

**CAMPUS INNENSTADT** Medizinische Klinik und Poliklinik IV **Endocrine Research Department** 

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# Immunoexpressions of CYP11B2 and HSD3B2 in genetically characterized aldosterone producing adenomas

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### Introduction and Objectives

Somatic mutations of KCNJ5 (1), CACNA1D (2), ATP1A1 and ATP2B3 (3) have been shown to be involved in the formation of aldosterone producing adenoma (APA). This study aimed to investigate in paraffinembedded APA tissues, the immunoexpression of CYP11B2, the rate-limiting enzyme for aldosterone production and HSD3B2, the prevalent isoform of 3β-HSD found in APA, and to relate these findings to mutation status, histopathology of APA and biochemical and clinical outcome of adrenalectomy.

#### Methods

#### Subjects and human adrenal tissue

Patients (n= 59) were diagnosed with PA by AVS and CT-scan according to the Endocrine Society Practice Guidelines (4) and were included in the German Conn's Registry. After unilateral adrenalectomy the adrenal tissue was histologically confirmed as APA. All patients gave written informed consent and the study was approved by the ethics committee of the University of Munich. Biochemical and clinical data was prospectively collected from PA diagnosis date up until 16.8 ±11.3\* months following adrenalectomy.

#### Histological and immunohistochemical staining analysis

32 out of 59 genotyped APA presented available paraffin-embedded tissues and were selected for functional immunohistochemical staining analysis. Hematoxylin and eosin (H&E) staining was performed with routine protocol to determine cell composition. Immunostaining was performed for CYP11B2 using the ImmPRESS Reagent (Vector laboratories) and for HSD3B2 using the Vectastain ABC Elite Kit (Vector laboratories). Primary antibodies used in immuno-histochemistry are listed in table 1. The antigenantibody complex was visualized with 3,3'-diaminobenzidine solution (DAB) and counterstained with hematoxylin. Double immunostaining was performed using Polink DS-MR-Hu C1 kit (GBI Labs). Semi-quantitative immunohistochemical evaluation was assessed using McCarty's H-scoring system and statistical analysis was performed using Mann-Whitney multiple comparison test.

#### APA Genotyping

Genotyping was performed by direct bidirectional Sanger sequencing or whole exome sequencing. Tissues were divided into five groups according to their mutation status: KCNJ5, CACNA1D, ATP1A1, ATP2B3 or Wild type (WT - defined as the absence of candidate gene mutation).

Table 1 Primary antibodies used in immunohistochemical studies of APAs.

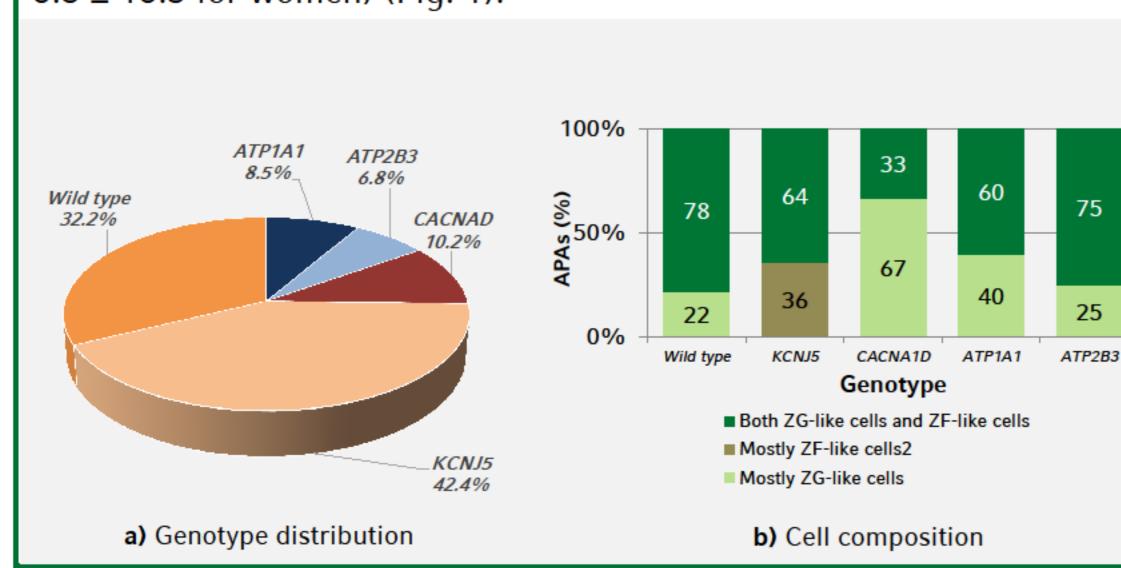
Antibody	Source
Anti-hCYP11B2	Celso Gomez-Sanchez et al., 2014
clone 41-17 (Mouse monoclonal)	(5)
Anti-hHSD3B2	Abcam
ref. Ab154385 (Rabbit polyclonal)	

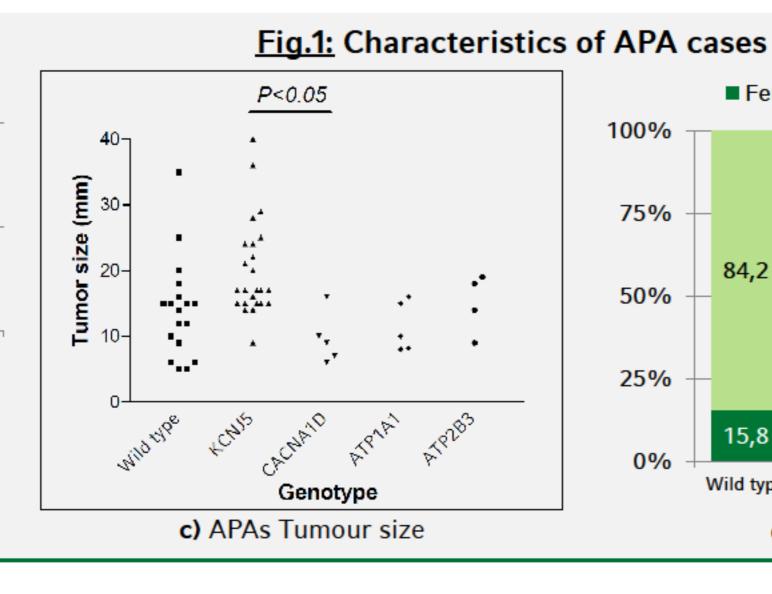
\* (values are expressed as mean ± SD unless specified otherwise)

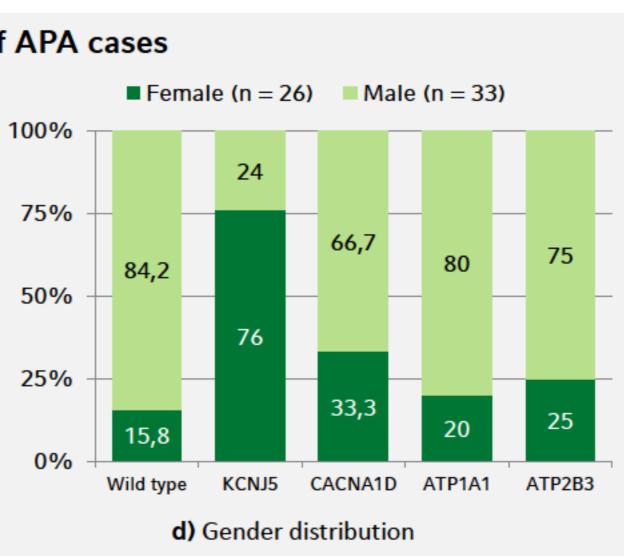
### Results

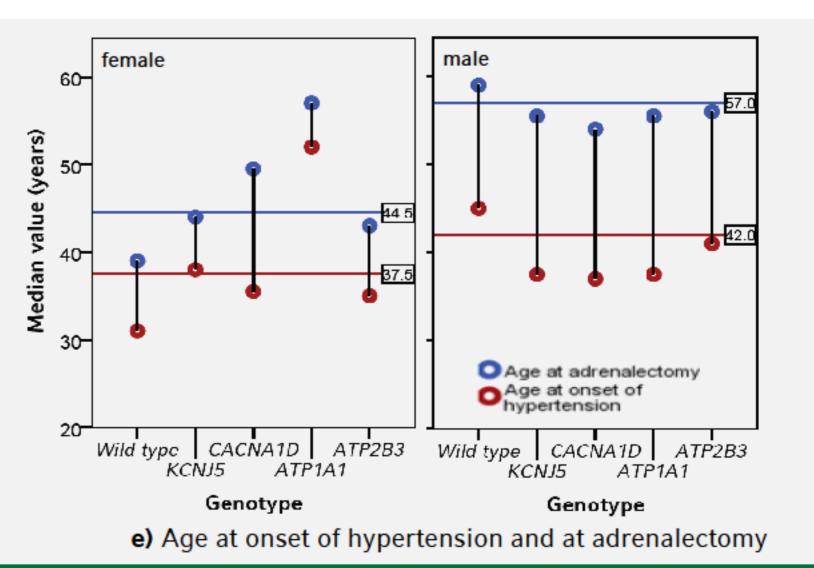
#### Genetic and histopathological characterization of APA cases

68% of APAs presented with a somatic mutation, most frequently KCNJ5 (42.4%). Histomorphological analysis showed that a majority of APAs consisted of both zona fasciculata-like (ZF -like) clear cortical cells and zona glomerulosa-like (ZG-like) compact cells, except for CACNA1D-mutated APAs, where 67% presented only ZG-like cells. CACNA1D-mutated tumours presented significantly smaller tumour size (P<0.05) than KCNJ5-mutated APAs. The gender distribution shows that mostly men are affected by APAs, except for KCNJ5-mutated APA, which were predominantly seen in women (76%). Earlier onset of hypertension is observed in women (median age: 37.5 ± 8.3 years vs 42 ± 9.4 for men), while longer duration of hypertension before adrenalectomy is observed in men (median duration: 13 ± 16.3 years vs  $6.5 \pm 16.3$  for women) (Fig. 1).



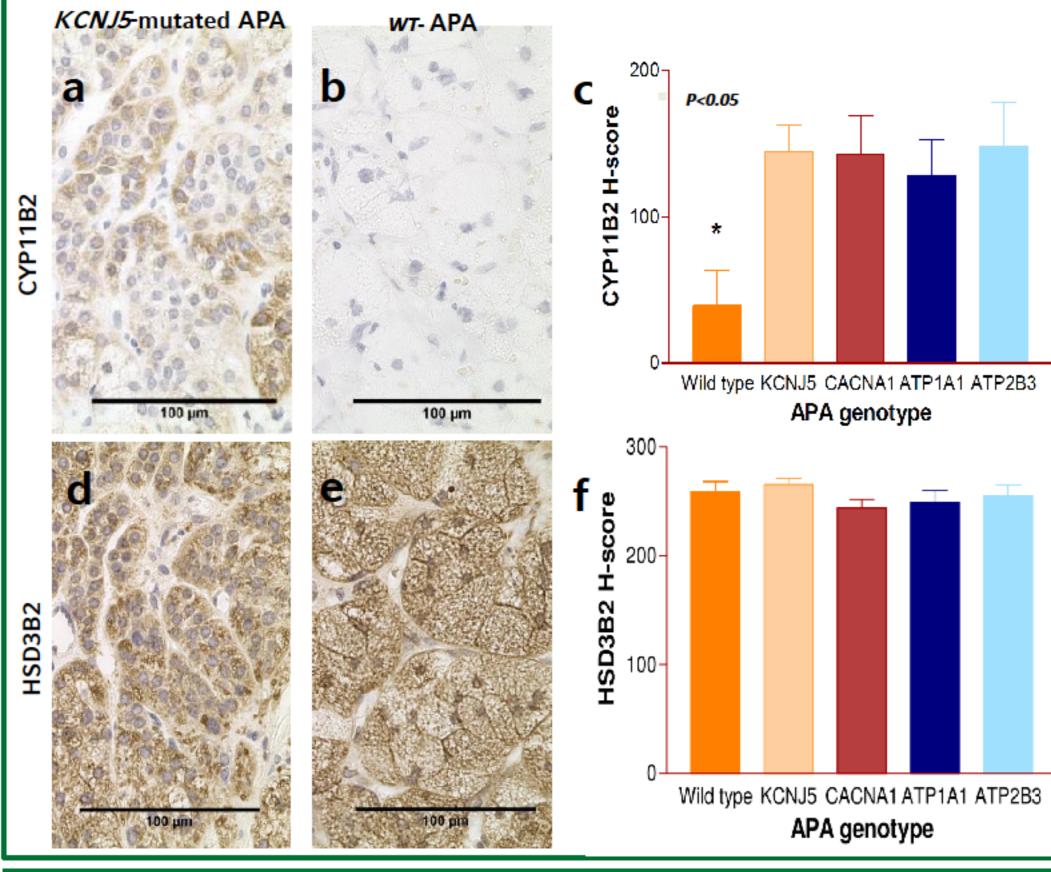


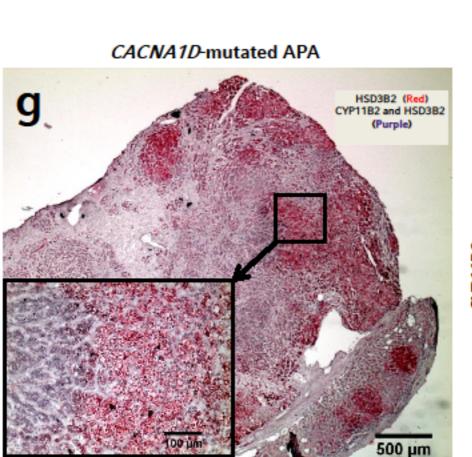


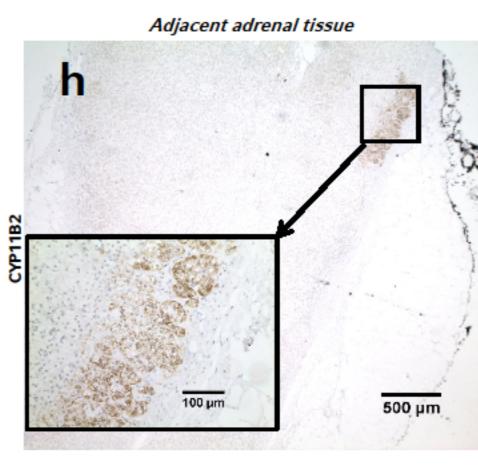


#### Immunoexpression of CYP11B2 and HSD3B2 in genetically characterized APAs

APAs with a somatic mutation presented CYP11B2 positive clusters or scattered cells (Fig. 2a). WT-APAs were weakly or negatively stained for CYP11B2 (Fig. 2b) while the adjacent ZG presented CYP11B2-positive clusters (Fig. 2h). The mean Hscores for CYP11B2 (Fig. 2c) was significantly different between WT-APAs and mutated APAs (P<0.05). APA lesions presented wider distribution of HSD3B2 independently of their mutation status (Fig. 2 d,e) and no significant difference was noted in HSD3B2 immunoexpression regarding the APAs genotype (Fig. 2f). Expression patterns of CYP11B2 and HSD3B2 in APA were visualized using double immunostaining (Fig. 2g).







#### Fig. 2: Immunohistochemical analysis of genetically characterized APAs.

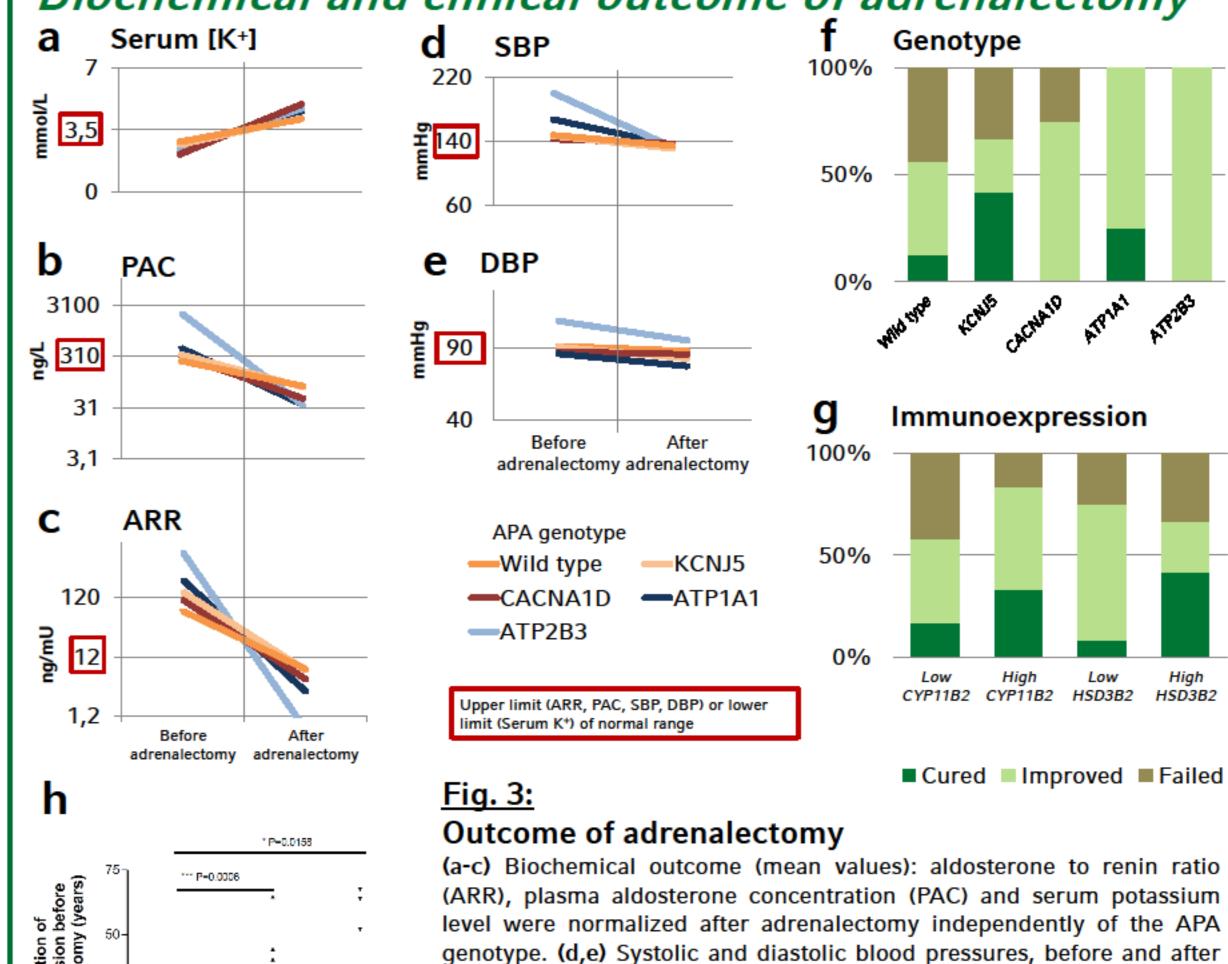
(a) CYP11B2 expression in a KCNJ5-mutated APA. (b) CYP11B2 expression in a WT-APA. (c) Mean H-scores of CYP11B2 immunoexpression. (d) HSD3B2 expression in a KCNJ5-mutated APA. (e) HSD3B2 expression in a WT-APA. (f) Mean H-scores of HSD3B2 immunoexpression. (g) Double immunostaining of CYP11B2 and HSD3B2 in a CACNA1D-mutated APA: red staining indicates immunoexpression of HSD3B2, purple cytoplasmic staining illustrates coimmunoexpression of CYP11B2 and HSD3B2.(h) CYP11B2 immunoexpression in adjacent adrenal tissue of a WT-APA.

### Conclusions

Our findings suggest that mutated APAs present a significantly higher CYP11B2 immunoreactivity, compared to wild type-APAs. In our analyzed APA cases, neither immunoreacivities of HSD3B2 or CYP11B2, nor genotype seem to be correlated with postadrenalectomy outcome in primary aldosteronism.

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# Biochemical and clinical outcome of adrenalectomy



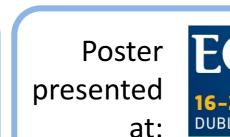
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Outcome of adrenalectom



andrenalectomy. (f) Comparison of adrenalectomy outcome depending

on APA genotype. (g) Comparison of adrenalectomy outcome in APA

cases with low (≤ median H-score) vs high (> median H-score)

immunoexpression of CYP11B2 and HSD3B2. (h) Clinical outcome was

significantly better in patients presenting a shorter duration of