

Measuring adrenal autoantibody response: Interlaboratory concordance in the first international serum exchange for the determination of 21-hydroxylase autoantibodies $\stackrel{\circ}{\sim}$

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KEYWORDS

Adrenal insufficiency; Autoimmune Addison's disease; Autoimmune polyendocrine syndrome; Radioimmunoassay; Standardization Abstract 21-hydroxylase autoantibodies (210HAb) are the gold standard immune marker to identify patients with clinical or subclinical autoimmune Addison's disease (AAD). No assessment of interlaboratory concordance has been made for 210HAb measurement. Serum samples from 267 patients with primary adrenal insufficiency and from 83 healthy control subjects were distributed to four independent laboratories that determined presence and titer of 210HAb, by using radiobinding assays with either *in vitro* translated ³⁵S-radiolabelled or ¹²⁵I-radiolabelled autoantigen. Cohen's κ of inter-rater agreement ranged from 0.857 to 0.983, showing a very good concordance of the positive/negative score among the four laboratories. Passing–Bablok regression showed a good agreement of 210HAb titers arranged by ranks, but important discrepancies emerged at the Bland–

Abbreviations 210HAb, 21-hydroxylase autoantibodies; AAD, Autoimmune Addison's Disease; ACA, Adrenal Cortex Autoantibodies; AHC, adrenal hypoplasia congenita; ALD, adrenoleukodystrophy; APS, autoimmune polyendocrine syndrome; AUC, area under ROC curve; CI, confidence interval; IAN, Italian Addison Network; ICC, intra-class correlation coefficient; PAI, primary adrenal insufficiency; RC, repeatability coefficient; ROC, receiver-operating characteristic; TBC, tuberculosis; T1DM, type 1 diabetes mellitus.

* Disclosure statement: RSR Ltd is a manufacturer of medical diagnostics including kits for 210H autoantibodies.

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Altman plot, as the repeatability coefficient was much higher than the laboratory cut-offs, which indicates that results from different laboratories cannot be used interchangeably. A standardization international program for 210HAb measurement is strongly needed. © 2011 Elsevier Inc. All rights reserved.

1. Introduction

Primary adrenal insufficiency (PAI) affects approximately 1 in 7000 individuals [1,2]. The primitive adrenocortical deficit has an heterogeneous etiopathogenesis and may develop through several distinct mechanisms, including autoimmunity, infiltrative adrenalitis, adrenoleukodystrophy (ALD), genetic disorders, metastasis, adrenal hemorrhage, surgery, sepsis, infections or toxic agents [3,4]. Autoimmune Addison's Disease (AAD) is caused by an autoimmune process responsible for the selective destruction of adrenal cortex cells [5,6]. Although AAD is the result of a T-cell mediated process, adrenal autoimmunity is made evident by the appearance of circulating adrenal cortex autoantibodies (ACA), that represent the best immune marker to identify patients with AAD [5,6]. ACA have little or no pathogenetic role [5-8], but their detection in human serum is clinically useful for both the etiological classification of PAI [9] and the identification of subjects at high-risk for future development of clinical AAD [10-16].

From 1963 until the middle of the 1990s, ACA were exclusively detected by means of the indirect immunofluorescence technique on cryostatic sections of human or animal adrenal glands [17]. The identification of the steroidogenic enzyme 21-hydroxylase as the main autoantigen identified by ACA [18,19] led to the development of sensitive and specific immunoassays for the detection of 21-hydroxylase autoantibodies (210HAb) in human serum [20-22]. Several subsequent studies have demonstrated that 210HAb is the gold standard immune marker for diagnosis of clinical and pre-clinical AAD [5,6,10–16,20–29]. In Europe, from 1974 to 2010, over 2000 patients with PAI have been reported in the literature as

Table 1	210HAb positivity in different clinical forms of	
PAI, in 4 d	fferent laboratories.	

	PG	RM	CAR	DEN
Idiopathic	168/205	168/205	173/205	170/205
	(82%)	(82%)	(84.4%)	(82.9%)
APS1	14/15	14/15	13/15	14/15
	(93.3%)	(93.3%)	(86.7%)	(93.3%)
ALD	0/8	0/8	2/8 (25%)	0/8
AHC	0/1	0/1	0/1	0/1
Post-sepsis	0/1	0/1	0/1	0/1
Post-surgical	0/1	1/1	0/1	0/1
Post-TBC	0/36	0/36	1/36	1/36
			(2.8%)	(2.8%)
Healthy control subjects	0/83	0/83	2/83 (2.4%)	0/83

AHC: adrenal hypoplasia congenita; ALD: adrenoleukodystrophy; APS1: autoimmune polyendocrine syndrome type 1; Post-TBC: post-tuberculosis primary adrenal insufficiency.

having been evaluated for adrenal autoantibodies [5,6]. The prevalence of adrenal autoantibodies ranged from 44.5 to 94% in different studies [5,6].

The Italian Addison Network (IAN) has developed a comprehensive flow-chart for the etiological classification of PAI which takes into consideration immunological, biochemical and imaging data [9]. In this flow-chart, the analysis of adrenal autoantibodies, and more specifically of 210HAb, plays a major role to discriminate autoimmune from nonautoimmune forms of the disease [9]. Studies on subjects with organ-specific autoimmune diseases have shown that not only the presence, but also levels, of 210HAb may have clinical relevance in the estimates of future risk for development of the clinical signs of AAD [13].

In the IAN study already mentioned [9], a comparison of 210HAb assays in two independent laboratories and of ACA in two other independent laboratories was performed. The concordance rate of the two laboratories that performed 210HAb assays was higher than that of the two laboratories that performed ACA assays [9]. As an extension of the previous IAN study, a novel study was planned in which four independent laboratories (two from Italy and two from other countries) performed 210HAb assays in a large series of samples from patients with PAI and healthy control subjects. The aim of the present study was specifically that of comparing 210HAb results generated by different laboratories using their own internal standard sera and in-house calculated cut-offs, to define the need for future standardization programs aimed at harmonizing the methods and identifying a common standard serum. Accordingly, serum samples were consecutively collected from PAI patients irrespective of disease duration or etiology.

2. Materials and methods

2.1. Serum samples

Serum samples from 267 patients (107 males and 160 females) with PAI and from 83 healthy control subjects (35 males and 48 females, median age 39 years, range 21-55 years) were collected by the IAN, as previously described [9], and stored at -70 °C until subsequently used for the present study. At the time of sample collection, the median patient age was 49 years (range, 6-87 years) and the median disease duration was 5 years (range, 0-53 years). In all patients, clinical symptoms and signs of PAI were associated with low basal cortisol ($<3 \mu g/dl$) and high basal ACTH (>100 pg/ml) levels. For statistical purposes, PAI patients were subdivided into two groups: a) clinically idiopathic/APS1 (n=220; this group included 15 cases with autoimmune polyendocrine syndrome type 1), and b) with demonstrated non-autoimmune etiology (such as postinfiltrative adrenalitis, n=36, ALD, n=8, adrenal hypoplasia

Table 2	Cohen's κ of inter-rater agreement among the four laboratories.					
	RM	CAR	DEN			
PG RM	0.983 95% CI: 0.963–1.000	0.880 95% CI: 0.830–0.929 0.862	0.971 95% Cl: 0.946–0.996 0.954			
CAR		95% CI: 0.809–0.915	95% Cl: 0.923–0.986 0.885 95% Cl: 0.836–0.934			

congenita, n=1, sepsis, n=1 or surgical adrenalectomy, n=1). All patients gave their written informed consent for this study.

All sera were coded by the laboratory of the University of Perugia, Italy and redistributed, blindly, to all the participating laboratories, as frozen aliquots shipped in dry ice. Altogether, four independent laboratories were involved in this study for testing 210HAb by different techniques: Perugia (Italy) (lab code: PG), Rome (Italy) (lab code: RM), Denver (CO, USA) (lab code: DEN) and Cardiff (UK) (lab code: CAR).

2.2. 210HAb assays

Three laboratories (PG, RM and DEN) used a radiobinding assay with *in vitro* translated recombinant human ³⁵S-21OH and immunoprecipitation of the immunocomplexes using protein A-Sepharose beads [21]. Autoantibody levels were evaluated by counting the immunoprecipitated radioactivity in a multiwell-plate scintillation counter (Top Count, Canberra Packard) and expressed as a relative index based on the analysis of one positive and two negative standard sera in each assay (210HAb index) using the formula: (cpm sample-mean cpm negative controls) / (cpm positive control-mean cpm negative controls). The cut-offs defined by the three laboratories were 0.060, 0.060 and 0.150 for PG, RM and DEN, respectively. Our laboratory in Perugia reported previously an intra-assay CV of 7-12% [21].

Our laboratory in Cardiff used a radiobinding assay with 1251 and the immunoprecipitation of immunocomplexes by protein A [22]. The radiobinding was counted on a gammacounter. The cut off was established as 1 unit/mL (arbitrary units). Interassay precision on 25 determinations of the same serum sample was mean of 7.5 units/mL with SD=0.49 and CV=6.58%. Intraassay precision obtained with 25 determinations of the same serum sample was mean of 5.7 units/mL with SD=0.32 and CV of 5.6%.

2.3. Statistical methods

The degree of concordance for the dichotomous variable presence/absence of 210HAb between two different laboratories was estimated using the Cohen's κ -test of interrater agreement [30] (κ -value ranging from 0 to 1), with Fleiss–Cuzick extension [31] when appropriate. Interpretation of strength of agreement was made according to the Landis and Koch's gradation: <0.2 = poor, 0.2–0.4 = fair, 0.4–0.6 = moderate, 0.6–0.8 = good, 0.8–1.0 = very good [32]. Passing–Bablok regression analysis [33] was performed for each participating laboratory using the median rank of each sample in the four laboratories and the rank of each sample

in each laboratory as variables. This type of analysis does not depend on the assignation of each variable as x or y and enables calculation of the regression slope and y axis intercept, which are then used to determine if they differ significantly from the ideal slope=1 and intercept=0. Linearity was evaluated by CUSUM test which rejects the null hypothesis of parameter constancy whenever the range of cumulated ordinary least squares residuals becomes too large [34].

The Bland–Altman plot technique [35], or difference plot technique, was used to compare quantitative 210HAb assay results in two independent laboratories. This method was used only for the three assays that expressed 210HAb levels as a relative index (210HAb index). In this graphical method, the differences between the two techniques were plotted against the averages of the two techniques. Intra-class correlation coefficient (ICC value ranging from 0 to 1), defined as the proportion of variance of an observation due to between-subject variability in the true scores, assesses rating reliability by comparing the variability of different ratings of the same subject to the total variation across all ratings and all subjects. The type of ICC was 2,1 [36]. A high ICC indicates that there is little variation between the scores given to each item by the raters. Kendall's τ correlation coefficient between intra-subject standard deviation and intra-subject mean, used to assess the interdependence of these two parameters, were also calculated. The repeatability coefficient (RC) was used as a precision measure which represents the value below which the absolute difference between two repeated test results may be expected to lie with a probability of 95%.

Diagnostic sensitivity (as determined in patients with clinically idiopathic PAI/APS1, n=220) and specificity (as determined in healthy control subjects, n=83) were estimated by using receiver-operating characteristic (ROC) curves with cut-offs corresponding to the best combination of high sensitivity and high specificity. Differences in area under ROC curves (AUC) were tested with modified Z-test [37]. In all analysis, a *p* value<0.05 was considered significant. Statistical analyses were performed using Predictive Analytic Software (PASW) release 17.0.2, (SPSS Inc., Chicago, USA, 2009), and StatsDirect version 2.7.2 (StatsDirect Ltd, Altrincham, Cheshire, UK, 2008).

3. Results

Prevalence of 210HAb in different forms of PAI is reported in Table 1. 210HAb were detected in 82–84.4% patients with idiopathic PAI, 86.7–93.3% patients with APS1, 0 to 25% patients with ALD and 0 to 2.8% patients with post-TBC PAI.

One serum from a post-surgical PAI patient was found negative in three assays and positive in one assay. On the other hand, the serum from a patient with post-sepsis PAI and the serum from a patient with adrenal hypoplasia congenita were found negative in all four assays. Positivity of 210HAb in 83 healthy control subjects ranged from 0 to 2.4%. To evaluate qualitative concordance among the four autoantibody assays, namely classification of subjects as positive or negative, we used the Cohen's κ of inter-rater agreement. The general agreement over the two categories with four raters per subject (evaluated by using the Fleiss–Cuzick extension) was very good (κ =0.923, 95% CI: 0.880–0.965, p<0.0001). A very good agreement was also observed in the comparisons of each laboratory with each other (Table 2).

Globally, 175 samples were found positive and 151 were found negative in all the four laboratories (representing 93% of the total samples having a 100% concordance in classification as positive or negative).

To evaluate diagnostic sensitivity and specificity of the four 210HAb assays, ROC curves were also generated (Fig. 1), and AUC for each laboratory was calculated. AUC was high in all the four laboratories ranging from 0.900 (95% CI: 0.865–0.935) to 0.969 (95% CI: 0.952–0.987). Statistically significant differences in AUC were observed between DEN and PG (p=0.002), DEN and CAR (p<0.001) and between RM and CAR (0.011).

To analyze quantitative results, conversion of 210HAb values as ranks was needed, because one of the laboratories

expressed its results using different measure units. When rank in each laboratory was plotted against median rank in the four laboratories, no particular trend related to single laboratories was observed (Fig. 2). A clear subdivision in two distinct subpopulations (positive and negative values) was evident. The two subpopulations showed a different pattern, as negative results (left side of the graph) appeared more dispersed than positive results (right side of the graph).

Passing-Bablok test was then performed for each laboratory. No significant deviation from linearity was observed in any of the tested laboratories (CUSUM test, p>0.10). Slope of the regression line resulted 0.967 (95% CI: 0.927-1.009) for PG, 0.990 (95% CI: 0956-1.024) for RM, 1.021 (95% CI: 0.980-1.062) for CAR and 1.025 (95% CI: 1.000-1.057) for DEN. Intercept resulted 13.11 (95% CI: 3.165-23.33) for PG, 5.450 (95% CI: -0.8095 to 12.0263) for RM, -4.08 (95% CI: -13.92-5.636) for CAR and -5.612 (95% CI: -12.44-0) for DEN. PG data were more dispersed than those generated in the other laboratories, as 0 was not included in the 95% CI of the intercept for that laboratory. We hypothesized that this was likely caused by a higher dispersion of negative values than in the other laboratories. To test this hypothesis, we repeated the Passing-Bablok analysis taking into consideration only the positive results (Fig. 3). When negative samples were excluded, slope of the regression line was 0.980 (95% CI: 0.936-1.023) for PG, 1.000 (95% CI: 0.938-1.062) for RM, 0.944 (95% CI: 0.876-1.006) for CAR and 1.000 (95% CI: 0.959-1.053) for DEN. On the other hand,



Figure 1 Receiving-operator-characteristic (ROC) curves for PG (panel A), RM (panel B), CAR (panel C) and DEN (panel D). AUC: Area under the ROC curve.



Figure 2 210HAb data generated by PG (circles), RM (squares), CAR (triangles) and DEN (diamonds) expressed as median rank in the four laboratories (x axis) vs. rank in the single laboratory (y axis).

intercept results were 9.687 (95% CI: -1.454-22.10) for PG, 2.000 (95% CI: -12.5000 to 17.9538) for RM, 16.72 (95% CI: 0.770-35.01) for CAR and 1.000 (95% CI: -11.45-10.76) for DEN. Accordingly, analysis of positive results was associated with an improvement of the regression line of PG (which supports the hypothesis that most of the dispersion was due to negative values, in that laboratory), but not of CAR, for which a significant deviation from linearity appeared (p < 0.05).

Agreement of the continuous 210HAb index among PG, RM and DEN was tested using the Bland–Altman plot analysis. Although all the three intra-class correlation coefficients were rather high, being 0.709 for the comparison between PG and RM, 0.765 for the comparison between PG and DEN and 0.849 for the comparison between RM and DEN (which indicates that high levels in one laboratory tend to correspond to high levels also in the other). 95% limits of agreement of 210HAb index were very broad, ranging from -0.675 to 0.888 for the comparison between PG and RM (Fig. 4A), from -0.708 to 0.832 for the comparison between PG and DEN (Fig. 4B) and from -0.219 to 0.555 for the comparison between RM and DEN (Fig. 4C). In all cases, cut-offs of the three laboratories were included within the 95% limits of agreement, which indicates that differences between laboratories may potentially have clinical relevance and the different 210HAb tests cannot be used interchangeably. Similarly, the repeatability coefficient was much higher than the three laboratory cut-offs, corresponding to a 210HAb index of 0.808 for the comparison between PG and RM, 0.779 for the comparison between PG and DEN and 0.508 for the comparison between RM and DEN. Correlation between mean values in the same subject and the correspondent standard deviation was highly significant (p < 0.0001) in all three comparisons (Figs. 4D–F), which indicates that the error was strongly dependent on the size of the measure, being higher for samples with higher 210HAb index.

4. Discussion

210HAb is the best single immune marker useful to both identify patients with clinical AAD and subjects at high-risk to develop clinical and biochemical signs of adrenal insufficiency



Figure 3 Passing–Bablok regression test for PG, RM, CAR and DEN. Data are expressed as median rank in the four laboratories (x axis) vs. rank in the single laboratory (y axis).



Figure 4 Bland-Altman plot analysis: agreement plot (panel A, B and C) and repeatability plot (panels D, E, F).

[5,6,9–29]. More specifically, the unequivocal diagnosis of AAD is based on the detection of 210HAb in serum samples. Accordingly, imprecisions in adrenal autoantibody tests may have important effects on clinical management of PAI, as a low diagnostic specificity may determine an erroneous classification of ALD patients as AAD, or, in the case of a low diagnostic sensitivity, individuals with pre-clinical AAD may go undiagnosed until an Addisonian crisis occurs. Hence, it is important that currently available tests for determination of 210HAb be compared and standardized. Although 210HAb have been tested in several studies in patients with PAI or other diseases, no attempt at standardizing this test has yet been made. The long-term experience of several workshops for the standardization of islet autoantibody determinations [38–41] is highly valuable as a model for the future organization of standardization programs for 210HAb and other steroid-cell autoantibodies. A complicating factor is that diagnosis of AAD depends on the detection of adrenal autoantibodies and is not independently formulated on pure clinical grounds, as it is in the case of T1DM. Accordingly, in our present study, we tested diagnostic sensitivity in patients with clinically idiopathic PAI, compared to normal control subjects, after exclusion of known non-autoimmune causes of the disease. This may have resulted in an underestimation of the actual diagnostic sensitivity of 210HAb for AAD. In addition, patients with long-term disease duration were also enrolled in our study, and this may further have reduced diagnostic sensitivity, as we cannot exclude that some samples scored negative because of the disappearance of previously present autoantibodies. However, the aim of the present study was not to estimate the frequency of this marker in newly diagnosed AAD patients, but exclusively that of comparing the autoantibody results among different laboratories, independently from etiological classification of PAI and disease duration, and the above limitations have little impact on the observed results. In addition, the study was not aimed at identifying a common standard serum to be used in laboratories throughout the world, but at providing initial information on the agreement for 210HAb assays in selected research laboratories with long-term experience in the field. More specifically, we did not attempt to harmonize the assays before the samples were tested, nor to address traceability using common materials, as our study intended to use the data as they were generated by the laboratories using internal standards and in-house calculated cut-offs. Accordingly, we did not calculate a common study cut-off, because this would mean attempting to standardize the different methods, which was beyond the scope of our present investigation and is more appropriate with a larger number of participating laboratories.

In a previous IAN study [9], 210HAb and ACA assays were each compared in two independent Italian laboratories. That study demonstrated the higher inter-laboratory concordance and higher diagnostic accuracy of 210HAb for the diagnosis of AAD, as compared to ACA [9]. Subsequently, a large series of serum samples from patients with PAI and healthy control subjects collected by IAN were randomized and blindly distributed to four laboratories, which have been very active in adrenal and other autoantibody research in the past. Our present study represents the first international serum exchange for comparison of 210HAb determination.

Statistical analyses were aimed at evaluating concordance on two distinct aspects of 210HAb determination: qualitative results (positive/negative score) and quantitative data (210HAb titer). However, not all the laboratories expressed 210HAb titer using similar scale units and some analyses could not be performed for all laboratories.

The Cohen's k-test showed very good inter-laboratory agreement on positive/negative scores, which confirms that diagnostic accuracy of 210HAb for AAD is extremely high, as shown by previous publications from the participating laboratories as well as from other groups. To evaluate diagnostic sensitivity and specificity of the four assays we constructed ROC curves, using the results obtained with sera from idiopathic/APS1 AAD patients and from healthy control subjects, and we calculated the area under each curve (AUC). All four ROC curve results were very good and the AUC was higher than 0.9 in all cases (Fig. 1). Again, as previously discussed, each calculated AUC might not be totally accurate, as diagnostic sensitivity was determined using patients for which an autoimmune origin could not be demonstrated in all cases. Nevertheless, one aim of the study was to compare the AUCs generated by the different laboratories, and a good agreement was observed with this analysis. The results of our study are in line with those observed for other organ-specific autoantibodies tested using similar assay types [40]. A slightly lower AUC was observed for CAR as compared to RM and DEN, which is mostly due to a somewhat lower diagnostic specificity for that laboratory as compared to the other three participating laboratories, even though it showed the highest diagnostic sensitivity (Table 1). However, the simple calculation of the area under the ROC curve may lead to erroneous conclusions as this value may be influenced by variables (such as dispersion of negatively scored samples) that have little impact on clinical application. Accordingly, the statistically significant differences observed in the ROC analysis among different laboratories do not necessarily correspond to actual differences in diagnostic sensitivity and specificity reported by each laboratory (Table 1).

To compare quantitative data, two separate statistical analyses were performed. In the first series of analysis, the results from each laboratory were plotted vs. the global results of the four laboratories. As the participating laboratories expressed their data using different scale units, samples were transformed in ascending rank (from the lowest to the highest) in each laboratory and the median rank from all laboratories was also determined. This type of analysis has previously been used also to compare data for islet autoantibodies from different laboratories [38]. It was evident that a large dispersion of the values below the cut-offs was present among the different laboratories (Fig. 2). More specifically, PG had the highest dispersion of the values of the samples scored as "negative" for 210HAb. Passing-Bablok regression analysis [33] was chosen because it does not depend on the assignation of each variable as x or y and enables calculation of the regression slope and y axis intercept. Using this test, all the laboratories had regression slopes and y axis intercepts which were not significantly different from 1 to 0, respectively, with the exception of PG, for which 95% confidence intervals of the intercept did not include 0, mostly because of the already mentioned high dispersion of samples scored negative. When the analysis was repeated taking into consideration only samples scored positive for 210HAb, a significant deviation from linearity was detected only for CAR. This lack of linearity of CAR was however caused by only 2 samples considered positive by this lab and clearly negative for the remaining three (Fig. 3C). Globally, all the laboratories performed highly satisfactorily in the Passing-Bablok analysis and the observed small deviations from the expected, ideal line were due either to the negative samples (for which the actual 210HAbindex has no clinical relevance) or to a very limited number of samples discordant with other laboratories.

The final analysis aimed at comparing 210HAbindex between two laboratories. CAR was not included in the Bland-Altman plot test because of the different scale units used. This analysis revealed major discrepancies among the three laboratories that may potentially have clinical relevance. In all comparisons, the repeatability coefficient was much higher than the three laboratory cut-offs, thus showing that the results generated by the three laboratories cannot be used interchangeably. Interestingly, the higher the 210HAbindex the higher was the observed difference among laboratories. This phenomenon may in part reduce the clinical impact of the inter-laboratory discrepancies, as one may expect that for samples with low 210HAbindex a lower discrepancy might exist, but the lack of agreement in 210HAbindex requires additional, larger international programs of standardization of 210HAb determination.

5. Conclusions

We have reported the results of the first international serum exchange for the determination of 210HAb. The four participating laboratories showed a high agreement as positive/negative score, but some major discrepancies emerged for the quantitation of 210HAb titer. The organization of a standardization study aimed at identifying a standard serum that may enable expression of results in common international units is strongly recommended. The identification of the international standard serum will also be instrumental for the harmonization of the different assays and the calculation of a uniform cut-off level.

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