

# Body Iron Stores and Glucose Intolerance in Premenopausal Women

Role of hyperandrogenism, insulin resistance, and genomic variants related to inflammation, oxidative stress, and iron metabolism

M. ANGELES MARTÍNEZ-GARCÍA, PHD<sup>1</sup>  
MANUEL LUQUE-RAMÍREZ, MD, PHD<sup>1</sup>

JOSÉ L. SAN-MILLÁN, PHD<sup>2</sup>  
HÉCTOR F. ESCOBAR-MORREALE, MD, PHD<sup>1</sup>

**OBJECTIVE**— Increased serum ferritin levels and iron stores may be involved in the development of abnormal glucose tolerance in women presenting with obesity and/or polycystic ovary syndrome (PCOS). We aimed to study the determinants of serum ferritin levels in premenopausal women among indexes of insulin resistance, adiposity, hyperandrogenism, and genotypes pertaining to inflammation, oxidative stress, and iron metabolism.

**RESEARCH DESIGN AND METHODS**— A total of 257 premenopausal women, classified depending on the presence or absence of PCOS, obesity, and/or abnormal glucose tolerance, underwent a complete metabolic evaluation, serum ferritin, haptoglobin, and C-reactive protein (CRP) measurements, and genotyping for proinflammatory and prooxidant variants and mutations in the *HFE* gene.

**RESULTS**— Serum ferritin concentrations were increased in women presenting with PCOS and/or abnormal glucose tolerance, independent of obesity. A stepwise multivariate linear regression analysis ( $R^2 = 0.18$ ,  $P < 0.0001$ ) retained menstrual dysfunction ( $\beta = 0.14$ ,  $P = 0.035$ ), free testosterone ( $\beta = 0.14$ ,  $P = 0.052$ ), insulin sensitivity index ( $\beta = -0.12$ ,  $P = 0.012$ ), the His63Asp variant in *HFE* ( $\beta = 0.16$ ,  $P = 0.008$ ), and abnormal glucose tolerance ( $\beta = 0.15$ ,  $P = 0.015$ ) as significant predictors of the logarithm of ferritin levels, whereas CRP, haptoglobin, waist-to-hip ratio, or variants in the *TNFA*, *TNFRSF1B*, *IL6*, *IL6ST*, *IL6R $\alpha$* , *PON1*, and *HFE* Cys282Tyr mutation exerted no influence.

**CONCLUSIONS**— Androgen excess (partly because of hyperandrogenemia and partly because of menstrual dysfunction), insulin resistance, abnormal glucose tolerance, and the *HFE* His63Asp variant correlate with ferritin levels in premenopausal women.

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Mounting evidence suggests that increased body iron stores are involved in the pathogenesis of insulin-resistant disorders such as the metabolic syndrome and type 2 diabetes in the general population (1). This involvement appears to be bidirectional, because not only does iron accumulation favor insulin resistance and may contrib-

ute to pancreatic  $\beta$ -cell dysfunction and diabetes, but also insulin resistance may in turn facilitate iron accumulation within the body (2,3). Interestingly, oxidative stress and inflammation are involved in the interplay between iron overload and insulin resistance (2,4).

Although most of the evidence linking iron metabolism and disorders of glu-

ucose metabolism has been provided from the study of middle-aged or older men and of postmenopausal women (2,5), body iron stores also influence insulin resistance and glucose metabolism in premenopausal women (6,7).

Ferritin is the cellular storage protein for iron. We have reported that serum ferritin levels are increased in women presenting either with obesity or polycystic ovary syndrome (PCOS) and especially when both conditions are present in the same patient (6). Therefore, both androgen excess and insulin resistance may underlie this finding. In addition, because these changes occurred independent of changes in serum inflammatory markers, the increased ferritin level indicates that body iron stores are actually increased in these women and do not result from the secondary role of ferritin as an acute-phase marker (6).

Of note, serum ferritin levels were clearly increased in the small subset of these women presenting with abnormal glucose tolerance, both in the PCOS and nonhyperandrogenic subgroups (6). This finding suggests that increased body iron stores could be related to the development of abnormalities in glucose metabolism in these patients, because progressive iron accumulation in the pancreas is a recognized pathogenic mechanism of disorders of glucose tolerance in patients with iron overload (8).

We hypothesized that the reduced menstrual losses due to the oligomenorrhea present in most women with PCOS could contribute to their increased iron stores, yet recent data from our group suggest that insulin resistance is actually one of the major players explaining their increased serum ferritin levels: whereas serum ferritin levels did not change after restoring regular menses by using an oral contraceptive for 6 months, these levels decreased markedly after insulin sensitization with metformin (3).

The present study was undertaken with the aim of identifying, in a large series of premenopausal women, the determinants of increased serum ferritin levels

From the <sup>1</sup>Department of Endocrinology, Hospital Universitario Ramón y Cajal and Universidad de Alcalá, Madrid, Spain and Centro de Investigaciones Biomédicas en Red de Diabetes y Enfermedades Metabólicas Asociadas, Barcelona, Spain; and <sup>2</sup>Department of Molecular Genetics, Hospital Universitario Ramón y Cajal and Universidad de Alcalá, Madrid, Spain and Centro de Investigaciones Biomédicas en Red de Diabetes y Enfermedades Metabólicas Asociadas, Barcelona, Spain.

Corresponding author: Héctor F. Escobar-Morreale, hescobarm.hrc@salud.madrid.org.

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M.L.-R. is currently affiliated with the Department of Endocrinology, Hospital Universitario de La Princesa, Madrid, Spain.

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(an index of body iron stores and a risk factor for the development of abnormalities of glucose tolerance) among markers of hyperandrogenism, adiposity, insulin resistance, and genomic variants related to chronic inflammation, oxidative stress, and iron metabolism.

## RESEARCH DESIGN AND METHODS

A total of 257 premenopausal women were included. The group was composed of consecutive patients reporting to the Department of Endocrinology because of PCOS and/or weight excess and of healthy nonhyperandrogenic nonobese volunteers recruited from the staff of Hospital Universitario Ramón y Cajal. Women were classified according to their BMI into obese (BMI  $\geq 30$  kg/m<sup>2</sup>,  $n = 128$ ) and nonobese (BMI  $< 30$  kg/m<sup>2</sup>,  $n = 129$ ) subgroups. PCOS was diagnosed in 149 women presenting with clinical and/or biochemical hyperandrogenism in addition to oligo-ovulation as reported previously (9), thereby fulfilling all of the current definitions of the syndrome (10–12), whereas 108 women showed no sign of hyperandrogenism, had normal androgen levels, and had regular ovulatory menstrual cycles.

Menstrual and ovulatory dysfunction were defined by the presence of oligomenorrhea (cycles longer than 35 days) or amenorrhea (absence of menstrual bleeding for at least three usual cycle lengths) or, in women presenting with regular menstrual cycles (cycles between 26 and 35 days), by lack of ovulation according to body temperature charts and/or serum progesterone levels  $< 12.7$  nmol/l during the luteal phase of the menstrual cycle (9).

None of the women had a personal history of hypertension, diabetes, or cardiovascular events. Women who took oral contraceptives, antiandrogens, insulin sensitizers, iron supplements, or drugs that might interfere with blood pressure regulation, lipid profile, or carbohydrate metabolism within the previous 6 months or who were referred for any medical reason aside from androgen and/or weight excess were automatically excluded. Written informed consent was obtained from all of the participants, and the study was approved by the ethics committee of Hospital Universitario Ramón y Cajal.

Clinical and anthropometric variables, including the hirsutism score, BMI, waist circumference, and waist-to-hip ratio (WHR) were determined. WHR was calculated by dividing the minimal waist circumference by the hip circumference

at the level of the greater trochanters, using a nonstretchable measuring tape.

Whole blood, serum, and plasma samples were obtained between days 5 and 10 of the menstrual cycle, or during amenorrhea after pregnancy was excluded. After a 3-day 300-g carbohydrate diet and 12-h overnight fasting, samples were obtained early in the morning for the measurement of total testosterone, sex hormone-binding globulin, 17-hydroxyprogesterone, androstenedione, dehydroepiandrosterone sulfate, luteinizing hormone, follicle-stimulating hormone, estradiol, thyrotropin, and prolactin. A complete hemogram and serum biochemistry and lipid profiles were also obtained. Then, a 75-g oral glucose tolerance test was performed, and samples were obtained for measurement of serum insulin and plasma glucose at 0, 30, 60, 90, and 120 min. Samples were immediately centrifuged, and serum and plasma were separated and frozen at  $-20^{\circ}\text{C}$  until assayed.

The assays used for these measurements have been described in detail elsewhere (9). Free testosterone concentrations were calculated from total testosterone and sex hormone-binding globulin levels (9). Serum ferritin and C-reactive protein (CRP) concentrations were measured by automated immunochemiluminescence (Immulite 2000 Ferritin and High Sensitivity CRP; Diagnostic Products Corporation, Los Angeles, CA) with lower limits of detection of 0.88 pmol/l and 0.1 mg/l, respectively, and intra- and interassay coefficients of variation  $< 10\%$ . The circulating concentrations of haptoglobin were assayed by a commercial immunonephelometry method (Dade Behring, Marburg, Germany), calibrated against the international CRM 470 reference material. The composite insulin sensitivity index was calculated from circulating glucose and insulin concentrations during the oral glucose tolerance test as described by Matsuda and DeFronzo (13). Disorders of glucose tolerance were diagnosed following the recommendations of the American Diabetes Association (14).

## DNA analysis

Genomic DNA was obtained from whole blood samples using a Nucleon BACC3 DNA isolation kit (Amersham, Buckinghamshire, U.K.) and was used to genotype several polymorphisms related to inflammation or oxidative stress that had been previously found to be associated with metabolic disorders such as PCOS,

obesity, diabetes, and insulin resistance and variants involved in iron metabolism. Genotyping of the dinucleotide CA repeat in intron 8 of the interleukin (IL)-6 receptor- $\alpha$  (*IL6R $\alpha$* ) locus (rs57636717) was performed by PCR using a fluorescently labeled forward primer (15). Amplified fragments were resolved by capillary electrophoresis on an ABI Prism 3100 automated genetic analyzer (Applied Biosystems, Foster City, CA), and their sizes were determined using GeneMapper 4.0 software. We detected 13 different alleles with sizes ranging from 143 to 169 bp, of which the 149-bp allele was the most frequent. For statistical analyses, subjects were genotyped as homozygous for 149-bp alleles, heterozygous for 149-bp allele (15). Several genomic variants were analyzed by PCR-restriction fragment-length polymorphism as described previously: variants His63Asp (c.187C>G) and Cys282Tyr (c.845G>A) in exons 2 and 4 (rs1799945 and rs1800562, respectively) of the *HFE* gene (7); polymorphism  $-308\text{G}>\text{A}$  (rs1800629) in the promoter of the tumor necrosis factor (TNF)- $\alpha$  (*TNF $\alpha$* ) gene (16); variant Met196Arg (c.587T>G; rs1061622) in exon 6 of the TNF receptor 2 (*TNFRSF1B*) gene (17); polymorphism  $-174\text{G}>\text{C}$  (rs1800795) in the IL-6 gene promoter (18); variant Gly148Arg (c.442G>C; rs2228044) in exon 5 of the IL-6 signal transducer (*IL6ST*) gene (15); and polymorphism  $-108\text{C}>\text{T}$  (rs705379) in the paraoxonase (*PON1*) gene (19).

## Statistical analysis

Data are shown as means  $\pm$  SD unless otherwise stated. The Kolmogorov-Smirnov statistic was applied to continuous variables. Logarithmic or square-root transformations were applied as needed to ensure normal distribution of the variables. Univariate general linear models were used to evaluate the influence of obesity, PCOS, and glucose tolerance on clinical and biochemical variables, with introduction of age as a covariate to control for possible age differences among the groups. A multivariate linear regression analysis was used to determine the influence of clinical, biochemical, metabolic, and genetic variants on the logarithm of serum ferritin concentrations as described below. Depending on the median of the whole population of premenopausal women, serum ferritin levels were categorized into values below, or equal to or above, the median value. The associa-

Table 1—Influence of obesity and PCOS on clinical and biochemical variables of premenopausal women

n	Nonobese women (n = 129)		Obese women (n = 128)		PCOS (n = 149) vs. control (n = 108)		Obese (n = 128) vs. nonobese (n = 129)		Interaction	
	PCOS patients	Nonhyperandrogenic women	PCOS patients	Nonhyperandrogenic women	F/ $\chi^2$	P value	F/ $\chi^2$	P value	F	P value
Age (years)	24 ± 6	29 ± 7	27 ± 7	33 ± 8	43.97	<0.001	11.79	0.001	0.19	0.664
BMI (kg/m <sup>2</sup> )	24.4 ± 3.5	24.4 ± 4.1	36.9 ± 5.5	36.7 ± 5.4	1.73	0.190	404.80	<0.001	0.09	0.764
Waist circumference (cm)	73 ± 10	76 ± 11	97 ± 13	96 ± 13	0.94	0.334	212.62	<0.001	2.17	0.142
WHR	0.75 ± 0.07	0.76 ± 0.08	0.82 ± 0.08	0.80 ± 0.08	8.37	0.004	26.50	<0.001	2.30	0.131
Hirsutism score	11 ± 6	2 ± 2	10 ± 6	1 ± 2	217.83	<0.001	3.36	0.068	2.27	0.133
Free testosterone (pmol/l)	37 ± 18	19 ± 8	54 ± 25	24 ± 10	116.10	<0.001	33.60	<0.001	1.82	0.179
Estradiol (pmol/l)	134 ± 80	245 ± 233	180 ± 129	231 ± 198	10.80	0.001	0.414	0.521	4.17	0.042
Luteinizing hormone (units/l)	6.6 ± 3.9	5.3 ± 3.1	6.1 ± 3.8	4.9 ± 3.1	5.20	0.023	0.902	0.343	0.01	0.995
Follicle-stimulating hormone (units/l)	5.7 ± 4.2	6.4 ± 6.1	6.2 ± 4.5	5.8 ± 1.8	0.361	0.548	0.01	0.948	0.59	0.442
Fasting insulin (pmol/l)	75 ± 58	58 ± 46	137 ± 75	84 ± 44	15.19	<0.001	46.69	<0.001	1.77	0.185
Fasting glucose (mmol/l)	4.9 ± 0.4	5.0 ± 0.4	5.3 ± 0.5	5.2 ± 0.5	0.02	0.890	20.08	<0.001	3.86	0.051
Insulin sensitivity index	5.4 ± 3.0	7.0 ± 3.8	2.8 ± 2.1	4.8 ± 3.6	27.34	<0.001	52.23	<0.001	3.53	0.061
CRP (mg/l)	1.3 ± 1.7	1.3 ± 1.6	6.0 ± 5.6	5.6 ± 5.1	1.68	0.196	161.77	<0.001	0.29	0.558
Haptoglobin (μmol/l)	12 ± 4	12 ± 4	17 ± 5	17 ± 5	0.01	0.969	66.13	<0.001	0.38	0.539
Regular menstruation	16 (20)	47 (100)	10 (15)	61 (100)						
Oligomenorrhea	47 (57)	0 (0)	35 (52)	0 (0)	171.00	<0.001	2.45	0.294	—	—
Amenorrhea	19 (23)	0 (0)	22 (33)	0 (0)						
Normal glucose tolerance	74 (90)	40 (85)	46 (70)	43 (70)	0.512	0.474	13.72	<0.001	—	—
Abnormal glucose tolerance	8 (10)	7 (15)	21 (31)	18 (30)						

Data are means ± SD or raw numbers (%). Continuous data were used for univariate general linear models and, because age was different between patients with PCOS and nonhyperandrogenic women and between obese and nonobese women, age was introduced as a covariate in the analysis of all the other variables. Categorical data were analyzed by  $\chi^2$  tests.

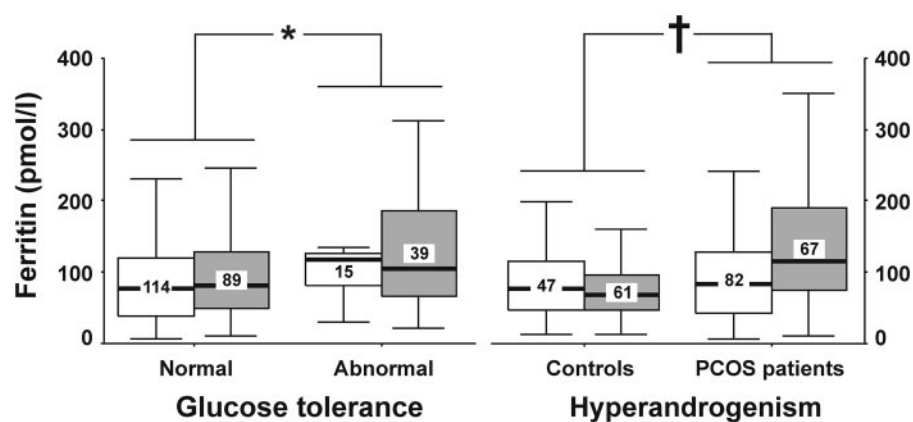
tions of these two groups with categorical and continuous variables were analyzed by  $\chi^2$  and Fisher's exact tests as appropriate.  $P < 0.05$  was considered statistically significant. Analyses were performed using SPSS Statistics 17 (SPSS, Chicago, IL).

**RESULTS**— The influence of PCOS and of obesity on clinical and biochemical variables is summarized in Table 1. Because patients with PCOS were younger than nonhyperandrogenic women and obese women were older than nonobese women, the influence of age was controlled by introducing this variable as a covariate in the comparisons described below.

Patients with PCOS had increased serum androgen and luteinizing hormone levels and decreased estradiol concentrations and were insulin resistant and had increased WHR compared with their nonhyperandrogenic counterparts irrespective of obesity. Obesity was characterized by increased indexes of insulin resistance, increased frequency of abnormalities in glucose tolerance, and increased serum CRP, haptoglobin, and free testosterone levels compared with those in nonobese women, both in patients with PCOS and in nonhyperandrogenic women.

Serum ferritin levels were influenced independently by PCOS and glucose intolerance (Fig. 1). Patients with PCOS presented with increased serum ferritin levels compared with those in nonhyperandrogenic women. When considered as a whole, women presenting with abnormal glucose tolerance had higher serum ferritin concentrations compared with those showing normal glucose values during fasting and after an oral glucose tolerance test. Of note, these results occurred both in the nonobese and in the obese subgroups, and obesity did not influence serum ferritin concentrations after controlling for both PCOS and glucose tolerance (Fig. 1).

Because serum ferritin levels were not distributed normally, their logarithm was introduced as a dependent variable in multiple linear regression models using age; BMI; glucose tolerance (codified as normal or abnormal); menstrual cycles (codified as regular menstruation, oligomenorrhea, or amenorrhea); serum free testosterone, CRP, and haptoglobin levels; the insulin sensitivity index; and the genomic variants related to iron metabolism, inflammation, and oxidative stress as independent variables. The model that considered all independent variables ex-



**Figure 1**—Serum ferritin levels depending on the presence or absence of abnormalities of glucose tolerance and PCOS in nonobese □ and obese ■ women. The box plot includes the median (horizontal line) and the interquartile range, and the whiskers indicate the minimum and maximum data values, unless outliers are present, in which case the whiskers extend to a maximum of 1.5 times the interquartile range. The figures inside the boxes are the numbers of women in each subgroup. The logarithm of serum ferritin levels was used in a general linear model in which glucose tolerance, PCOS, and obesity were introduced as independent variables and age was introduced as a covariate to correct for the difference in age between patients and control subjects and between obese and nonobese women. No interaction was found among independent variables. \* $P = 0.001$  between women presenting with or without abnormalities of glucose tolerance. † $P = 0.034$  between patients with PCOS and nonhyperandrogenic control women.

plained 22% of the variability in the logarithm of serum ferritin concentrations ( $R^2 = 0.22$ ,  $P < 0.0001$ ) (Fig. 2).

Conversely, when the independent variables were introduced using a stepwise method (probability to enter  $\leq 0.05$ ; probability to remove  $\geq 0.10$ ), the model ( $R^2 = 0.18$ ,  $P < 0.0001$ ) retained only menstrual dysfunction ( $\beta = 0.14$ ,  $P = 0.035$ ), serum free testosterone levels ( $\beta = 0.14$ ,  $P = 0.052$ ), insulin sensitivity index ( $\beta = -0.12$ ,  $P = 0.012$ ), His63Asp variant in *HFE* ( $\beta = 0.16$ ,  $P = 0.008$ ), and abnormal glucose tolerance ( $\beta = 0.15$ ,  $P = 0.015$ ) as significant predictors of the logarithm of serum ferritin levels, whereas all the other clinical, biochemical, and genomic variables were excluded (Fig. 2). Finally, a similar stepwise regression method ( $R^2 = 0.17$ ,  $P < 0.0001$ ) adjusted for age and BMI (these variables were manually entered into the model) retained only menstrual dysfunction ( $\beta = 0.18$ ,  $P = 0.003$ ), abnormal glucose tolerance ( $\beta = 0.17$ ,  $P = 0.013$ ), His63Asp variant in *HFE* ( $\beta = 0.17$ ,  $P = 0.004$ ), and insulin sensitivity index ( $\beta = -0.15$ ,  $P = 0.031$ ) as predictors of the logarithm of serum ferritin levels (Fig. 2).

When considering the median of the serum ferritin concentrations of the whole population (83 pmol/l) as a cutoff value, premenopausal women presenting with a serum ferritin level above the median had an odds ratio (OR) for abnormal

glucose tolerance of 2.4 (95% CI 1.3–4.4,  $\chi^2 = 7.420$ ,  $P = 0.009$ ) and an OR for PCOS of 2.2 (1.3–3.7,  $\chi^2 = 9.524$ ,  $P = 0.002$ ).

**CONCLUSIONS**— The metabolic consequences of iron overload are exemplified by the development of abnormalities of glucose tolerance in primary or secondary hemochromatosis. However, less severe iron overload also plays an important role in the development of abnormalities in glucose tolerance (2), as demonstrated by the improvement in insulin resistance and glucose tolerance in type 2 diabetic patients after the iron depletion achieved by repeated blood letting (20) or the higher insulin sensitivity associated with reduced iron stores in frequent blood donors from the general population (21).

Our present results demonstrate that serum ferritin levels, an accurate marker of body iron stores in the absence of acute inflammatory syndromes (2), are also related to abnormal glucose tolerance in premenopausal women. Serum ferritin levels above the median were associated with a 2.4-fold OR for presenting with abnormal glucose tolerance, and ferritin concentrations were clearly higher in women in whom fasting or postload glucose levels were above the normal range. Furthermore, our present results suggest that the body iron stores of premeno-

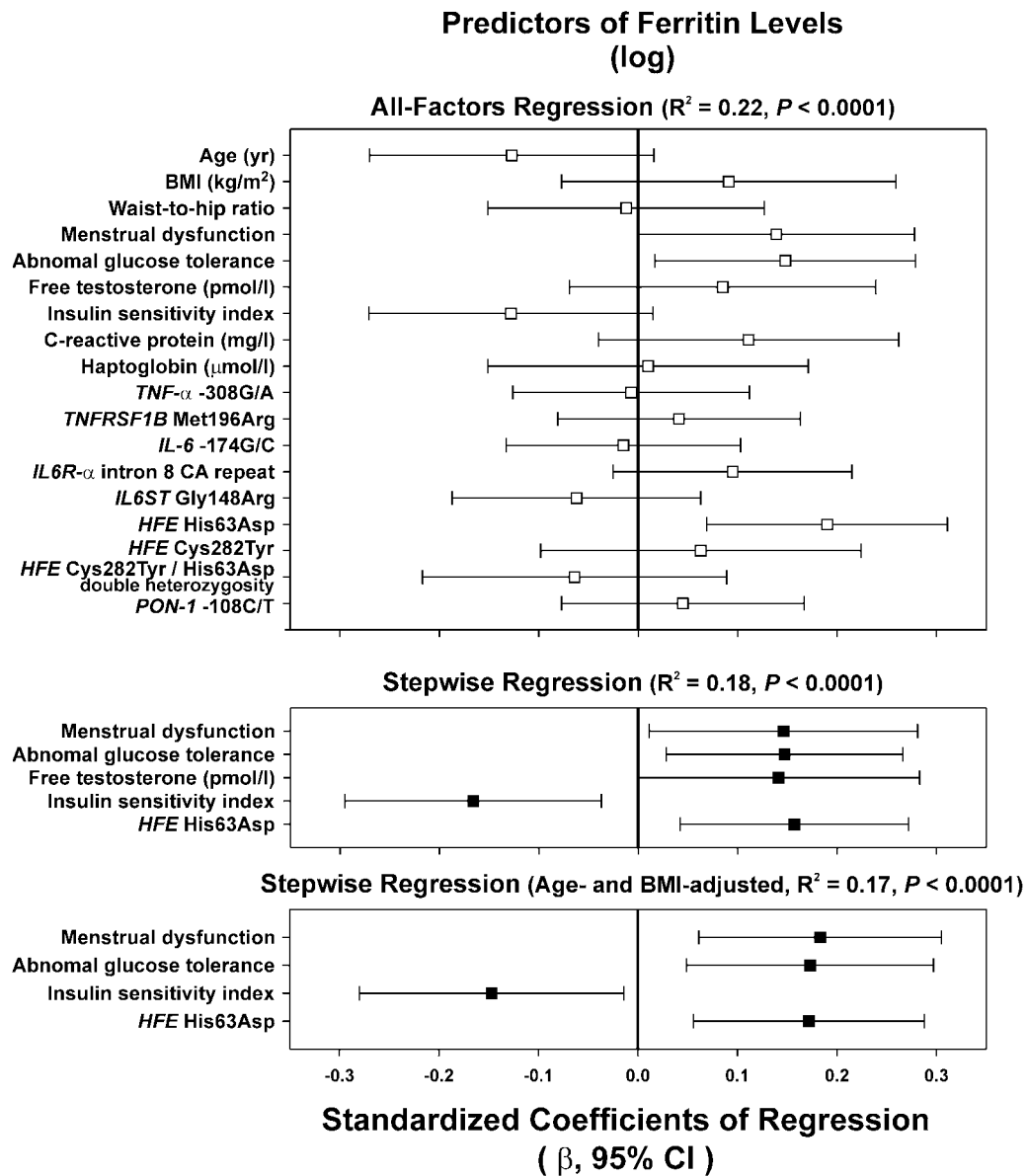
pausal women are associated with several factors including menstrual dysfunction, insulin resistance, and the His63Asp variant in *HFE* even after controlling for the difference in age among patients with PCOS and nonhyperandrogenic control subjects and between obese and nonobese women.

We have reported previously, in a much smaller series, that body iron stores are increased in premenopausal women presenting with PCOS and obesity and that these associations were independent of mutations in *HFE* and were not influenced by changes in markers of chronic inflammation (6,7). Our present findings confirm that serum ferritin levels are increased in patients with PCOS and that, accordingly, women presenting with serum ferritin levels above the median have a 2.2 OR of having PCOS. Yet when controlling for the presence or absence of abnormal glucose tolerance, the previously reported association with obesity disappears, suggesting that it is the increased prevalence of disordered glucose tolerance in obese women and not weight excess by itself that is responsible for increased body iron stores in these women.

According to the regression analyses presented here, menstrual dysfunction, increased androgen levels, and insulin resistance, together with the *HFE* His63Asp variant and abnormal glucose tolerance, are among the strongest predictors of serum ferritin levels. On the contrary, these levels are not influenced by indexes of global or abdominal adiposity or by genomic variants related to chronic inflammation and oxidative stress.

Therefore, it appears that androgen excess and insulin resistance, which are present in many patients with PCOS, collaborate in increasing body iron stores in premenopausal women. The effects of insulin resistance and hyperinsulinism on body iron stores might depend on a direct insulin stimulation of intestinal iron absorption by upregulating activity of hypoxia-inducible factor-1 $\alpha$  and downregulating hepcidin expression (22,23) and may be counteracted in patients with PCOS by administering insulin sensitizers (3).

The effect of androgen excess on body iron stores might result from the well-known stimulatory effect of androgens on erythropoiesis, thereby increasing intestinal iron absorption (24) but may also result from the iron-sparing effect of reduced menstrual losses due to the chronic menstrual dysfunction of PCOS.



**Figure 2**—Multiple linear regression analysis of the logarithm of serum ferritin concentrations. The squares are the standardized regression coefficients ( $\beta$ , the change in terms of SDs in the dependent variable that results from a change of 1 SD in an independent variable), and the error bars indicate the 95% CI of  $\beta$ . Menstrual history and genomic variants were coded as dummy variables: regular menstruation was coded 0, 1 was used for oligomenorrhea, and 2 was used for amenorrhea. Variants in *TNF $\alpha$* , *TNFRSF1B*, *IL6*, *IL6ST*, *HFE*, and *PON1* loci were coded as 0 for homozygosity for wild-type alleles, 1 for heterozygosity, and 2 for homozygosity for mutant alleles. The *IL6R $\alpha$*  polymorphism was coded 0 for homozygosity for 149-bp alleles, 1 for subjects carrying only one 149-bp allele, and 2 for subjects carrying two non-149-bp alleles. Finally, *HFE* His63Asp/Cys282Tyr double heterozygotes were coded 1 and subjects without double heterozygosity were coded 0.

This iron-sparing mechanism may take years to result in increased iron stores, explaining why regularization of menstrual bleeding in patients with PCOS by administering antiandrogenic contraceptive pills for 6 months had no evident impact on serum ferritin levels in our previous report (3).

Our present results also suggest that the His63Asp variant in *HFE* influences body iron stores in premenopausal women, in conceptual agreement with

the partial loss of *HFE* function induced by this mutation in animal models, leading to a variable degree of hepatic iron loading (25). However, more importantly, the relatively strong association of increased serum ferritin levels with abnormal glucose tolerance raises the possibility that increased iron stores played some pathogenic role in the development of such metabolic derangements, given that progressive iron accumulation in the pancreas contributes to  $\beta$ -cell dysfunction

and abnormal glucose tolerance in syndromes of iron overload (8). However, because association does not imply causality, this hypothesis is speculative.

In summary, body iron stores, as reflected by serum ferritin concentrations, are interrelated with androgen excess, insulin resistance, and the His63Asp variant in *HFE* in premenopausal women and are associated with the development of abnormal glucose tolerance in this particular population.

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