

# **ADRENOCORTICAL HORMONE PROFILES DO NOT PREDICT THE MOLECULAR** ETIOLOGY IN NON-CAH PRIMARY ADRENAL INSUFFICIENCY

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## **PATIENTS AND DESIGN**

- Forty-one children (19 females, median age: 3 months, range: 0-8 years) with non-CAH PAI of unknown etiology from 16 tertiary pediatric endocrinology clinics.
- Patients with CAH, adrenoleukodystrophy, autoimmune adrenal insufficiency or obvious syndromic PAI on clinical and biochemical assessment were excluded.
- Genetic analysis was performed using either targeted gene panel or whole-exome sequencing.
- Plasma adrenal steroids were quantified by liquid chromatography-mass spectrometry and compared to that of controls.

# RESULTS

- Molecular diagnosis was found in 11 genes in 29 (70.7%) cases. The range of genetic etiologies found in this cohort were: StAR (n=6; 20%), MC2R (n=6; 20%), NNT (n=3; 10%), NROB1/DAX1 (n=3; 10%), CYP11A1 (n=2; 7%), MRAP (n=2; 7%), SGPL1 (n=2; 7%), ABCD1 (n=1; 3%), AIRE (n=1; 3%), AAAS (n=1; 3%), and HSD3B2 (n=2; 7%).
- In 12 patients whose genetic etiology could not be found by TPS, further diagnosis was not be achieved by WES.

## **Table 1.** Deficiency of adrenocortical steroids according to underlying molecular defect in patients with rare causes of PAI.

Gene (Chromosome locus)	Variant	n	Age of diagnosis	GC deficie
MC2R (18p11.21)	c.455C>A (p.T152K) c.560del (p. V187fs) c. 476C>A (p.T159K)	2 2 2	6.25 years – 4 days 4 days – 3 months 2.48 years – 14 days	6/6
STAR (8p11.23)	c.505G>A (p.E169K) c.470T>C (p.L157P)	3 3	3 days – 5 days – 5 months 1 month – 2 months – 2 months	6/6
NNT (5p12)	exon 12-14 deletion c.259C>T (p.Q87*) c.2507G>A (p.G836D)	1 1 1	1.5 years 6 months 1.48 years	3/3
NR0B1 (Xp21.2)	c.1075C>T (p.Q359*) c.1210C>T (p. Q404*) exon 1-2 deletion	1 1 1	1.5 years 2.5 years 1.5 years	3/3
CYP11A1 (15q24.1)	<b>c.461T&gt;C (p.L154P)</b> c.1351C>T (p.R451W)	1 1	2 days 1.66 years	2/2
MRAP (21q22.11)	c.106 +1del <b>c.106 + 2dupT</b>	1 1	2 days 1.4 years	2/2
SGPL1 (10q22.1)	c.1018C>T (p.R340W) <b>c.518T&gt;A (p.L173Q)</b>	1 1	5 months 4 months	2/2
ABCD1 (Xq28)	c.1772G>T (p.R591L)	1	8 years	1
AIRE (21q22.3)	c.415C>T (p.R139*) / c.260T>C (p.L87P)	1	4.5 years	1
AAAS (12q13.13)	c.1333C>T (p.R478*)	1	4.5 years	1
HSD3B2 (1p12)	c.1003C>T (p.R335*) <b>c.939del (p.F314Sfs*54)</b> / c.745C>T (p.R249*)	1 1	1 month 1 month	2/2

Sex steroid deficiency IUGR / Short MC **Elevated** deficiency TSH stature Gonada Adrena 0/2 0/1 2/2 1/2 2/2 0/0 0/2 1/2 0/6 0\*/2 2/2 2/2 0/0 3/3 3/3 1/3 1/0 6/6 3/3 0/0 3/3 0/3 0/1 0/0 0/1 1/1 1/0 0/1 1/1 0/1 0/0 1/1 0/1 0/0 1/1 1/1 1/0 3/3 1/1 0/1 1/1 1/1 0/1 0/0 1/1 0/1 0/0 1/1 2/2 1/0 0/1 1/1 0/1 0/1 0/1 0/0 1/1 0/2 0/0 1/1 1/1 \*/1 0/1\*\* 1/1 1/1 1/1 2/2 0/1 0/1\*\* 1/1 1/1 0/0 0/0 0/1 1/1 0/1 1/1 1/1 1/1 0/1 \*/1 1/1 0/0 0/0 0/1 0/1 0/1 2/2 0/1 1/1 1/0 0/1

	Patients (n=29)		Control (n=324)		
(ng/mL)	median	range	median	range	р
Idosterone	0.06	0.007-0.70	0.198	0.002-0.982	0.0002
ortisol	0.891	0.021-77.5	31.08	1.681-248.8	0.0003
HEA-S	4.22	0.308-2757	132.1	5.609-978.1	0.5301
orticosterone	0.057	0.002-5.12	0.926	0.025-19.62	0.0059
1-Deoxycortisol	0.029	0.001-2.03	0.231	0.028-2.094	0.0021
ndrostenedione	0.013	0.001-0.42	0.073	0.001-0.824	0.2815
1- Deoxycorticosterone	0.017	0.001-0.55	0.035	0.002-3.085	0.4476
HEA	0.112	0.01-35.9	0.626	0.012-28.07	0.5164
<b>70H-Pregnenolone</b>	0.105	0.034-46.55	0.506	0.033-9.487	0.0043
70H-Progesterone	0.013	0.002-1.934	0.214	0.001-2.728	0.0852
rogesterone	0.021	0.003-0.174	0.027	0.001-1.684	0.2966
ndrosterone	0.44	0.004-10.01	0.97	0.004-24.56	0.9453
regnenolone	0.126	0.017-51.97	0.421	0.042-10.19	0.0024
ortisone	0.426	0.026-23.09	29.3	1.376-165.9	<0.0001
<b>1-Deoxycortisol</b>	0.031	0.003-0.211	0.086	0.001-0.944	0.022

Table 3. Com patients with s

Age of diagnosis **Consanguinity** of MC deficiency ( Sex steroid defi

Karyotype (n)

Legal sex (n)

ACTH (N=10-60 Cortisol (µg/dL) Cortisone (ng/m **Corticosterone** 11-Deoxycortico

# RESULTS

## **RESULTS**

RESULIS								
parison of clinical and biochemical characteristics of PAI solved or unsolved etiology.								
	PAI solved (n=29)	PAI unsolved (n=12)	р					
s (yrs)	1.3 ± 0.3	5.7 ± 2.1	0.005**					
of parents (n)	23	3	0.001**					
n)	17	7	0.98					
iciency (n)								
Gonadal	9	1	0.12					
Adrenal	28	11	0.5					
46,XX	10	2	0.25					
46,XY	19	10	0.25					
Female	15	2	0.03*					
Male	14	10						
pg/mL)	1132 ± 55.14	992.7 ± 204.8	0.38					
	0.8 ± 0.3	0.4 ± 0.1	0.002**					
nL)	3.8 ± 1.6	24.2 ± 5.1	0.0001***					
(ng/mL)	0.20 ± 0.06	1.38 ± 0.40	<0.0001****					
osterone (ng/mL)	0.02 ± 0.007	0.10 ± 0.05	0.02*					

- molecular etiology is low.

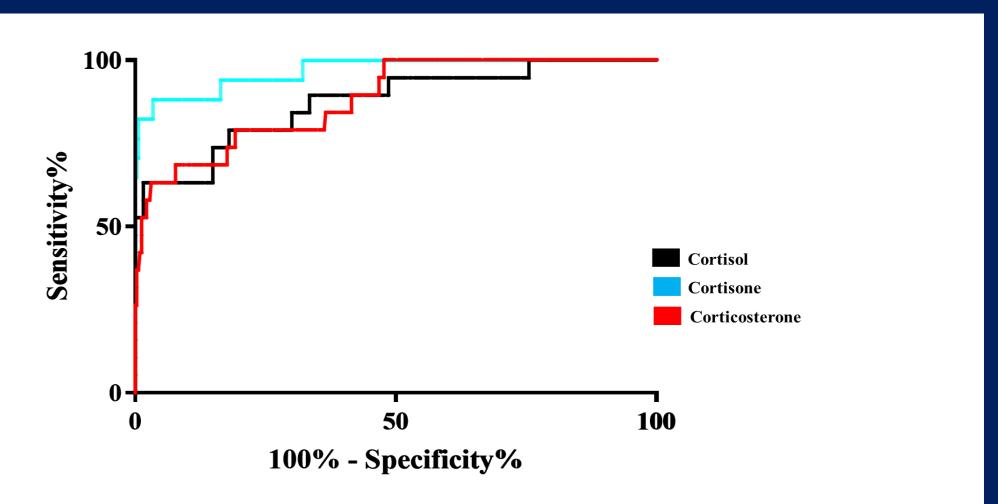
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itients with solved molecular etiologies had lower ncentrations of steroids than healthy infants, who has sysiologically low steroids. The difference was most for aldosterone, cortisol, cortisone, nificant rticosterone, pregnenolone, 170H-pregnenolone, 11eoxycortisol, 21-deoxycortisol.

asma cortisol<4 ng/mL, cortisone<11 ng/mL, and rticosterone<0.11 ng/mL had >95% specificity to gregate non-CAH PAI patients compared to control oups (p < 0.0001, area under the ROC curve: 0.96, 0.88, 87, respectively)



**ure 1.** Specificity and sensitivity of cortisol, cortisone and rticosterone in the diagnosis of PAI

## CONLUSION

Adrenocortical hormone profiles determined by LC-MS/MS which demonstrate significantly low glucocorticoids, mineralocorticoids and adrenal androgens, are highly sensitive for the recognition of non-CAH PAI even at early infancy, but the specificity to indicate a definitive

Targeted gene panel sequencing is a first-line approach in the molecular diagnosis of non-CAH PAI with high efficacy.

Nevertheless, lower ACTH, absent parental consanguinity, late-onset of diagnosis and a less severe deficiency profile of steroid hormones decreases the probability of achieving the genetic diagnosis.

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