

Clinical and Genetic Heterogeneity, Overlap with Other Tumor Syndromes, and Atypical Glucocorticoid Hormone Secretion in Adrenocorticotropin-Independent Macronodular Adrenal Hyperplasia Compared with Other Adrenocortical Tumors

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Objective: ACTH-independent macronodular adrenal hyperplasia (AIMAH) is often associated with subclinical cortisol secretion or atypical Cushing's syndrome (CS). We characterized a large series of patients of AIMAH and compared them with patients with other adrenocortical tumors.

Design and Patients: We recruited 82 subjects with: 1) AIMAH (n = 16); 2) adrenocortical cortisol-producing adenoma with CS (n = 15); 3) aldosterone-producing adenoma (n = 19); and 4) single adenomas with clinically nonsignificant cortisol secretion (n = 32).

Methods: Urinary free cortisol (UFC) and 17-hydroxycorticosteroid (17OHS) were collected at baseline and during dexamethasone testing; aberrant receptor responses was also sought by clinical testing and confirmed molecularly. Peripheral and/or tumor DNA was sequenced for candidate genes.

Results: AIMAH patients had the highest 17OHS excretion, even when UFCs were within or close to the normal range. Aberrant receptor expression was highly prevalent. Histology showed at least two subtypes of AIMAH. For three patients with AIMAH, there was family history of CS; germline mutations were identified in three other patients in the genes for *menin* (one), *fumarate hydratase* (one), and *adenomatosis polyposis coli (APC)* (one); a *PDE11A* gene variant was found in another. One patient had a *GNAS* mutation in adrenal nodules only. There were no mutations in any of the tested genes in the patients of the other groups.

Conclusions: AIMAH is a clinically and genetically heterogeneous disorder that can be associated with various genetic defects and aberrant hormone receptors. It is frequently associated with atypical CS and increased 17OHS; UFCs and other measures of adrenocortical activity can be misleadingly normal. (*J Clin Endocrinol Metab* 94: 2930–2937, 2009)

Endogenous Cushing's syndrome (CS) is due to primary adrenal disease in approximately 15–20% of cases; in these patients, CS is caused mainly by unilateral adenomas (1). Bilateral adrenal lesions occur only in 10–15% of adrenal CS and include primary pigmented nodular adrenocortical disease (PPNAD) and ACTH-independent macronodular adrenal hyperplasia (AIMAH), also known as massive macronodular adrenal disease (2, 3). PPNAD is most frequently associated with Carney complex and is caused mainly by inactivating germline mutations of *PRKARIA* and allelic losses of its locus at the 17q22-24 chromosomal region (4, 5).

AIMAH is a rare cause of CS, accounting for less than 1% of adrenal CS. Since the review of AIMAH by Lieberman *et al.* in 1994 (6), a greater number of cases have been reported (7). Several studies demonstrated that the regulation of cortisol secretion in AIMAH (as well as in some unilateral, single adenomas) is mediated by the aberrant expression and function of membrane-bound G protein-coupled receptors such as those for gastric inhibitory polypeptide (GIP), vasopressin, catecholamines, LH/human chorionic gonadotropin, serotonin, IL-1, and leptin (8–14).

Establishing the diagnosis of AIMAH with CS is based on, first, the clinical phenotype of CS; this is followed by the demonstration of ACTH-independent hypercortisolism and bilateral adrenal nodular enlargement on radiologic imaging. More often than not, diagnosis of AIMAH is difficult because hypercortisolism usually develops slowly over several years, may be cyclical and is frequently associated with subtle clinical manifestations. Radiologic imaging is helpful, but occasionally adrenal nodularity is indistinguishable from that in normal elderly persons (15). Biochemically, too, there is no specific testing for the diagnosis of AIMAH: whereas most patients with PPNAD respond to dexamethasone with a paradoxical (unexpected) increase in glucocorticoid excretion during Liddle's test, AIMAH patients respond to dexamethasone as patients with the common adrenal cortisol-producing tumors do, requiring final confirmation of the diagnosis by histological examination (16).

In this investigation, we studied genetically and clinically a large series of patients with AIMAH and compared them with patients with other adrenocortical tumors. The data prove the clinical and genetic heterogeneity of the condition in these patients but also demonstrate the large genetic component of the condition. Finally, our study proposes that urinary 17-hydroxycorticosteroid (17OHS) is the best screening test for hypercortisolism in AIMAH.

Patients and Methods

A total of 102 patients were admitted to the National Institutes of Health Warren Magnuson Clinical Center from 2000 to 2008 for the work-up and treatment of adrenocortical tumors under protocol 00CH160. Eighty-two of 102 patients (28 males and 54 females) met the following criteria: 1) evidence for the existence of an adrenocortical tumor, as indicated by imaging studies or biochemical investigation of hormonal secretion; 2) exclusion of Carney complex or other diseases associated with micronodular forms of adrenocortical hyperplasia. The National Institute of Child Health and Human Development Institutional Review Board approved this study; informed consents were obtained from all subjects.

The 82 subjects whose data were included here were divided into four groups on the basis of their final histological examination or, for those with nonsecreting tumors, biochemical testing and radiological imaging: 1)

AIMAH (n = 16); 2) adrenocortical cortisol-producing adenoma with CS (ACS) (n = 15); 3) aldosterone-producing adenoma (APA) (n = 19); and 4) single adenomas with clinically nonsignificant cortisol secretion (SCA) (n = 32).

The following data were analyzed for all subjects, as we have reported elsewhere for the investigation of CS (17): diurnal variation in plasma ACTH and cortisol levels; cortisol levels before and after the overnight 8-mg dexamethasone test; and plasma ACTH and cortisol levels before and after iv administration of 1 μ g/kg ovine CRH (oCRH; corticorelin). Urine was collected for 24 h for free cortisol excretion (UFC) and 17-hydroxycorticosteroid (17OHS), and their data are presented as the mean of two consecutive measurements.

Finally, in all patients with AIMAH, a review of their pathological reports (*i.e.* macroscopic appearance and weights) and histology was conducted. We recorded size and number of nodules and the presence of hyperplasia or atrophy of nonnodular adrenal cortex, as we have reported elsewhere (18).

Provocative tests for the detection of aberrant receptor expression

Provocative tests for the identification of aberrant receptor expression was employed in the AIMAH and ACS subgroups over a 3-d period, as described previously (19, 20). The protocol is based on monitoring plasma levels of steroids at 30- to 60-min intervals for 2–3 h during tests that transiently modulate the levels of ligands for potentially aberrant receptors. Initial tests included a posture test performed in a supine position for baseline, followed by a 2-h ambulatory period (to evaluate potential modulation by vasopressin, catecholamines, angiotensin II, and others); this was followed by a standard mixed meal to evaluate the response to fluctuations of gastrointestinal hormones including GIP. On the second day, the administration of 100 μ g GnRH iv (gonadorelin, 1 μ g/kg, iv, testing for modulation by FSH, LH, GnRH) was followed by 200 μ g TRH iv (modulation by TSH, prolactin, TRH) and GHRH (sermorelin, 1 μ g/kg, iv). Responses to 1 mg glucagon iv, 10 IU arginine vasopressin im were tested sequentially on the third day. A change of 50% or greater of plasma cortisol was defined as a positive response; a 25–49% change, as a partial response; and a change of less than 25%, as no response; these cutoffs and their justification have been published elsewhere (19, 20).

Dexamethasone testing

A 6-d Liddle's test was conducted for patients with AIMAH, as described elsewhere (16, 17, 21). After 2 d of baseline measurement of urinary steroid excretion, dexamethasone, 0.5 mg, was given orally every 6 h for 2 d starting at 0600 h; the dose of dexamethasone was then increased to 2 mg every 6 h for the last 2 d of the test. The 24-h UFC and urinary 17OHS were measured on each day and percentage changes from baseline were calculated. 24-h UFC were corrected for body surface area and 17OHS rates were corrected for creatinine excretion (per day per gram creatinine). In all patients, dexamethasone levels were measured on the last day of the test, and this ensured adequate absorption of the medication; in addition, ACTH levels were suppressed (<5 pmol/liter) in all patients on the same day of the test.

Hormone assays

Plasma cortisol and ACTH levels were measured as described elsewhere (22). UFC excretion was measured by direct RIA (23). The intraassay coefficient of variation was 5%, and the interassay coefficient of variation was 10%; urinary 17OHS excretion was measured by using a modification of the colorimetric method as previously described by Murphy (24). The intraassay and interassay coefficients of variation were 6 and 11%, respectively.

Genetic and other molecular tests

Genomic DNAs were obtained from the blood leukocytes in all subjects and the adrenocortical tissues when they underwent surgery (25, 26). Mutation analysis was performed for the *menin* (*MEN1*), fumarate hydratase (*FH*), adenomatous polyposis coli (*APC*), *GNAS*, and *PDE11A* (phosphodiesterase 11A) genes (26–28).

TABLE 1. Clinical and laboratory data for all patients with ACTH-independent macronodular adrenal hyperplasia and other adrenocortical tumors

	AIMAH (n = 16)	ACS (n = 15)	APA (n = 19)	SCA (n = 32)	P value
Gender (F:M)	11:5	11:4	7:12	25:7	
Age (yr)	46.8 ± 9.8	44.3 ± 14.8	51.2 ± 10.0	49.0 ± 10.7	0.32
8 AM cortisol (nmol/liter)	535.2 ± 198.6 ^a	538.0 ± 215.2 ^a	433.2 ± 82.8	389.0 ± 140.7	0.03
12 midnight cortisol (nmol/liter)	435.9 ± 242.8 ^{b,c}	471.8 ± 355.9 ^{b,c}	168.3 ± 57.9	129.7 ± 69.0	<0.01
ACTH (pmol/liter)	1.9 ± 2.5 ^b	0.7 ± 0.4 ^d	4.8 ± 3.7	3.5 ± 2.3	<0.01
Cortisol, before DEX (nmol/liter)	527.0 ± 157.3	405.6 ± 126.9	485.6 ± 209.7	449.7 ± 182.1	0.35
Cortisol, after DEX (nmol/liter)	320.0 ± 298.0 ^{a,b}	303.5 ± 223.5 ^a	93.8 ± 88.3	85.5 ± 60.7	<0.01
UFC (nmol/d · g creatinine)	611.9 ± 320.7 ^{c,d}	625.7 ± 448.5 ^{c,d}	183.5 ± 123.6	135.2 ± 64.0	<0.01
17OHS (mmol/d · g creatinine)	43.1 ± 20.4 ^{c,d}	29.8 ± 13.8 ^b	15.2 ± 10.8	18.2 ± 12.7	<0.01
Change at d 6 in Liddle's test (%)					
UFC	18.0 ± 48.0 ^{b,c}	44.6 ± 109.7 ^{b,c}	−98.7 ± 0.1	−90.3 ± 15.6	<0.01
17OHS	14.2 ± 47.1 ^{b,c}	10.6 ± 44.9 ^a	−61.2 ± 7.5	−57.5 ± 18.8	<0.01

Conversion factors to metric units are as follows: cortisol, 0.036 to micrograms per deciliter; ACTH, 5.0 to picograms per milliliter; UFC, 0.362 to micrograms per day per gram creatinine.

^a $P < 0.05$ compared with SCA.

^b $P < 0.05$ compared with APA.

^c $P < 0.01$ compared with SCA.

^d $P < 0.01$ compared with APA.

Aberrant receptor expression that was suggested by clinical testing was confirmed by molecular testing as previously published (19, 20). The snap-frozen tissue (50–100 mg) was homogenized with 1 ml TRIZOL reagent (Invitrogen, Carlsbad, CA). The extracted RNA was treated with deoxyribonuclease digestion to eliminate contaminating genomic DNA using ribonuclease-free deoxyribonuclease set (QIAGEN, catalog no. 79254; Valencia, CA). The RNA was further cleaned up with RNeasy minikit (QIAGEN, catalog no. 74104). The integrity and concentration of the RNA were determined by agarose gel electrophoresis with ethidium bromide and ND-1000 spectrophotometer (NanoDrop, Wilmington, DE). Two micrograms of RNA was reverse transcribed to cDNA using the oligo(dT) primers in the SuperScript III first-strand synthesis system (Invitrogen), according to the procedures by the manufacturer. PCRs were performed with Biolase DNA polymerase (Bioline, Randolph, MA) and primers specific to the genes of individual receptors and internal control, β -actin. The primers were designed for the receptors of ACTH, V1, V2, V3, GIP, and LH/human chorionic gonadotropin, either flanking the intron or on the exon/exon boundary to prevent the amplification from the genomic DNA (29–31). The amplicon products of the PCR were visualized and autographed in agarose gel electrophoresis with ethidium bromide under the Alphamiger 3400 (Imgen Technologies, Alexandria, VA).

Statistical analysis

All data were expressed as the mean ± SD (for demographic data) or mean ± SE (for all comparisons and the data presented in the figures). Data were analyzed by using SPSS 14.0 for Windows (SPSS Inc., Chicago, IL). The one-way ANOVA with *post hoc* follow-up (Tukey honestly significant difference) for multiple comparisons was used to test for any differences in demographics; for the percentage and frequency among groups in provocative tests, the χ^2 test and follow-up pairwise comparison tests were done with cross-tabulation procedure. The Kruskal-Wallis H test was used to assess differences in Liddle test; Mann-Whitney U test was used for *post hoc* test among groups. For all statistical comparisons, $P < 0.05$ was considered significant.

Results

Demographics

The mean age of AIMAH patients was 46.8 yr (46.8 ± 9.8 yr); this was not significantly different from the mean age of patients in

other groups (Table 1). The overall gender distribution (female to male) for this study was 54 to 28, a female predominance (65.9% females) that is similar to what has been reported for adrenal tumors in other studies. The only group of patients in which males ($n = 12$) outnumbered females ($n = 7$) was the APA group.

Clinical characteristics and baseline glucocorticoid hormone levels

Patients with AIMAH and ACS did not have any diurnal rhythmicity in their cortisol secretion; those with APA and SCA groups had the expected diurnal variation in their midnight to morning cortisol values. Morning cortisol values were higher for patients with AIMAH and ACS, 535.2 ± 198.6 and 538.0 ± 215.2 nmol/liter, respectively, *vs.* those in patients with SCA (389.0 ± 140.7 nmol/liter, $P < 0.05$) (Table 1). Patients with AIMAH and ACS had a trend of higher morning cortisol levels than the APA group, but this difference did not reach statistical significance.

UFCs were significantly different in patients with AIMAH and ACS *vs.* those with APA and SCA (611.9 ± 320.7 and 625.7 ± 448.5 nmol/d 183.5 ± 123.6 and 135.2 ± 64.0 nmol/d, respectively, $P < 0.01$). Patients in the AIMAH group had the highest urinary 17OHS excretion (43.1 ± 20.4 mmol/d · g crn); 17OHS were 29.8 ± 13.8 mmol/d · g crn in patients with ACS ($P = 0.08$) and 15.2 ± 10.8 and 18.2 ± 12.7 mmol/d · g crn ($P < 0.01$) in patients with APA and SCA, respectively (Table 1).

Dexamethasone and oCRH testing, and aberrant receptor expression studies

During the Liddle's test, patients with AIMAH and ACS failed to suppress their glucocorticoid hormone serum levels and urinary excretion rates, consistent with our data in an earlier cohort of similar patients that were studied in our institution (16). In contrast, all patients with APA and SCA suppressed (Fig. 1, A and B). The percentage changes on d 6 from baseline (d 0)

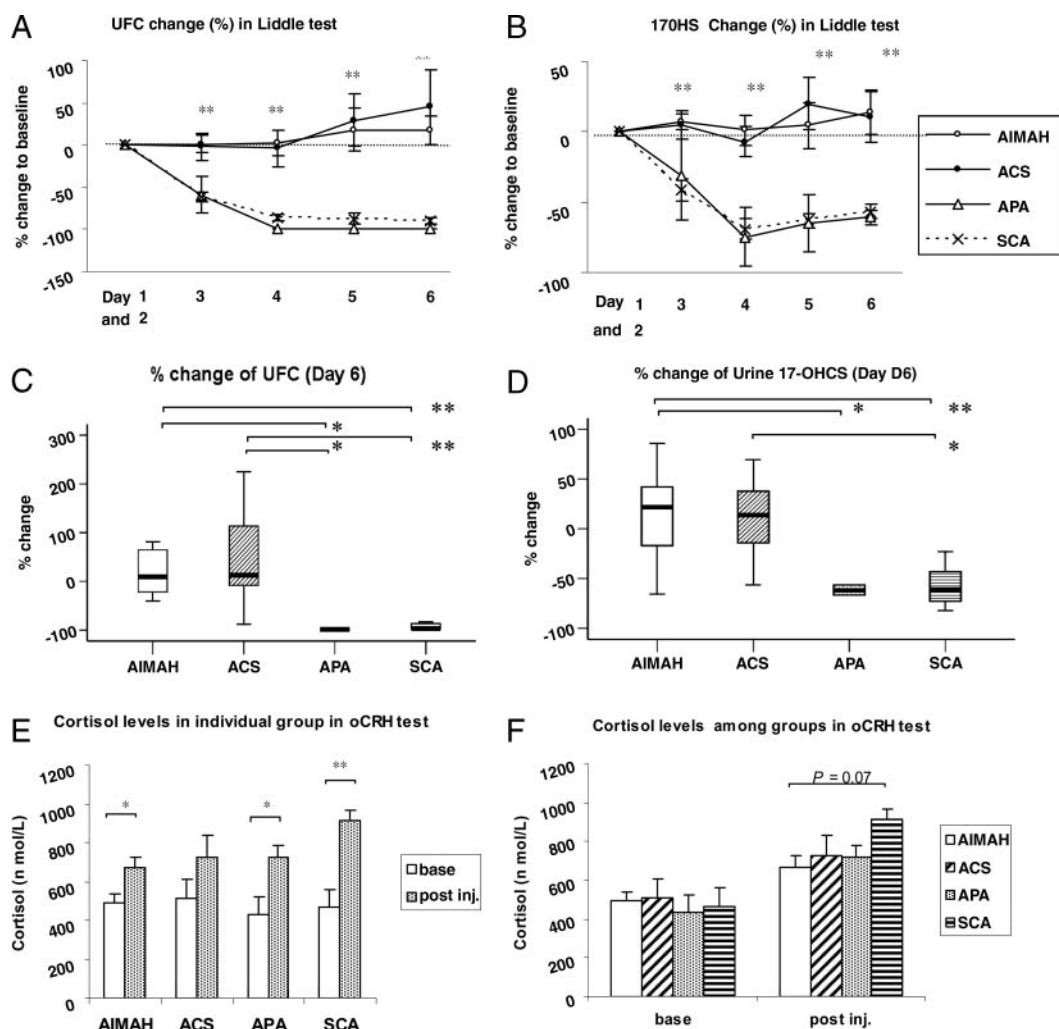


FIG. 1. A and B, Serial percentage changes (from d 1 to d 6) compared with the baseline of UFC and 17OHS during Liddle’s test. C and D, Percentage changes compared with baseline in UFCs and 17OHS on d 6 of the Liddle’s test. E and F, Changes in cortisol levels during the oCRH test. *, $P < 0.05$; **, $P < 0.01$.

were for patients with AIMAH, ACS, APA, and SCA; UFCs $18.0 \pm 48.0\%$, $44.6 \pm 109.7\%$, $-98.7 \pm 0.1\%$, and $-90.3 \pm 15.6\%$ ($P < 0.01$), respectively; and 17OHS, $14.2 \pm 47.1\%$, $10.6 \pm 44.9\%$, $-61.2 \pm 7.5\%$, and $-57.5 \pm 48.8\%$ ($P < 0.01$), respectively (Fig. 1, C and D).

Although cortisol levels responded to oCRH testing in all patients, patients with AIMAH, ACS, and APA had lower responses than what was seen in patients with SCA (670.4 ± 55.2 , 725.6 ± 110.4 , and 722.8 ± 60.7 nmol/liter *vs.* 915.9 ± 52.4 nmol/liter, respectively $P = 0.07$) (Fig. 1, E and F). It is note-

worthy that in almost none of our patients with CS there was a flat response to oCRH stimulation.

The response rates for the aberrant receptor provocative tests for patients with AIMAH and ACS are shown on Table 2; there were no statistically significant differences in response rates between these two groups of patients with cortisol-producing tumors. In all cases, expression of the aberrant receptor was confirmed by RT-PCR (Fig. 2) and as previously published (19, 20). There was no significant difference in the expression of the aberrant receptors within the AIMAH group.

Clinical and molecular genetics

There were three unrelated patients among the 16 with AIMAH who had family history of adrenocortical tumors and/or ACTH-independent CS; the relatives of these patients, however, were not available for investigation. Inheritance of the condition was suggested to be in an autosomal dominant manner based on the constructed pedigrees (data not shown). There were no familial cases in any of the other groups of patients (Table 3).

We identified three germline mutations in known genes in three other patients with AIMAH. Interestingly, none of these

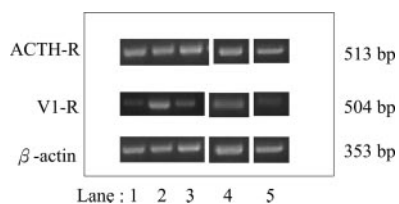


FIG. 2. The RT-PCR analysis of ACTH receptor and V1 receptor expression in various adrenal tissue samples compared with the β-actin. Lanes 1 and 2, Type 1 (BMAH) AIMAH patients; lane 3, type 2 (diffuse hyperplasia) AIMAH patient; lane 4, ACS patient; and lane 5, normal control adrenal tissue.

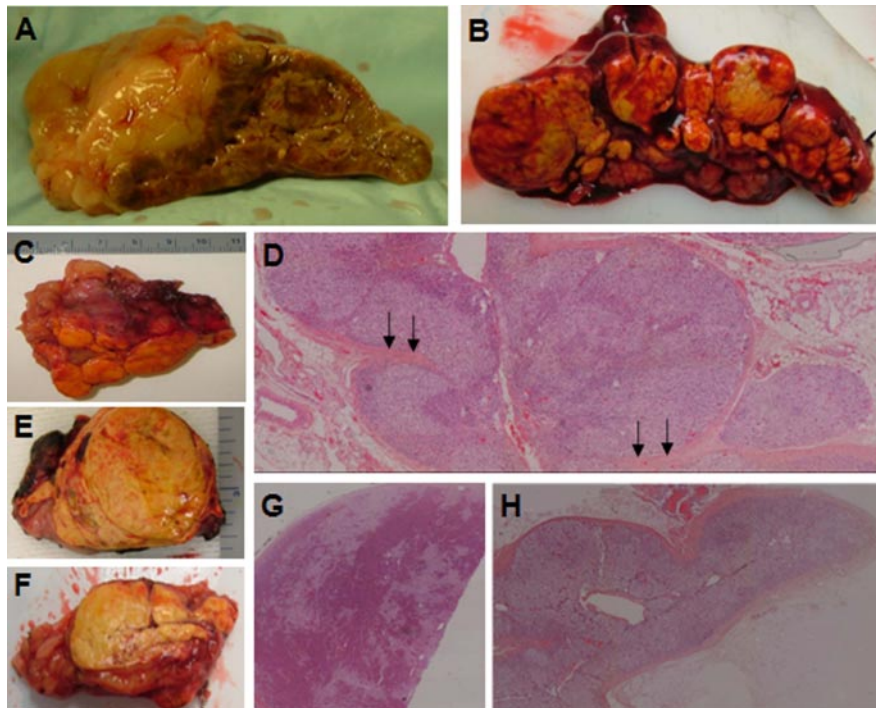


FIG. 3. A, Corticotropin-induced hyperplasia. The adrenal is large and nodular, but the overall anatomy and shape of the gland is largely preserved. B and C, Two examples, right and left adrenal gland, respectively, with multiple nodules or discrete adenomas and intervening atrophic cortical tissue (type I AIMAH, BMAH). D, Hematoxylin and eosin (H&E) staining (magnification, $\times 5$) of the tissue from C; multiple nodules are clearly visible, and the intervening cortex is atrophic and even sandwiched between adenomas (arrows). E and F, Two examples, right and left adrenal gland, respectively, with diffuse cortical hyperplasia and no residual normal or surrounding atrophic adrenal cortex (type II AIMAH). G, Histology of tissue from E showing a large adenoma in the context of diffuse hyperplasia (H&E staining, $\times 5$). H, Histology of tissue from F showing diffuse cortical hyperplasia (H&E staining, $\times 5$).

patients had family history of ACTH-independent CS or any other endocrine condition. One (ADT43.01) presented with AIMAH and hyperparathyroidism, and he was found to have a *MEN1* mutation (Pro494Leu) (28); another (ADT06.01) presented with AIMAH and the hereditary leiomyomatosis and renal cancer syndrome; she had a heterozygous fumarate hydratase mutation (32); a third patient (ADT52.02) had an *APC* gene mutation (4393_4394delAG) also in the heterozygote state, and her history included polyps and desmoids tumors. This mutation has been previously reported in patients with familial adenomatous polyposis and is predicted to produce a truncated protein (see <http://www.hgmd.cf.ac.uk/ac/index.php>). There was one additional patient who was found to have a somatic

TABLE 2. Response to testing for aberrant receptor expression in patients with AIMAH and ACS

	AIMAH (n = 14)		ACS (n = 12)	
		%		%
Posture	4/11	36.4	3/12	25.0
Meal	1/12	8.3	0/11	0.0
GnRH	1/6	16.7	0/8	0.0
TRH	1/3	33.3	1/8	12.5
GHRH	2/11	18.2	0/10	0.0
Glucagon	0/10	0.0	2/11	18.2
Vasopressin	5/11	45.5	4/10	40.0

The denominator stands for the number of patients being tested; the numerator stands for the number of patients with positive response (>150% increasing).

GNAS mutation (Arg201His) in her adrenocortical tumor tissue only; the patient did not have any other signs of McCune-Albright syndrome and was in that sense similar to the patients reported by Frago *et al.* (33). There were no pathogenic mutations in the *PRKAR1A* gene; one of the three familial AIMAH cases was a carrier of the R867G *PDE11A* gene polymorphism (27, 34). None of the other patients, in any of the groups, was found to carry a mutation in any of the tested genes in the peripheral or tumor DNA.

Tumors in other organs in our AIMAH, ACS, APA, and SCA patients were found in five, two, one, and two cases, respectively. In their families, there were three AIMAH patients with family history of tumors other than in the adrenal gland; the SCA groups had three such patients (Table 3).

Histology of patients with AIMAH

It was recognized that patients with AIMAH could generally be subgrouped in two categories (18) (Fig. 3): those with multiple nodules or discrete adenomas and intervening atrophic cortical tissue [type I AIMAH, bilateral adenomata, or bilateral macronodular adrenocortical hyperplasia (BMAH)] and those with dif-

fuse hyperplasia and no residual normal or surrounding atrophic adrenal cortex (type II AIMAH). Most patients with AIMAH belonged to the second category; the three familial cases and the patients with germline *MEN1* and *APC* and the one with the somatic *GNAS* mutation belonged to the first group.

Discussion

Although the majority of our patients with AIMAH presented in the fifth decade of life (mean age of 46.8 yr), most had an insidious and atypical form of CS for a number of years. It is characteristic that by the time these patients were operated at the National Institutes of Health and despite their lack of suppression in response to dexamethasone as well as relative loss of their diurnal rhythm in cortisol secretion, almost all of them retained their response to oCRH. As a group, AIMAH patients had higher morning cortisol levels and lower oCRH responses than patients with APAs or SCAs, but their hypothalamic-pituitary-adrenal axis was not completely suppressed. This is a phenomenon that we have described before in patients with PPNAD (17, 35), suggesting that oCRH testing is not useful in the investigation of bilateral adrenocortical hyperplasias (BAH) that are generally associated with atypical, milder, chronic, or cyclical forms of CS.

In the absence of the typical paradoxical rise in glucocorticoid hormone excretion rates in response to dexamethasone for mi-

TABLE 3. Mutations and other tumors in patients with ACTH-independent macronodular adrenal hyperplasia and other adrenocortical tumors

	Familial cases	Mutation	Other tumors (number of cases)	Other tumors in family members (number)
AIMAH (n = 16)	3	<i>MEN1</i> (Pro494Leu) <i>FH</i> (c.781del7) <i>APC</i> (c. 4393_4394delAG) <i>GNAS</i> (Arg201His), somatic	Thyroid adenoma (1) Lymphoma (1) Uterine fibroids (5) Parotid tumor (3) Parathyroid adenoma (1)	Thyroid cancer (1) in M Prostate cancer (1) in F Lung cancer (1) in M
ACS (n = 15)	None	None	Parathyroid adenoma (1) Nodular goiter (1)	None
APA (n = 19)	None	None	Thyroid nodule (1)	None
SCA (n = 32)	None	None	Thyroid nodule (1) Parathyroid adenoma (1)	Pancreatic Ca (F, GF on maternal side) Uterine Ca (M) Cervical Ca (S) Breast Ca (A) Pituitary tumor (F)

M, Mother; F, father; GF, grandfather; S, sister; A, aunt.

cronodular forms of BAH (16–18, 35), how can one diagnose AIMAH in its early stages? The suggestion of BAH on radiological imaging is the first diagnostic criterion, as we proposed elsewhere (16, 36). The present investigation identified another useful diagnostic feature for patients with AIMAH: these patients had high urinary 17OHS excretion, even when their UFC levels were comparable or even lower than those of the ACS group. Urinary 17OHS has long been known to represent the fractions of the corticosteroids possessing a hydroxyl group at position 17 of the steroid structure, including cortisol, cortisone, 6 β -hydroxycortisol, tetrahydrocortisol, allotetrahydrocortisol, and tetrahydrocortisone (37, 38). Thus, patients with AIMAH appear to excrete in their urine various glucocorticoid metabolites that are not detected by the assays for UFCs; measurement of 17OHS can therefore be a useful and early diagnostic test for patients with suspected atypical or early CS due to AIMAH.

It should be noted that in two patients with ACS, there was a rise in glucocorticoid secretion during Liddle's test, albeit modest compared with what has been seen in PPNAD and other micronodular forms of BAH (16, 35, 39). This is consistent with our previous data (16) and may indicate that at least some patients with ACS may be affected by a mild form of micronodular BAH or that *PRKAR1A* mutations or related genetic defects that were not detected by routine sequencing were present at the tissue level, as we have shown before in at least some cases of AIMAH or ACS (26, 40).

Nine AIMAH patients showed aberrant cortisol responses to at least one stimulus; this response rate was consistent with what has been reported in other studies that used a similar testing protocol (10–12, 14, 41, 42). In addition, patients with ACS also showed aberrant receptor expression; again, this has also been reported to be a relatively frequent phenomenon (14, 41–45). To date, the unexpected regulation of cortisol secretion by receptors for various neuroendocrine substances in ACTH-independent CS remains a biological puzzle; both mouse and human studies indicate that in at least some of these patients, this phenomenon is a primary defect, causative of the CS phenotype (41).

Our clinical and molecular genetic data pointed to significant heterogeneity among our cohort of patients with AIMAH: there were three cases with some (but not extensive) family history of adrenocortical tumors. There are so far seven reports of familial AIMAH, all pointing to an autosomal dominant transmission (31, 41, 45–47). Evidence for heterogeneity included the type of aberrant receptor expression involved in the AIMAH phenotype, the association with various tumors, and various types of germline mutations. First, the aberrant receptors associated with familial AIMAH have most frequently been reported to be the vasopressin (V_1 and V_2) and the β -adrenergic receptors; the serotonin receptor was present in one family (31, 41, 45–47). In our study, two of the three familial cases had aberrant vasopressin receptors. Second, we found thyroid, parathyroid, and uterine leiomyomatous tumors that have all been previously reported in other patients with AIMAH (31, 41, 45–47); we also found three cases with parotid tumors that have not been previously reported in association with AIMAH.

Finally, an astonishing three of 16 patients (19%) had germline genetic defects in *MEN1*, *FH*, and *APC*, all previously reported pathogenic mutations in the respective genes. A fourth patient had a *GNAS* somatic (tumor) mutation (R201H) like two of the patients with AIMAH reported by Fragoso *et al.* (33); none of these patients had any signs of McCune-Albright syndrome. Another patient (with familial AIMAH) carried the R867G *PDE11A* gene variant (27, 34), which, although a relatively frequent gene variant, is located in a highly conserved region of the *PDE11A* gene and affects enzymatic activity *in vitro* (34).

Histologically we were able to subclassify our patients with AIMAH in two groups, those with multiple nodules or discrete adenomas and intervening atrophic cortical tissue (type I AIMAH) and those with diffuse hyperplasia (type II AIMAH). Most of our patients belonged to the second category; the three familial cases and the patients with germline *MEN1* and *APC* and somatic *GNAS* mutations belonged to the first group. However, we found no association of this histopathological observation with distinct expressed receptor pattern or any other clin-

ical feature; it remains unclear whether this observation reflects simply different stages of changes in the adrenal cortex during the development of a chronic disease, or it is in fact a direct effect of the underlying molecular etiology.

We conclude that AIMAH is a heterogeneous disorder that is often associated with genetic defects at both the germline or the somatic level. Determination of urinary 17OHS is a useful diagnostic test for the early detection of abnormal glucocorticoid secretion in this condition; oCRH testing, on the other hand, is not helpful. Histological subgrouping may assist in the future in further investigating the molecular causes of this fascinating condition.

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