified somatic and germline mutations in genes
9 coding for ion channels and ATPases as responsible for APA and familial forms of PA (5-12).
10 The discovery of these mutations highlighted the central role of calcium signaling in the
11 pathogenesis of the disease (13). Recently, the use of next generation sequencing (NGS)
12 performed on DNA from CYP11B2 positive nodules extracted from formalin fixed paraffin
13 embedded (FFPE) tissues has allowed the identification of somatic heterozygous mutations in
14 *KCNJ5* (coding for the potassium channel GIRK4), *ATP1A1* (coding for the $\alpha 1$ subunit of the
15 Na^+/K^+ -ATPase), *ATP2B3* (coding for the plasma membrane Ca^{2+} -ATPase, type 3 PMCA3),
16 *CACNA1D* (encoding the Cav1.3 voltage dependent calcium channel), *CACNA1H* (encoding
17 the Cav3.1 voltage dependent calcium channel), *CLCN2* (coding for the chloride channel ClC-
18 2), and *CTNNB1* (coding for β -catenin) in more than 88% of APA (14-17). However, the genetic
19 causes of BAH remain largely unknown. Analysis of 15 adrenals from patients with BAH has
20 shown that BAH may result from the accumulation or enlargement of aldosterone producing
21 cell clusters (APCC) harboring somatic mutations, particularly in the *CACNA1D* gene (18).

22 The goal of PA treatment is the normalization of blood pressure, together with correction
23 of hypokalaemia and biochemical abnormalities (aldosterone and renin) in unilateral forms, or
24 the efficient blockade of the mineralocorticoid receptor, since PA is associated with increased
25 risk of cardiovascular complications independently of blood pressure levels (19-23). Surgical

26 adrenalectomy is the recommended treatment for lateralized PA (4, 24) . For this purpose,
27 correct subtyping of PA is mandatory to identify patients with unilateral disease who can be
28 cured after adrenalectomy. Many studies have shown an improvement of hypertension, a
29 decrease in cardiovascular and metabolic risk factors, and a reduction in mortality after
30 adrenalectomy compared with mineralocorticoid receptor antagonist treatment alone (25) (26).
31 Despite the recommendations, approximately 6% of PA patients classified pre-surgery as
32 having unilateral PA do not achieve complete biochemical cure after adrenalectomy (27). These
33 patients exhibit lower lateralization index on adrenal vein sampling (AVS), suggesting bilateral
34 but asymmetric aldosterone production and a misdiagnosed BAH (28).

35 The objective of this study was to analyze clinical data of patients with APA without
36 biochemical success after adrenalectomy and to investigate the histological and genetic
37 characteristics of their adrenal glands.

38 **METHODS**

39 *Patients*

40 Among patients who underwent adrenalectomy between 2008 and 2018 at the Hôpital Européen
41 Georges Pompidou recruited within the COMETE-HEGP protocol, 12 patients with partial or
42 absent biochemical cure after adrenalectomy using the PASO criteria (29) were analyzed.
43 Partial biochemical success: correction of hypokalaemia (if present pre-surgery) and a raised
44 aldosterone-to-renin ratio with $\geq 50\%$ decrease in baseline plasma aldosterone concentration
45 and/or abnormal but improved post-surgery confirmatory test result. Absent biochemical
46 success: persistent hypokalaemia or persistent raised aldosterone-to-renin ratio, or both, with
47 failure to suppress aldosterone secretion with a post-surgery confirmatory test. Data from 39
48 PA patients with biochemical cure after adrenalectomy and complete clinical, biochemical,
49 histological and genetic exploration (16) were used as controls. The term “non-cured group”
50 used in this article corresponds to patients with partial or absent biochemical success. Seven
51 patients were classified as having absent biochemical success after adrenalectomy and five
52 patients exhibited partial biochemical success (Table S2). All patients exhibited visible adrenal
53 adenoma or adrenal enlargement on CT scan. The study was approved by the French Research
54 ethics committee (Comité de Protection des Personnes, CPP) under authorization number CPP
55 2012-A00508-35. Written consent has been obtained from each patient or subject after full
56 explanation of the purpose and nature of all procedures used. Methods used for screening and
57 subtype identification of PA were performed according to institutional and Endocrine Society
58 guidelines (4, 30-32). All patients underwent adrenal vein sampling (AVS) to differentiate
59 between unilateral and bilateral aldosterone hypersecretion. AVS sampling was performed
60 simultaneously in both adrenal veins and without pharmacologic stimulation (32, 33); the
61 complete protocol is described in the supplementary data. Baseline and follow up clinical and

62 biochemical characteristics of PA patients without biochemical success after adrenalectomy are
63 described in supplementary tables S1 and S2.

64 ***Immunohistochemistry and pathological analysis***

65 Each paraffined adrenal block from the non cured patients (absent biochemical success) was
66 analysed entirely to have a precise morphological and cellular analysis. Sections (4- μ m thick)
67 were deparaffinised in xylene and rehydrated through graded ethanol. Hematoxylin/Eosin, 11 β -
68 hydroxylase (CYP11B1) and aldosterone synthase (CYP11B2) immunohistochemistry were
69 performed as previously described (16). Images were acquired in a Lamina Slide Scanner from
70 Perkin Elmer and analysed on the Cochin Image Database (Institut Cochin, France). Zona
71 glomerulosa (ZG) hyperplasia was defined as the presence of a continuous character of the ZG,
72 or, in case of discontinuity of the ZG, ZG thickness $\geq 200 \mu\text{m}$ (16, 34). CYP11B1 and
73 CYP11B2 staining was quantified as previously described (16). Percentages of aldosterone-
74 synthase and 11 β -hydroxylase expressing cells are reported as: 0, absent; 1, 1%-33%; 2, 34%-
75 66%; 3, 67%-100%.

76

77 ***DNA isolation and sequencing***

78 Somatic DNA of APA was extracted from fresh frozen tissue (FFT) using QIamp DNA midi
79 kit (Qiagen, Courtaboeuf Cedex, France) and Sanger sequencing targeting hot spot regions of
80 *KCNJ5*, *CACNA1D*, *ATP1A1*, *ATP2B3* and *CTNNB1* genes was performed as previously
81 described (35). For negative samples on Sanger sequencing and samples not sequenced
82 previously, CYP11B2 immunohistochemistry-guided NGS (CYP11B2 IHC-guided NGS) was
83 performed as previously described (16). Before DNA/RNA extraction of FFPE tissue, APA and
84 aldosterone producing nodules were identified by CYP11B2 IHC. Based on the CYP11B2 IHC,
85 the areas of interest were delimited and isolated for DNA/RNA extraction by scraping unstained
86 FFPE sections guided by the CYP11B2 IHC slide using a scalpel under a Wild Heerbrugg or

87 Olympus microscope. DNA was extracted from FFPE sections using AllPrep DNA/RNA FFPE
88 kit (Qiagen). NGS was performed using a NGS kit, covering all coding exons and intron-exon
89 junctions of the *KCNJ5* (NM_000890), *ATP1A1* (NM_000701), *ATP2B3* (NM_0010001344),
90 *CTNNB1* (NM_001904), *CACNAID* (NM_001128839.2 and NM_000720), *APC*
91 (NM_000038.5), *CACNAIH* (NM_021098 and NM_001005407), *PRKACA* (NM_002730) and
92 *ARMC5* (NM_002730) genes (MASTR_PA kit, Multiplicom/Agilent, Santa Clara, CA USA)
93 (16).

94

95 ***Statistical analyses.***

96 Quantitative variables are reported as medians and interquartile range. Comparisons between
97 groups were done with unpaired t-test when Gaussian distribution or the Mann-Whitney test
98 when no Gaussian distribution. Categorical variables are reported as numbers and and
99 compared with Fisher's exact test. A p value < 0.05 was considered significant for comparisons
100 between 2 groups.

101

102 **Results**

103 **Comparison of clinical data between patients with or without biochemical success after**
104 **adrenalectomy**

105 Baseline and follow up clinical and biochemical data are summarized in Table 1. At follow up
106 (8.5 (interquartile range 4.2,15) months), patients with partial or absent biochemical success
107 exhibited higher systolic ($p=0.003$) and diastolic blood pressure ($p=0.008$), higher plasma
108 aldosterone levels ($p<0.0001$), lower renin levels ($p=0.007$), lower plasma K^+ concentration
109 ($p<0.0001$) and higher number of anti hypertensive drugs ($p=0.003$) compared with patients
110 with complete biochemical success. The proportion of female patients was lower, but not
111 significant, in the biochemically non-cured group compared with the cured group (16.7% vs
112 33%, $p= 0.47$). The duration of hypertension before PA diagnosis was longer in patients with
113 absent biochemical success compared with patients with complete biochemical success (11
114 (6,15) years vs 3 (1,8) years, $p= 0.002$). While no differences were observed in pre-operative
115 plasma aldosterone and plasma renin levels, patients with absent biochemical success displayed
116 a lower lateralization index at AVS ($p=0.02$). No significant differences were found in the other
117 baseline parameters.

118

119 ***Histological analysis***

120 No difference in the adenoma size was observed between patients with complete biochemical
121 success (10.5 (7,14) mm) versus absent biochemical success (14.5 (10,15) mm) after
122 adrenalectomy ($p=0.09$) (Table S3). Zona glomerulosa hyperplasia was observed in 7 out of 12
123 adrenals from biochemically non-cured PA patients (Table 2). CYP11B2 staining was
124 performed on the entire FFPE blocs of adrenals from biochemically non-cured patients in order
125 to identify aldosterone producing nodules. 10 adrenals showed one CYP11B2 positive adenoma
126 of at least 5 mm (Table 2). Micronodular hyperplasia was observed in two of these adrenals

127 (patients 1 and 4) and CYP11B2 positive secondary micronodules were observed in three
128 adrenals (patients 2, 8 and 9). Aldosterone producing cell clusters (APCCs) were observed in 8
129 out of 12 adrenals, ranging from 1 to 4 APCCs per adrenal (Table 2). In two adrenals, no
130 adenomas expressing CYP11B2 were observed. The first adrenal (patient 6) exhibited
131 CYP11B2 negative micronodular hyperplasia and three APCCs. In the second adrenal (patient
132 7) one adenoma non expressing CYP11B2 and two APCCs were observed.

133 Given the previously documented intratumoral heterogeneity of APA, the number of cells
134 expressing CYP11B2 and CYP11B1 was quantified in each APA (Table 2). Five APAs showed
135 between 67% and 100% of cells expressing CYP11B2, two exhibited 33% to 66% of cells
136 expressing CYP11B2, while in three APA, 1% to 33% CYP11B2 expressing cells were
137 observed. Concerning CYP11B1 expression, two adenomas were composed of 33% to 66% of
138 cells expressing CYP11B1 and in eight adenomas 1 to 33% of CYP11B1 expressing cells were
139 observed. In comparison with APA from PA patients with complete biochemical success after
140 adrenalectomy, there was no difference in the number of cells expressing CYP11B2 and
141 CYP11B1 (Table S3).

142

143 ***Identification of somatic mutations in the absent biochemical success group.***

144 Among the 10 CYP11B2 positive adenomas from the absent biochemical success group,
145 analysis of hot spot mutations in APA driver genes was performed by Sanger sequencing in
146 fresh frozen adenoma tissue (FFT) from four patients. One somatic *KCNJ5* mutation
147 (c.451G>C/p.Gly151Arg) was identified in one adenoma and somatic *CACNAID* mutations
148 were identified in three patients: p.Arg990Gly (c.2968C>G), p.Gly403Arg (c.1207G>C), and
149 p.Val1373Asp (c.4117G>A). CYP11B2 IHC-guided NGS was performed in CYP11B2 positive
150 FFPE samples from five adenomas without previous sequencing. *CACNAID* mutations were
151 identified in three patients (c.1207G>C/p.Gly403Arg, c.2968C>G/p.Arg990Gly, and

152 c.3458T>A/p.Val1153Asn), the same *KCNJ5* mutation (c.451G>C/P.Gly151Arg) was
153 identified in two patients, and the *ATP1A1* mutation p.Leu104Arg ((c.311T>G) was identified
154 in one patient. Targeted NGS was also performed in one CYP11B2 negative adenoma and no
155 mutations in APA driver genes were observed. In total, somatic mutations in APA driver genes
156 were observed in all CYP11B2 expressing adenomas from 12 patients (Table 3).
157 Differently from patients with complete biochemical success after adrenalectomy with a higher
158 prevalence of somatic *KCNJ5* mutations (38.4% within the 39 patients analyzed), *CACNAID*
159 mutations were the most prevalent genetic abnormality in APA from patients without
160 biochemical success (60.0%). Adenomas harbouring *CACNAID* mutations (median 9 mm)
161 were smaller than adenomas harbouring *KCNJ5* mutations (median 14 mm) (Table S4). No
162 correlation was observed between the number of CYP11B2 or CYP11B1 expressing cells and
163 the mutation status (Table S4).
164

165 **Discussion**

166 In the present study, we report for the first time the presence of somatic mutations in aldosterone
167 producing nodules from patients with absent biochemical success after adrenalectomy. Analysis
168 of clinical data from 12 PA patients who had undergone adrenalectomy with partial or absent
169 biochemical success after surgery also showed a decreased lateralization index of aldosterone
170 production on AVS but no differences were observed in the histological analysis of CYP11B2
171 positive adenomas and adrenals from patients completely cured or non-cured after
172 adrenalectomy.

173 Patients were recruited within a single referral center for hypertension in France and performed
174 histological and genetic characterization of their adrenals. The patients were selected
175 accordingly to an international multicentric consensus for classifying surgical outcome and
176 follow-up of patients with unilateral primary aldosteronism (PASO) (29). This consensus has
177 provided the uniformity of clinical and biochemical criteria of PA outcome in different
178 specialized centres, allowing to identify factors influencing the success of adrenalectomy. We
179 focused this study on patients with unilateral PA who did not reach complete biochemical
180 success after adrenalectomy. These patients could represent a group of patients with a bilateral
181 form of PA misdiagnosed during the AVS, explaining the absence of PA cure after surgery, as
182 suggested by some studies (28, 29).

183 It has been previously shown that younger patients and women displayed a more favourable
184 outcome after adrenalectomy (29, 36-38). In the present study, we observed a higher percentage
185 of men among patients with absent biochemical success compared with a previous published
186 cohort of 39 patients with unilateral PA with complete biochemical success after surgery,
187 confirming previous studies. In addition to female sex, other variables were described as
188 predictors for favourable clinical outcome, including short term hypertension (≤ 7 years),

189 absence of overweight, low number of antihypertensive drugs, higher baseline blood pressure
190 and no history of diabetes mellitus (29, 37). Although we did not observe differences in blood
191 pressure at baseline, number of hypertensive drugs or BMI between patients with or without
192 biochemical success after surgery, biochemically non-cured patients were diagnosed later and
193 presented a longer duration of hypertension. The duration of hypertension may indicate not only
194 a predictor for clinical but also for biochemical outcome of unilateral PA. In the present study,
195 we observed lower lateralization index of aldosterone at AVS in patients without biochemical
196 success. This finding was previously observed in a larger cohort of non-cured PA patients (28),
197 and together with the longer duration of hypertension before PA diagnosis suggests a bilateral
198 disease with asymmetric aldosterone production.

199 Remarkably, in 10 out of 12 cases, one functional adenoma expressing CYP11B2 was observed
200 in the resected adrenal of patients without biochemical success after adrenalectomy. This
201 finding confirms lateralization of aldosterone production at AVS in the majority of the
202 investigated patients, suggesting asymmetric bilateral aldosterone production; for two patients,
203 no adrenal adenoma expressing CYP11B2 was identified and the source of autonomous and
204 excessive aldosterone production was not identified, suggesting a misdiagnosis of PA subtyping
205 before surgery. Zona glomerulosa hyperplasia as well as heterogeneous CYP11B2 and
206 CYP11B1 expression in APA were observed in the adrenals from non-cured patients,
207 accordingly to previously described adrenals from lateralized PA (16, 39). In contrast to a
208 multicentric study of patients with lateralized PA biochemically not-cured after adrenalectomy
209 (28), we did not observe histological findings associated with biochemical success. The absence
210 of relationship between histopathology and biochemical cure was also observed in an Italian
211 cohort of PA patients (40). The difference between these studies may imply differences between
212 patients recruited in different centers, an observation that has already been reported for clinical

213 characteristics of patients with PA (35), or may represent less power to identify these
214 differences due to a smaller sample size.

215 Remarkably, we report a high prevalence of somatic mutations in APA driver genes in
216 adenomas from patients with absent biochemical cure. *CACNA1D* mutations were the most
217 frequent genetic abnormality in APAs from patients with partial or absent success after
218 adrenalectomy, which were not associated with ethnic background as previously reported (15).
219 A higher frequency of somatic *CACNA1D* mutations was observed previously in the context of
220 bilateral adrenal hyperplasia, with the accumulation or enlargement of APCCs harbouring
221 somatic mutations in APA-driver genes (18), suggesting common pathogenic mechanisms at
222 the origin of BAH and APA. Our data showing the presence of somatic mutations (in particular
223 *CACNA1D* mutations) in APA from patients not cured after adrenalectomy, also presenting
224 with lower lateralization index and longer duration of HT, support this hypothesis.

225 Only a small number of APA harboured somatic *KCNJ5* mutations, in contrast with what
226 observed in patients with complete biochemical cure in this study and studies analysing a large
227 number of patients from French or European cohorts (16, 35, 41-44). While the small number
228 of *KCNJ5* mutations may be associated with the higher number of male patients without
229 biochemical success, the association of *KCNJ5* mutations and favourable outcome was
230 described previously, independently from the favourable impact of female sex (45, 46). While
231 associations between the mutation status and histological findings are well characterized in
232 APA (16, 47), no associations were observed in the present study, probably due to the small
233 number of adenomas within each genotype.

234 In conclusion, 10 out of 12 adrenals from PA patients without biochemical success after
235 adrenalectomy exhibited *CYP11B2* expressing adenomas carrying somatic mutations,
236 confirming the diagnosis of APA. However, these patients were diagnosed later and exhibited

237 lower lateralization index of aldosterone production at AVS, suggesting asymmetric
238 aldosterone production in the context of BAH. The identification of somatic mutations in
239 adrenal glands from those patients suggests a possible continuum between bilateral and
240 unilateral disease in a subset of patients, supporting common mechanisms underlying BAH and
241 APA.

242

243 **Declaration of interest**

244 The authors have nothing to disclose

245

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Table 1. Baseline and follow up clinical and biochemical data of PA patients with or without biochemical success after adrenalectomy.

	Non biochemical success (12)	Biochemical success (39)	p value
Age (ys)	47 (39,51)	44 (35,50)	0.90
Gender F/M (F%)	2/10 (16.7%)	13/26 (33.3%)	0.03
BMI (kg/m ²)	31.2 (29,35)	28.5(25,33)	0.182
HT duration (ys)	12 (6,15)	3 (1,8)	0.0022
Pre-op parameters			
Aldosterone (pmol/L)	1 049 (530,1436)	883.5 (569,1254)	0.71
Renin (mU/L)	1.15 (1,2.75)	1.0 (1,2.075)	0.39
ARR	209 (106,287)	168 (109,234)	0.58
K ⁺ (mmol/L)	2.8 (2.5,3.1)	2.8 (2.5,3.1)	0.95
SBP (mmHg)	157 (144,163)	146 (138,152)	0.06
DBP (mmHg)	94 (88,98)	90 (83,96)	0.51
Lateralization Index	6.3 (5,10)	15 (8,26)	0.02
Antihypertensive drugs (N)	2.5 (2,3)	2 (2,3)	0.37
Follow up			
Aldosterone (pmol/L)	445.8 (333,487)	124.9 (80,168)	<0.0001
Renin (mU/L)	1.8 (1.1,3.7)	9.6 (7,13)	<0.0001
ARR	73 (67,103)	16 (9.4,23)	<0.0001
K ⁺ (mmol/L)	3.4 (3.1,3.9)	4.1 (3.7,4.3)	<0.0001
SBP (mmHg)	147 (142,157)	129 (116,134)	0.003
DBP (mmHg)	93.9 (89,96)	83.7 (79,86)	0.008
Antihypertensive drugs	1 (1,3)	0 (0,1)	0.003

Ys: years; F: female, M: male; BMI: body mass index; HT: hypertension; ARR; Aldosterone to renin ratio; SBP: systolic blood pressure, DBP: diastolic blood pressure. Values are expressed as median (interquartile range).

Table 2 Histological characteristics of adrenals from non-cured PA patients

Patient	CYP11B2 ⁺ adenoma size (mm)	APCC (N)	ZG hyperplasia	CYP11B2 expression (% of positive cells)‡	CYP11B1 expression (% of positive cells) ‡
1	25	0	Y	1	2
2	7	0	Y	3	1
3	8	0	Y	3	1
4	11	3	Y	1	1
5	5	0	Y	1	2
6*	No	3	Y	-	-
7**	No	2	N	-	-
8	10	3	N	2	1
9	14	2	N	3	1
10	10	4	N	2	1
11	15	1	Y	3	1
12	11	4	N	3	1

* Micronodular hyperplasia non expressing CYP11B2. **nodule 12mm non expressing CYP11B2. ‡ 1: 1%-33%; 2: 34%-66%; 3: 67%-100%. APCC: aldosterone producing cell clusters; ZG: zona glomerulosa; Y: yes; N: no.

Table 3. Genetic analysis of CYP11B2 positive APAs from PA patients without biochemical success after adrenalectomy

APA	Gene mutated Sanger sequencing	Gene mutated NGS	Read depth	VAF	Nucleotide	Protein
1	<i>KCNJ5</i>	-	-	-	c.451G>C	p.Gly151Arg
2	<i>CACNAID</i>	-	-	-	c.2968C>G	p.Arg990Gly
3	<i>CACNAID</i>	-	-	-	c.1207G>C	p.Gly403Arg
4	-	<i>KCNJ5</i>	1044	28%	c.451G>C	p.Gly151Arg
5	<i>CACNAID</i>	-	-	-	c.4117G>A	p.Val1373Asp
8	-	<i>CACNAID</i>	2384	15%	c.1207G>C	p.Gly403Arg
9	-	<i>KCNJ5</i>	1044	28%	c.451G>C	p.Gly151Arg
10	-	<i>ATP1A1</i>	625	17%	c.311T>G	p.Leu104Arg
11	-	<i>CACNAID</i>	1871	33%	c.3458T>A	p.Val1153Asn
12	-	<i>CACNAID</i>	646	27%	c.2968C>G	p.Arg990Gly

NGS; next generation sequencing. VAF, Variant Allele Frequency. *KCNJ5* (NM_000890), *ATP1A1* (NM_000701), *CACNAID* (NM_001128839.2 and NM_000720).

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