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Original Article

Association between Inflammatory and Obesity Markers in a Swiss Population-Based Sample (CoLaus Study)

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Key Words

IL-1 β · IL-6 · TNF- α · hs-CRP · BMI · Waist circumference · Body composition · Population study

Abstract

Objective: To assess the associations between obesity markers (BMI, waist circumference and %body fat) and inflammatory markers (interleukin-1β (IL-1β); interleukin-6 (IL-6); tumor necrosis factor-α (TNF-α) and high-sensitivity C-reactive protein (hs-CRP)). *Methods:* Population sample of 2,884 men and 3,201 women aged 35-75 years. Associations were assessed using ridge regression adjusting for age, leisure-time physical activity, and smoking. Results: No differences were found in IL-1\(\beta\) levels between participants with increased obesity markers and healthy counterparts; multivariate regression showed %body fat to be negatively associated with IL-1β. Participants with high %body fat or abdominal obesity had higher IL-6 levels, but no independent association between IL-6 levels and obesity markers was found on multivariate regression. Participants with abdominal obesity had higher TNF- α levels, and positive associations were found between TNF- α levels and waist circumference in men and between TNF- α levels and BMI in women. Obese participants had higher hs-CRP levels, and these differences persisted after multivariate adjustment; similarly, positive associations were found between hs-CRP levels and all obesity markers studied. Conclusion: Obesity markers are differentially associated with cytokine levels. %Body fat is negatively associated with IL-1β; BMI (in women) and waist circumference (in men) are associated with TNF- α ; all obesity markers are positively associated with hs-CRP. Copyright © 2012 S. Karger GmbH, Freiburg



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Introduction

Cytokines, such as interleukin 1 β (IL-1 β), interleukin 6 (IL-6), tumor necrosis factor (TNF- α) and C-reactive protein (CRP) have important effects in inflammation and atherosclerosis. Elevated proinflammatory cytokine levels and high-sensitive CRP (hs-CRP) have been associated with an increased risk of developing coronary heart disease [1] or diabetes [2].

It has been suggested that obesity is a chronic inflammatory condition [3], with increased production of cytokines and hs-CRP by adipocytes [4]. Still, the effects of each obesity marker (BMI, waist circumference or %body fat) on cytokine levels remain to be clarified. For instance, BMI does not differentiate between lean and fat mass [5], while waist circumference relates only to the abdominal accumulation of fat. Further, most studies which assessed the independent effects of each obesity marker on cytokine levels did so by multivariate linear regression [6, 7], a method that might lead to inflated estimates when multicollinearity (i.e. correlation) of the independent variables (i.e. obesity markers) exists. Multicollinearity can distort the interpretation of a model by increased inaccuracy of the regression coefficients and their standard errors [8]. Regression coefficients biased by collinearity might cause variables that have no significant relationship with the outcome (i.e. cytokine levels) when considered in isolation to become highly significant in conjunction with collinear variables, thus increasing type I error (false-positive results). Multiple regression coefficients might also show no statistical significance due to large confidence intervals, thus increasing type II error (false-negative results).

Hence, we used the data from the large, population-based CoLaus study to assess the association of different obesity markers with cytokine levels by ridge regression, a method whose goal is to circumvent the problem of collinearity between independent variables [9]. To our knowledge, this is currently the largest population study that has cytokine levels measured.

Participants and Methods

Recruitment

The CoLaus Study was approved by the Institutional Ethics Committee of the University of Lausanne. The CoLaus Study is a cross-sectional study aimed at assessing the prevalence and identifying the molecular determinants of cardiovascular risk factors in the Caucasian population of Lausanne, Switzerland, a town of 117,161 inhabitants, of which 79,420 are of Swiss nationality. The sampling procedure of the CoLaus Study has been described previously [10]. Briefly, the complete list of Lausanne inhabitants aged 35–75 years (n = 56,694) was provided by the population registry of the city. A simple, nonstratified random sample of 35% of the overall population was drawn. The following inclusion criteria were applied: i) written informed consent; ii) age 35–75 years; iii) willingness to take part in the examination and to donate blood samples; iv) Caucasian origin, defined as having both parents and grandparents born in a restricted list of countries (available from the authors). Recruitment began in June 2003 and ended in May 2006. Participation rate was 41%.

All participants attended the outpatient clinic of the University Hospital of Lausanne in the morning after an overnight fast. Data were collected by trained field interviewers in a single visit lasting about 60 min.

Lifestyle and Clinical Data

According to their smoking histories, participants were classified as never, current, or former smokers. Leisure-time physical activity was considered if the participant took part for at least 20 min in leisure physical activities (tennis, running, football) two or more times per week.





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Anthropometric Measurements and Bioimpedance

Body weight and height were measured with participants standing without shoes in light indoor clothes. Body weight was measured in kilograms to the nearest 100 g using a Seca® scale, which was calibrated regularly. Height was measured to the nearest 5 mm using a Seca® height gauge. Overweight was defined as a BMI \geq 25 and \leq 30 kg/m²; obesity was defined as a BMI \geq 30 kg/m².

Waist circumference was measured with a non-stretchable tape over the unclothed abdomen at the narrowest point between the lowest rib and the iliac crest. Two measures were made and the mean (expressed in centimeters) was used for analyses. Abdominal obesity was considered for a waist circumference >102 cm for men and >88 cm for women [11].

Body composition was assessed by electrical bioelectrical impedance analysis (BIA) using the Bodystat® 1500 analyzer (Isle of Man, British Isles, UK). This device has been validated against reference methods [12]. The coefficient of variation for repeated measurements of electrical impedance is 0.7% [12]; the reproducibility has been shown to be between 1% [12] and 5% [13]; the intraclass correlation coefficient has been estimated at 0.99 [14], and its correlation with reference methods such as isotope dilution is between 0.86 [13] and 0.96 [15]. All metallic adornments were removed, and measurement was performed after a 10-min rest in the supine position. Care was taken that the subject did not touch any metallic component of the bed and that the inner part of the thighs did not touch each other. The electrodes were positioned in the right side of the body according to the manufacturer's recommendations: injecting leads were connected to the electrodes behind the finger and toe, whereas measuring leads were connected to the electrodes on the wrist and ankle. Measurement was conducted with a test current of 400 µA (root mean square) at a frequency of 50 kHz, with an impedance measuring range of 30-1,000 Ω (accuracy 3 Ω). Conversely, no information was available regarding the equation used (proprietary). Body fat was expressed as percentage of total body weight (%body fat), and participants in the highest sex-specific quartile (>28% in men and >40% in women) were considered as presenting with high body fat.

Cytokine Measurement

Venous blood samples (50 ml) were drawn in the fasting state. hs-CRP was assessed by immunoassay and latex HS (IMMULITE 1000-High, Diagnostic Products Corporation, Los Angeles, CA, USA) with maximum intra- and interbatch coefficients of variation of 1.3% and 4.6%, respectively. Serum was preferred to plasma as it has been shown that different anticoagulants may affect absolute cytokine levels differently [16]. It has also been stated that cytokine levels in a single blood sample may be useful biomarkers of inflammation in population-based studies of obesity-related diseases [16]. Serum samples were kept at $-80\,^{\circ}$ C before assessment of the other cytokines and sent on dry ice to the laboratory. Cytokine levels were measured using a multiplexed particle-based flow cytometric cytokine assay, a methodology used in other studies [17]. Milliplex kits were purchased from Millipore (Zug, Switzerland). The procedures closely followed the manufacturer's instructions. The analysis was conducted using a conventional flow cytometer (FC500 MPL, BeckmanCoulter, Nyon, Switzerland). Lower detection limits for IL-1 β , IL-6, and TNF- α were 0.2 pg/ml in each case. Data regarding agreement between signal and cytokine levels, intra- and interassay coefficients of variation and correlation between repeated measurements have been reported elsewhere [18].

Statistical Analysis

Statistical analysis was conducted using SAS v.9.2 (SAS Inc, Cary, NC, USA), stratifying by gender. Quantitative variables (apart from inflammatory biomarkers) were expressed as mean \pm standard deviation and qualitative variables as number of participants and percentage. Inflammatory biomarkers were presented as percentage of values below detection limit, median and interquartile range of measured values, and as exponentiated multivariate-adjusted mean \pm standard error. Undetectable values were included in the first quartile. Between groups comparisons were performed using Student's t-test or Kruskall-Wallis nonparametric test for quantitative and chi-square test for qualitative variables. The bivariate association between cytokines and obesity markers was assessed by Spearman nonparametric correlation.

The association between the different obesity markers (BMI, waist circumference, and %body fat) and inflammatory biomarkers was assessed by multivariate analysis including all obesity markers in the same model. As age, leisure-time physical activity, and smoking have been associated with cytokine levels, they were included in all multivariate models. The association of each obesity marker taken separately





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was assessed by multivariate linear regression on log-transformed values of detectable IL-18, Il-6, TNF-α, and hs-CRP. As BMI, waist circumference, and %body fat are closely associated, their independent associations with inflammatory markers were assessed in a second multivariate analysis using ridge regression applied on all obesity markers. Ridge regression is a variant of ordinary multiple linear regression whose goal is to circumvent the problem of predictors' collinearity [9]. When the predictor variables are highly correlated amongst themselves, the coefficients of the resulting least squares fit may be very imprecise. By allowing a small amount of bias in the estimates, more reasonable coefficients may often be obtained. Briefly, ridge regression implies that a constant number δ (penalty) is added to the elements on the diagonal of the matrix to be inverted. Let X be an $n \times (p+1)$ matrix of the p independent variables obtained in n subjects; let X^T be the transposed matrix of X; let Y be an $n \times 1$ vector corresponding to the dependent variable (i.e. IL-1β levels). The usual linear regression estimates (β values) are obtained via the matrix product $(X^T \times X)^{-1} \times X^T \times Y$. The ridge regression estimates are obtained via the matrix product $(X^T \times X)^{-1} \times Y$. $(X + \delta I)^{-1} \times X^T \times Y$, where I is the p \times p identity matrix, X is an n \times p matrix of the centered independent variables, and δ is the penalty. It has been shown that ridge regression estimators have smaller mean square error than the ordinary least-squares estimators, provided δ is small enough [19]. Prior to analysis, the effect on the variance inflation factor (VIF) of ridge parameter values from 0 to 0.24 was assessed for each gender and the value of 0.18 (corresponding to a VIF close to unity) was chosen. Statistical significance was considered for p < 0.05.

Results

Clinical Characteristics of Participants

Of the 6,188 initial participants, 103 (1.7%) were excluded because of missing cytokine data, and a further 55 (0.9%) because of missing data for at least one of the anthropometric markers. The clinical characteristics of the remaining 6,030 (2,863 men and 3,167 women) are summarized in table 1. Men were significantly younger, had a higher BMI, and smoked more than women. Conversely, no gender differences were found for leisure-time physical activity.

Among the 6,030 participants, 2,289 (37.6%), 451 (7.4%) and 43 (0.7%) had IL-1 β , IL-6 and TNF- α levels below detection limits, respectively. The distribution of IL-1 β , IL-6, TNF- α , and hs-CRP levels stratified by gender according to the different obesity markers studied are summarized in tables 2 and 3.

Associations between Obesity Markers and Cytokine Levels

Men with high BMI had lower IL-1 β levels, but this association was no longer significant after multivariate analysis (table 2). In women, high %body fat and abdominal obesity were associated with lower IL-1 β levels, but this association was no longer significant after multivariate adjustment (table 2). %Body fat and waist circumference (men only) were significantly and negatively correlated with IL-1 β levels, while no association was found for BMI (table 4). On multivariate ridge regression, %body fat was independently and significantly associated with IL-1 β levels in women, with a borderline significant association (p = 0.053) in men (table 4).

In both genders, IL-6 levels increased with greater levels of BMI, waist circumference, and %body fat. After multivariate adjustment, these associations remained significant in women while in men only the association with abdominal obesity remained significant (table 2). In both genders, all obesity markers were individually and positively correlated with IL-6 levels (table 4), but this association was no longer significant after multivariate ridge regression (table 4).

In both genders, TNF- α levels increased with greater levels of BMI, waist circumference, and %body fat (table 3); after multivariate adjustment these associations remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained significant in women while in men only the association with abdominal obesity remained with the significant in women while in men only the association with a significant in women while while while while we will not be a significant while whil





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Table 1. Clinical characteristics of the sample, by gender^a

	Men (N = 2,863)	Women (N = 3,167)	Test
Age, years	52.6 ± 10.8	53.5 ± 10.7	3.52***
Smoking status (%)			
Never	921 (32.2)	1,496 (47.2)	
Former	1,106 (38.6)	887 (28.0)	147.6***
Current	836 (29.2)	784 (24.8)	
Leisure-time physical activity (%)	1,838 (64.2)	2,058 (65)	0.40 NS
BMI, kg/m ²	26.6 ± 4.0	25.1 ± 4.8	12.77***
BMI categories (%)			
Normal	1,078 (37.7)	1,812 (57.2)	
Overweight	1,306 (45.6)	902 (28.5)	246.4***
Obese	479 (16.7)	453 (14.3)	
Waist circumference, cm	96 ± 11	83 ± 12	40.30***
Abdominal obesity (%)	681 (23.8)	974 (30.8)	36.67***
%Body fat	23.8 ± 6.1	34.3 ± 8.2	56.20***
Dosable IL-1β (%)	132 (60.5)	225 (63.9)	7.60**
IL-1β, pg/ml	1.10 (0.45-3.97)	1.28 (0.52-3.76)	3.65 ^{NS}
Dosable IL-6 (%)	278 (93.5)	206 (91.8)	6.95**
IL-6, pg/ml	1.58 (0.77-3.85)	1.37 (0.67-3.23)	19.68***
Dosable TNF-α (%)	244 (99.3)	343 (99.2)	0.19^{NS}
TNF-α, pg/ml	3.06 (1.90-4.66)	2.75 (1.73-4.42)	23.65***
hs-CRP, mg/dl	1.2 (0.6-2.5)	1.3 (0.6-2.9)	6.74**

^aResults are expressed as mean \pm standard deviation (except cytokines), median and (interquartile range) of dosable values (cytokines) or as number of subjects and (percentage). Statistical analysis by Student's t-test or Kruskall-Wallis nonparametric test for quantitative variables and by chi-square for qualitative variables: ^{NS}not significant; *p < 0.05; **p < 0.01; ***p < 0.001.

icant (table 3). Taken individually, all obesity markers were significantly and positively correlated with TNF- α . Conversely, after multivariate ridge regression only BMI (in women) and waist circumference (in men) were significantly and positively associated with TNF- α (table 4).

In both genders, hs-CRP levels increased with greater levels of BMI, waist circumference, and %body fat, and these associations remained significant after multivariate adjustment (table 3). All obesity markers were significantly and positively associated with hs-CRP, both on bivariate correlation or on multivariate ridge regression (table 4).

Discussion

To our knowledge, this is currently the largest population-based sample in which the associations between different obesity markers and IL-1 β , IL-6, TNF- α , and hs-CRP have been assessed. Further, most studies [6, 7] assessed the independent associations of each obesity marker with cytokine levels by multivariate linear regression, which is prone to error when the independent variables are correlated In this study, the results of the linear regression were confirmed by a method which corrects for such a correlation and by using log-transformed values to reduce skewness.

IL-1β comes from the interleukin 1 family, which comprises 11 different proteins encoded by 11 distinct genes. In mice, IL-1 induces anorexia [20], although no differences in





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Table 2. IL-1 β and IL-6 distribution according to different markers of obesity, by gender¹

v L Men (N = 2,863)	L-1β values > LOD (N) ^a	median (IQR) ^b	multi-	IL-6 values >	median (IQR) ^b	multi-
Men (N = 2,863)		median (IQR) ^b		values >	median (IOR)b	multi
			adjusted ^c	LOD (N ⁾ a	median (rgiv)	adjusted ^c
DMI 8						
BMI categories §						
		1.24 (0.49-4.50)	1.59 ± 1.06	989 (91.7)	1.52 (0.71–3.87)	1.97 ± 1.04
	. ,	1.00 (0.45-3.93)	1.44 ± 1.06	1,225 (93.8)	1.50 (0.75-3.67)	1.88 ± 1.04
		0.94 (0.40-2.89)	1.30 ± 1.10	464 (96.9)	1.87 (0.93-4.11)	2.15 ± 1.06
	2.03 ^{NS}	6.21*	1.74 ^{NS}	14.68***	7.46*	1.72 ^{NS}
High %body fat ^{§§}						
		1.15 (0.46-4.13)	1.50 ± 1.04	2,004 (92.9)	1.48 (0.71–3.68)	1.90 ± 1.03
,		0.90 (0.43-3.11)	1.38 ± 1.08	674 (95.5)	1.90 (0.98-4.21)	2.15 ± 1.06
	9.69**	3.61 ^{NS}	0.74 ^{NS}	5.77*	17.34***	3.92*
Abdominal obesity§§§	1 2 4 5 (64 6)	1 11 (0 16 1 00)	1 45 1 1 0 4	2 020 (02 ()	1 40 (0.71 0.70)	100 100
		1.11 (0.46-4.09)	1.47 ± 1.04	2,020 (92.6)	1.49 (0.71–3.70)	1.89 ± 1.03
		1.00 (0.43-3.34) 0.75 ^{NS}	1.46 ± 1.08 0.01^{NS}	658 (96.6) 14.06***	1.90 (0.94–4.27) 15.92***	2.19 ± 1.05 6.10*
Test 5	0.05"	0.75	0.01	14.00	15.92	6.10"
Women (N = 3,167)						
BMI categories §						
S	1,179 (65.1)	1.36 (0.55-4.02)	1.57 ± 1.05	1,629 (89.9)	1.23 (0.60-3.23)	1.66 ± 1.04
		1.20 (0.47-3.74)	1.54 ± 1.06	840 (93.1)	1.30 (0.69-2.80)	1.65 ± 1.05
Obese (n = 453) 2	277 (61.2)	1.07 (0.49-2.93)	1.5 ± 1.09	437 (96.5)	2.03 (1.03-3.91)	2.34 ± 1.07
Test 2	2.82 ^{NS}	2.88 ^{NS}	0.11 ^{NS}	23.79***	43.37***	11.55***
High %body fat ^{§§}						
No $(n = 2,320)$	1,512 (65.2)	1.35 (0.53-4.09)	1.59 ± 1.04	2,093 (90.2)	1.25 (0.61-3.22)	1.68 ± 1.03
Yes (n = 847) 5	513 (60.6)	1.10 (0.47-3.17)	1.44 ± 1.07	813 (96.0)	1.60 (0.88-3.25)	1.93 ± 1.05
	5.71*	6.06*	1.59 ^{NS}	27.32***	18.55***	5.10*
Abdominal obesity ^{§§§}						
		1.37 (0.54-4.08)	1.61 ± 1.04	1,978 (90.2)	1.22 (0.60-3.04)	1.63 ± 1.03
	. ,	1.00 (0.47-3.21)	1.43 ± 1.06	928 (95.3)	1.70 (0.87–3.54)	2.03 ± 1.05
Test 3	3.34 ^{NS}	6.63**	2.64 ^{NS}	23.03***	34.99***	15.34***

 1 Results are expressed as number of subjects (percentage) with values above the lower limit of detection (LOD), median and interquartile range (IQR) for values above LOD and exponentiated multivariate-adjusted average \pm standard error for values above LOD. Statistical analysis by a chi-square, b Kruskall-Wallis test and c general linear model on log-transformed values adjusting for age (continuous), smoking (3 groups), and leisure-time physical activity (2 groups): NS not significant; *p < 0.05; **p < 0.01; ***p < 0.001.

IL-1 β levels between patients with anorexia nervosa and normal-weight subjects have been found [21]. In this study, negative associations between IL-1 β levels and obesity markers were found, with %body fat being the only obesity marker associated with IL-1 β levels on multivariate analysis in women with a similar borderline significant (p = 0.053) trend in men. The lack of significance among men might be related to a lower statistical power due to a small sample size; indeed, the multivariate-adjusted estimate of the slope in men was comparable to that in women (table 4). These findings are in agreement with the previously reported effect of IL-1 in mice, i.e., lower adipocyte numbers and reduced dietary intake,



[§]Normal, BMI < 25 kg/m²; overweight, $25 \le BMI < 30 \text{ kg/m}^2$; obese, BMI $\ge 30 \text{ kg/m}^2$.

^{§§}Defined as a %body fat ≥28% in men and ≥40% in women.

^{§§§}Defined as a waist > 102 cm in men and > 88 cm in women.



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Table 3. TNF- α and CRP distribution according to different markers of obesity, by gender¹

	TNF-α			CRP		
	values > LOD (N) ^a	median (IQR) ^b	multi- adjusted ^c	values > LOD (N ⁾ a	median (IQR) ^b	multi- adjusted ^c
Men (N = 2,863)						
BMI categories [§] Normal (n = 1,078) Overweight (n = 1,306) Obese (n = 479) Test	1,069 (99.2) 1,298 (99.4) 477 (99.6) 0.97 ^{NS}	2.87 (1.86-4.37) 3.10 (1.90-4.63) 3.46 (2.13-5.48) 19.32***	3.04 ± 1.03 3.12 ± 1.02 3.37 ± 1.04 2.45^{NS}	1,078 (100) 1,306 (100) 479 (100) NA	0.8 (0.4–1.6) 1.4 (0.7–2.7) 2.2 (1.3–4.3) 343.3***	0.93 ± 1.03 1.47 ± 1.03 2.24 ± 1.05 140.6***
High %body fat ^{§§} No (n = 2,157) Yes (n = 706) Test Abdominal obesity ^{§§§}	2,144 (99.4) 700 (99.2) 0.49 ^{NS}	2.94 (1.87–4.46) 3.40 (2.09–5.48) 15.48***	3.10 ± 1.02 3.21 ± 1.03 0.80 ^{NS}	2,157 (100) 700 (100) NA	1.0 (0.5–2.1) 2.0 (1.1–4.3) 224.2***	1.17 ± 1.02 1.98 ± 1.04 126.2***
No (n = 2,182) Yes (n = 681) Test	2166 (99.3) 678 (99.6) 0.59	2.93 (1.86-4.46) 3.50 (2.12-5.67) 26.48***	3.04 ± 1.02 3.41 ± 1.03 8.66**	2182 (100) 681 (100) NA	1.0 (0.5-2.0) 2.2 (1.2-4.3) 261.3***	1.15 ± 1.02 2.12 ± 1.04 184.5***
Women (N = 3,167)						
BMI categories [§] Normal (n = 1,812) Overweight (n = 902) Obese (n = 453) Test High %body fat ^{§§}	1,797 (99.2) 897 (99.5) 449 (99.1) 0.71 ^{NS}	2.59 (1.64-4.08) 2.90 (1.77-4.62) 3.29 (2.12-5.08) 39.49***	2.78 ± 1.02 2.96 ± 1.03 3.43 ± 1.04 9.91***	1,812 (100) 902 (100) 453 (100) NA	0.8 (0.5–1.8) 1.9 (0.9–3.6) 3.8 (1.9–7.0) 647.8***	0.95 ± 1.03 1.88 ± 1.03 3.53 ± 1.05 $337.5***$
No (n = 2,320) Yes (n = 847) Test Abdominal obesity ^{§§§}	2,300 (99.1) 843 (99.5) 1.25 ^{NS}	2.59 (1.63-4.16) 3.20 (2.11-4.92) 48.92***	2.81 ± 1.02 3.22 ± 1.03 12.12***	2,320 (100) 847 (100) NA	1.0 (0.5-2.2) 2.6 (1.4-5.1) 396.1***	1.13 ± 1.02 2.55 ± 1.04 315.9***
No (n = 2,193) Yes (n = 974) Test	2,177 (99.3) 966 (99.2) 0.08 ^{NS}	2.57 (1.66-4.14) 3.20 (2.04-5.01) 48.11***	2.79 ± 1.02 3.22 ± 1.03 16.27***	2,193 (100) 974 (100) NA	1.0 (0.5-2.1) 2.6 (1.4-5.2) 462.6***	1.08 ± 1.02 2.51 ± 1.04 410.0***

 1 Results are expressed as number of subjects (percentage) with values above the lower limit of detection (LOD), median and interquartile range (IQR) for values above LOD and exponentiated multivariate-adjusted average \pm standard error for values above LOD. Statistical analysis by a chi-square, b Kruskall-Wallis test and c general linear model on log-transformed values adjusting for age (continuous), smoking (3 groups), and leisure-time physical activity (2 groups): NS not significant; *p < 0.01; **p < 0.001.

although we have no information on the former. Overall, our results suggest that increased IL-1 β levels are associated with lower %body fat levels, probably due to a reduced adipocyte differentiation.

It has been estimated that approximately 30% of circulating IL-6 comes from adipose tissue [22]. In this study, obese subjects had higher IL-6 levels, a finding also reported by others [23]. Interestingly, after multivariate adjustment, abdominal obesity was significantly associated with IL-6 levels in both genders, a finding which is in agreement with recent literature [7, 24]. Still, although all obesity markers were significantly and positively correlated with IL-6 levels in both genders, these associations were no longer significant



[§]Normal, BMI < 25 kg/m²; overweight, 25 ≤ BMI < 30 kg/m²; obese, BMI ≥ 30 kg/m².

^{§§}Defined as a %body fat ≥28% in men and ≥40% in women.

^{§§§} Defined as a waist > 102 cm in men and > 88 cm in women.



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Table 4. Relationships between inflammatory and obesity markers, by gender¹

	IL-1β	IL-6	TNF-α	hs-CRP		
	ть-тр	IL-U	TIVIŒ	IIS-GIVI		
Men (N)	1,732	2,678	2,844	2,862		
Spearman correlation						
BMI	-0.057^{NS}	0.053**	0.095***	0.359***		
%Body fat	-0.079**	0.072***	0.114***	0.365***		
Waist circumference	e -0.063**	0.074***	0.122***	0.381***		
Linear regression ^a						
BMI	-0.011 (0.009) ^{NS}	0.015 (0.006)*	0.012 (0.004)**	0.082 (0.005)***		
%Body fat	-0.015 (0.007)*	$0.005(0.005)^{NS}$	0.007 (0.003)*	0.055 (0.004)***		
Waist circumference	e –0.003 (0.003) ^{NS}	0.007 (0.002)**	0.006 (0.001)***	0.031 (0.002)***		
Ridge regression ^b						
BMI	-0.002 (0.020) ^{NS}	$-0.001 (0.014)^{NS}$	-0.004 (0.009) ^{NS}	0.025 (0.010)*		
%Body fat	$-0.019 (0.010)^{NS}$	$-0.007(0.007)^{NS}$	$-0.001 (0.004)^{NS}$	0.019 (0.005)***		
Waist	0.004 (0.007) ^{NS}	$0.009 (0.005)^{NS}$	0.008 (0.003)*	0.016 (0.004)***		
Women (N)	2,025	2,906	3,143	3,167		
Spearman correlation						
BMI	-0.048^{NS}	0.095***	0.122***	0.475***		
%Body fat	-0.082***	0.080***	0.122***	0.433***		
Waist circumference	e –0.055 ^{NS}	0.109***	0.141***	0.462***		
Linear regression ^a						
BMI	-0.005 (0.007)*	0.026 (0.005)***	0.019 (0.003)***	0.106 (0.004)***		
%Body fat	-0.010 (0.004) ^{NS}	0.010 (0.003)**	0.006 (0.002)**	0.058 (0.003)***		
Waist circumference	e -0.004 (0.003) ^{NS}	0.011 (0.002)***	0.007 (0.001)***	0.040 (0.002)***		
Ridge regression ^b						
BMI	0.029 (0.015) ^{NS}	$0.020 (0.012)^{NS}$	0.021 (0.007)**	0.055 (0.008)***		
%Body fat	-0.017 (0.006)**	$-0.007 (0.005)^{NS}$	$-0.007(0.003)^*$	0.017 (0.004)***		
Waist circumference		0.007 (0.004) ^{ŃS}	0.003 (0.003) ^{ŃS}	0.015 (0.003)***		

 $^1\text{Only}$ for participants with values above lower levels of detection. All analyses were performed on log-transformed variables, adjusting for age, smoking status and leisure-time physical activity. For linear and ridge regression, results are presented as slope (standard error): $^{\text{NS}}$ not significant; *p < 0.05; **p < 0.01; ***p < 0.001.

^aFor each obesity marker separately. ^bAll obesity markers taken together.

after multivariate ridge regression. Our results are partly in agreement with one study [25] which found no significant correlation between abdominal fat distributions measured by ultrasonography and circulating IL-6 levels. Conversely, the lack of association between %body fat and IL-6 levels is not in agreement with a previous study [7]. A possible explanation for this discrepancy is that the previous study included only obese subjects, thus limiting the generalization of the findings. Overall, our results suggest that abdominal obesity is associated with increased IL-6 levels in both genders, but that this increase in IL-6 levels is not mediated by any specific obesity marker (BMI, waist circumference, or %body fat). Nevertheless, further studies are needed to better assess the specific effects of selected obesity markers on IL-6 levels.

TNF- α is mainly secreted by macrophages after stimulation by bacterial lipopolysaccharides and stimulates the production of other cytokines such as IL-6 [24]. Adipocytes have also been shown to produce TNF- α [26], and high TNF- α levels have been found in obese subjects [27]. In both genders, subjects with abdominal obesity had higher TNF- α levels, and these differences persisted after multivariate adjustment; conversely, increased BMI and %body fat were associated with increased TNF- α levels in women only. Only BMI





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(in women) and waist circumference were independently and positively associated with TNF- α levels, a result which is partly in agreement with some smaller studies [23] but not with others [24, 25]. Again, no independent association between %body fat and TNF- α levels was found, a finding in contradiction with a previous study [7]. In women, BMI, but not waist circumference, was associated with TNF- α , a finding which is in agreement with another study [25]. Finally, obese subjects have a higher prevalence of diabetes, a condition which increases inflammatory markers. Still, further adjustment on diabetic status (defined as fasting plasma glucose \geq 7 mmol/l and/or presence of antidiabetic treatment) did not change the results (supplementary tables 1 and 2). Overall, our results suggest that both BMI and waist circumference might be associated with an increased production of TNF- α , while the effect of %body fat awaits further investigation.

CRP is an acute-phase reactant mainly synthesized in the liver. Several studies have shown that hs-CRP is associated with most obesity markers [7]. Indeed, all obesity markers were significantly and positively associated with hs-CRP levels, but likely due to differing metabolic pathways. For instance, the high fat mass (overall or in the abdominal region) might lead to increased production of TNF- α and IL-6, which in turn increase hepatic production of hs-CRP [28], while the relationship with BMI could be simply due to increased arthritis or hyperuricemia [29].

The major strength of our study is that we used a large, population-based sample representative of the Swiss population, that a precise characterization of the participants was performed, and that we used a statistical method correcting for correlations amongst independent variables. However, we should also mention some limitations of our study. Although the participation rate was similar to other epidemiological studies [30], it was rather low (41%), which might limit the generalization of findings. It has also been shown that African Americans have lower visceral adipose tissue but similar or higher concentrations of inflammatory biomarkers than Caucasians [31]; as only data from Caucasian subjects was analyzed, our findings may not apply to other ethnicities. Further, all measures of association (Spearman correlations, linear or ridge regression coefficients) were relatively low for all inflammatory markers, except for hs-CRP, suggesting that obesity markers exert only a modest impact on IL-1 β , IL-6, and TNF- α levels.

In summary, our results suggest that obesity markers are significantly, moderately and differentially associated with cytokine levels. %Body fat is negatively associated with IL-1 β levels; abdominal obesity is associated with increased IL-6 levels in both genders, but that this increase in IL-6 levels is not mediated by any specific obesity marker; both BMI and waist circumference are associated with an increased production of TNF- α , while the effect of %body fat awaits further investigation. Conversely, all obesity markers are strongly associated with hs-CRP levels.

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Disclosure Statement

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