

Brain-Derived Neurotrophic Factor and Extracellular Vesicle-Derived miRNAs in an Italian Cohort of Individuals With Obesity: A Key to Explain the Link Between Depression and Atherothrombosis

Patrizia Amadio¹, Chiara Macchi², Chiara Favero³, Marta Zarà¹, Giulia Solazzo³, Laura Dioni³, Leonardo Sandrini¹, Luisella Vigna⁴, Maria Francesca Greco², Massimiliano Buoli^{5,6}, Cesare R. Sirtori², Angela Cecilia Pesatori^{3,4}, Alessandro Ieraci⁷, Massimiliano Ruscica^{2*}, Silvia Stella Barbieri^{1*†} and Valentina Bollati^{3†}

¹ Brain-Heart Axis: Cellular and Molecular Mechanisms Unit, Centro Cardiologico Monzino Istituti di Ricovero e Cura a Carattere Scientifico (IRCCS), Milan, Italy, ² Department of Biomolecular and Pharmacological Sciences, University of Milan, Milan, Italy, ³ EPIGET LAB, Department of Clinical Sciences and Community Health, University of Milan, Milan, Italy, ⁴ Occupational Health Unit, Fondazione Istituti di Ricovero e Cura a Carattere Scientifico (IRCCS) Ca' Granda Ospedale Maggiore Policlinico, Milan, Italy, ⁵ Department of Neurosciences and Mental Health, Fondazione Istituti di Ricovero e Cura a Carattere Scientifico (IRCCS) Ca'Granda Ospedale, Maggiore Policlinico, Milan, Italy, ⁶ Department of Pathophysiology and Transplantation, University of Milan, Milan, Italy, ⁷ Department of Pharmaceutical Sciences, University of Milan, Milan, Italy,

Background: Obesity and depression are intertwined diseases often associated with an increased risk of cardiovascular (CV) complications. Brain-Derived Neurotrophic Factor (BDNF), altered in the brain both of subjects with depression and obesity, provides a potential link between depression and thrombosis. Since the relationship among peripheral BDNF, depression and obesity is not well-defined, the aim of the present report has been to address this issue taking advantage of the contribution played by extracellular vesicle (EV)-derived miRNAs.

Research Process: Associations among circulating BDNF, depression and EVderived miRNAs related to atherothrombosis have been evaluated in a large Italian cohort of obese individuals (n = 743), characterized by the Beck Depression Inventory (BDI-II) score.

Results: BDI-II was negatively associated with BDNF levels without a significant impact of the rs6265 BDNF polymorphism; this association was modified by raised levels of IFN- γ . BDNF levels were linked to an increase of 80 EV-derived miRNAs and a decrease of 59 miRNAs related to atherosclerosis and thrombosis. Network analysis identified at least 18 genes targeted by these miRNAs, 7 of which involved in depression and CV risk. The observation of a possible link among BDNF, depression, and miRNAs related to atherothrombosis and depression in obesity is novel and may lead to a wider use of BDNF as a CV risk biomarker in this specific subject group.

Keywords: BDNF, obesity, depression, atherothrombosis, EV-miRNA, cardiovascular disease

OPEN ACCESS

Edited by:

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Reviewed by:

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*Correspondence:

Massimiliano Ruscica massimiliano.ruscica@unimi.it Silvia Stella Barbieri silvia.barbieri@ccfm.it

[†]These authors have contributed equally to this work and share last authorship

Specialty section:

This article was submitted to General Cardiovascular Medicine, a section of the journal Frontiers in Cardiovascular Medicine

> Received: 28 March 2022 Accepted: 16 June 2022 Published: 13 July 2022

Citation:

Amadio P, Macchi C, Favero C, Zarà M, Solazzo G, Dioni L, Sandrini L, Vigna L, Greco MF, Buoli M, Sirtori CR, Pesatori AC, Ieraci A, Ruscica M, Barbieri SS and Bollati V (2022) Brain-Derived Neurotrophic Factor and Extracellular Vesicle-Derived miRNAs in an Italian Cohort of Individuals With Obesity: A Key to Explain the Link Between Depression and Atherothrombosis. Front. Cardiovasc. Med. 9:906483. doi: 10.3389/fcvm.2022.906483

INTRODUCTION

Obesity prevalence has increased exponentially over the last few decades reaching epidemic proportions. Between 2017 and 2018, 42.4% of adult Americans were found to have a body mass index (BMI) exceeding 30 (1). Obesity is a complex condition, with serious social and psychological facets. It may affect virtually all age and socioeconomic groups and it may threaten to overwhelm the health systems of both developed and developing countries. Contrary to conventional wisdom, the obesity epidemic is not restricted to industrialized societies; in developing countries, it is estimated that over 115 million people suffer from obesity-related health problems (2).

Although in men generally there is a higher rate of overweight, women have instead higher rates of obesity, linked to numerous health consequences, including diabetes and diseases of the cardiovascular (CV) system, such as hypertension, stroke, atherosclerosis, and thrombosis (3). In addition, systematic reviews and meta-analyses have confirmed the relationship between obesity and depression, both indicating that obesity raises the risk of depression, and that depression predisposes to obesity (4). This reciprocal link establishes a potential vicious circle that can lead to or aggravate the development of metabolic disorders. In particular, it is well-known that depression increases markedly the risk to develop CV complications in obese patients (5). Understanding the biochemical bases of obesity/depression, is thus crucial in order to identify new potential biomarkers of CV diseases (CVDs).

In this complex area of research, the investigations on circulating extracellular vesicles (EVs)-derived miRNAs have achieved a great interest. In particular, changes in circulating EV-derived miRNAs have been described in atherosclerosis, thrombosis and obesity, suggesting that EVs might play a role in the crosstalk between metabolic homeostasis and CV complications (6, 7).

Genome-wide association studies (GWAS) have identified 24 potential pleiotropic genes, overlapping between mood disorders and cardiometabolic diseases (8). Among these genes, brain derived neurotrophic factor (BDNF) has been recently found in both obesity and depression (9). BDNF is a neurotrophin playing a pivotal role in several physiological and pathological conditions, including neuroplasticity, energy homeostasis, and cardiovascular function (10). It is involved in the regulation of neurogenesis and neuroprotection, food intake, atherosclerotic plaque variability and morphology, and it has been related to inflammation, fibrin clotting stability, platelet and monocyte activation, and adipocyte maturation (11–14). Altogether, the hypothesis that altered levels of this neurotrophin may predispose to a raised CV risk is supported by many Authors (15).

Polymorphisms in the BDNF gene, among these the rs6265, have been associated with depression (16) and acute coronary syndromes (17, 18); low circulating BDNF levels have been reported in coronary artery disease (12, 15, 19) and in depressed patients (20). In contrast, the relationship between peripheral BDNF and obesity is not well-defined (21). Indeed, owing to small sample sizes of clinical studies (<100 individuals), bias in

subjects' recruitment, procedures of sampling/storage and BDNF measurement, as well as the lack of adjustment for confounders (e.g., platelet number, physical exercise) it is difficult to interpret results in literature (22).

Aim of the present study was, therefore, to investigate the association among BDNF, depression and EV-derived miRNAs related to atherothrombosis in a cohort of 743 obese individuals. In addition, the impact of the rs6265 BDNF polymorphism (23) on these associations has been explored.

MATERIALS AND METHODS

Study Design and Patient Population

Seventy hundred and forty-three individuals of the crosssectional Susceptibility to Particle Health Effects, miRNAs and Exosomes (SPHERE) study were randomly selected (24). These individuals were recruited at the Center for Obesity and Work-Activity (Fondazione IRCCS Ca' Granda Ospedale Maggiore Policlinico in Milan, Lombardy, Italy). The eligibility criteria of the SPHERE study were: (a) older than 18 years at enrolment; (b) overweight/obese according to body mass index (BMI): overweight, BMI between 25 and 30 kg/m²; obese: BMI of 30 kg/m² or more; (c) resident in the Lombardy Region at the time of recruitment. Exclusion criteria were: diagnosis of cancer, heart diseases, stroke, other chronic diseases in the last 5 years. The study conformed to the Declaration of Helsinki and each participant provided written informed consent approved by the Ethics Committee of Fondazione IRCCS Cà Granda Ospedale Maggiore Policlinico (approval number 1425).

Samples Collection

All patients underwent fasting blood sampling around 9 a.m. Peripheral venous blood samples were collected into vacutainer tubes containing EDTA (ethylenediaminetetraacetic acid) disodium salt (9.3 mM; Vacutainer System, Becton Dickinson, Franklin Lakes, NJ, US.); they were centrifuged at 1,200 g for 15 min. Plasma and buffy coat fractions was transferred to separate cryovials, aliquoted and immediately frozen at -80° C until use. Only one aliquot of 1.5 ml of fresh plasma was centrifuged to prepare EV pellets at 1,000, 2,000, and 3,000 g for 15 min at 4°C. The pellet was discarded to remove cell debris.

Laboratory Measurements and Beck Depression Inventory II (BDI-II) Evaluation

Demographic, clinical and biochemical data were obtained in all subjects as previously described (25). All participants were evaluated according to the Beck Depression Inventory II (BDI-II), considered as an appropriate tool to evaluate depressive symptoms in subjects with medical comorbidities such as obesity (26). The following scores correspond to the different severity of depressive symptoms: minimal changes = 0-13, mild depression = 14-19, moderate depression = 20-28, and severe depression = 29-63.

BDNF and PCSK9 Evaluation

Plasma BDNF and PCSK9 concentrations were measured by commercial ELISA kits (Biosensis, South Australia and R&D Systems, MN, respectively) as previously described (12, 25).

BDNF Polymorphism

DNA was extracted from each buffy coat by the Wizard Genomic DNA purification kit (Promega, Madison, WI, USA), according to the manufacturer's instructions. BDNF polymorphism rs6265 (C/T) was selected for the current study, and analyzed using TaqMan® SNP Genotyping Assay (C_11582758_10, Applied Biosystems, San Diego, CA, USA). In a volume of 7 μ l, containing 15 ng gDNA sample (3 μ l), 0.35 μ l TaqMan® Assay (20X), 3.5 μ l TaqPathTM ProAmpTM Master Mix and 0.15 μ l Nuclease-Free Water, the Polymerase Chain Reaction (PCR) was carried out as follows: pre-read 60°C for 30 s, initial denaturation 95°C for 5 min, followed by 40 cycles of denaturing at 95°C for 15 s and annealing/extension at 60°C for 60 s; a final step post-read 60°C for 30 s.The reactions and analysis were conducted on Quant Studio 12KFlex Real-Time PCR System (Applied Biosystems, San Diego, CA, USA).

Extracellular Vesicles-miRNAs Isolation and Analysis

To prepare the EV pellets for miRNA extraction, fresh plasma deprived of cell debris was transferred to a 10.4-ml polycarbonate ultracentrifuge tube (Beckman Coulter), filled with PBS. Plasma was ultracentrifuged (BeckmanCoulter Optima-MAX-XP) at 110,000 g for 75 min at 4°C and decanted. The EV pellet was stored at -80°C until use. Isolation of miRNAs from plasma extracellular vesicles (EVs) was performed with the combination of miRNeasy kit and RNeasy Cleanup Kit (Qiagen), according to the manufacturer's protocol. MiRNAs were eluted in 20 µL of Nuclease-Free Water and stored at -80° C, until use. MiRNAs reverse transcription (RT) and preamplification reactions, followed by real-time RT-PCR analysis with the QuantStudioTM 12K Flex OpenArray® Platform (Applied Biosystems), were previously described (27). Gene Expression Suite Software (Applied Biosystems) was used to process miRNA expression data from the "TaqManTM OpenArrayTM Human MicroRNA panel" (ThermoFisher) analysis. To elucidate the possible role of the EV-miRNAs, we performed a miRNA target analysis using SpidermiR by R software (v 4.0.4). Then, we compared genes obtained by the miRNA target analysis, with genes associated with thrombosis, atherosclerosis, atherothrombosis, cardiovascular disease, and depression downloaded from DisGeNET v 7.0. Finally, genes targeted by identified miRNAs and associated with atherothrombosis and depression were used to create a network on STRING v 11.5.

Statistical Analysis

Descriptive statistics were performed on all variables. Continuous data were expressed as the mean \pm standard deviation (SD) or as the median and interquartile range (Q1–Q3), as appropriate. Categorical data were presented as frequencies and percentages. We applied univariate and

multivariable negative binomial regression models for overdispersed count observations to evaluate the relationship between BDI-II score and BDNF levels. We tested the presence of over-dispersion based upon the Lagrange Multiplier (LM) test. Continuous variables were tested for normality and linearity. Multivariable analyses were adjusted for variables significantly related to the outcome in univariate analysis (p < 0.05). Given the existence of multicollinearity among predictor variables (e.g., HOMA-IR, QUICKI) the variance inflation factor (VIF) statistic was calculated. To determine the best-performing model, we ran several regression equations separately, that included one or more significant explanatory variables used to predict BDI-II score. For each model, we separately estimated the respective β coefficient and standard error (SE) for each variable in the model, p-value, VIF statistics, as well as goodness of fit of the model (R2). Finally, the best model selected to predict the association between BDI-II score and BDNF levels was adjusted for gender, BMI, education, occupation, lifestyle, BDNF rs6265, use of antidepressant drugs, PCSK9, MAP, glycated hemoglobin, HOMA-IR, use of diabetes medications, fibrinogen, and platelets. Estimated effects were described as a percent variation associated with an increase of 1 pg/mL in BDNF concentration. The percent variation was defined as [incidence rate ratio (IRR)-1]*100, IRR correspond to $exp(\beta)$.

To examine the potential effect modification by IFN- γ , we added the interaction term BDNF*IFN- γ to the multivariable selected model. We evaluated whether the effect of BDNF on depression score differs depending on IFN- γ concentrations. Linear regression models were used to evaluate the associations between BDNF levels and demographics and clinical characteristics of participants.

Multivariable linear regression models were applied to verify the association between miRNA expression and BDNF levels. miRNA expression values were log2-transformed to achieve a normal distribution. The regression models were adjusted for age, gender, BMI, smoking habits. Due to the high number of comparisons, we applied multiple comparison correction methods based on the Benjamini-Hochberg False Discovery Rate (FDR) to calculate the FDR *P*-value. A volcano plot of Δ % vs. – log10 *P*-values was used to display results. All statistical analyses were performed with SAS software (version 9.4; SAS Institute Inc., Cary, North Carolina, USA). A two-sided *p*-value of 0.05 was considered as statistically significant.

RESULTS

Study Population

As shown in **Table 1**, the majority of the 743 individuals (aged 50.8 ± 13.5) were females (73.6 vs. 26.4%). Mean BMI was 33.3 ± 5.4 kg/m² and the waist circumference (WC) was 100.7 ± 13.2 cm. Those not exercising physically were 60.2%, compared to 37.7% physically active. Blood pressure was in the normal range, 36.6% of participants being on hypotensive medications. The mean values of total cholesterol (TC), low-density lipoprotein cholesterol (LDL-C) and non-high-density lipoprotein cholesterol (non-HDL-C) were in the upper range of normal (TC 212.6 \pm 39.8 mg/dL, LDL-C 133.4 \pm 35.3 mg/dL, and

TABLE 1 | Demographic and clinical characteristics of the study participants (N = 743).

Characteristics	Value
Age, years	50.8 ± 13.5
Males	196 (26.4%)
BMI, Kg/m ²	33.3 ± 5.4
WC, cm	100.7 ± 13.2
TC, mg/dl	212.6 ± 39.8
HDL-C, mg/dL	58.8 ± 14.9
LDL-C. ma/dL	133.4 ± 35.3
non-HDL, ma/dL	153.8 ± 40.1
LDL/HDL	2.4 ± 0.9
TC/HDI	3.8 ± 1.2
TG mg/dl	102 [75:144]
CBP mg/l	0.3 [0.1.0.6]
Liric acid ma/dl	5.14 ± 1.35
Fibringgen mg/dl	333 5 + 59 1
Serum creatining, mg/dl	0.8 ± 0.3
AST 11/1	10[16:24]
	19 [10,24]
	22 [10;30]
	10 [13;29]
	11.2 ± 4.7
TSH, U/mL	1.8 [1.2;2.4]
Glucose, mg/dL	91 [86;100]
HbA1C, mmoL/moL	38.7 [36;42]
Insulin, U/mL	12.4 [8.5,17.8]
HOMA-IR score	2.8 [1.9;4.2]
QUICKI score	0.14 ± 0.01
PCSK9, ng/mL	276.3 ± 102.6
BDNF pg/mL	9.4 ± 5.6
Blood count	
White blood cells ($\times 10^3/\mu$ L)	6.8 ± 1.7
Red blood cells ($\times 10^6/\mu$ L)	4.8 ± 0.5
Platelets (×10 ³ /µL)	249.3 ± 59.4
Mean Corpuscolar Volume (fL)	85.0 ± 6.4
Hemoglobin (g/dL)	13.7 ± 1.3
Hematocrit (%)	40.5 ± 3.3
Smoking status	
Never smoker	358 (48.2%)
Former smoker	261 (35.1%)
Current smoker	124 (16.7%)
Occupation	447 (60.2%)
Employee	
Unemployed	63 (8.5%)
Pensioner	164 (22.0%)
Housewife	61 (8.2%)
Missing	8 (1.1%)
Education	
Primary school or less	60 (8.1%)
Secondary school	204 (27.5%)
High school	289 (38.9%)
University or more	171 (23.0%)
Missing	19 (2.5%)
	(Continued)

TABLE 1 | Continued

Characteristics	Value
Lifestyle	
Sedentary	447 (60.2%)
Physical activity	280 (37.7%)
Missing	16 (2.1%)
Heart rate, bpm	67.3 ± 10.2
Blood pressure, mmHg	
Systolic	124.9 ± 15.8
Diastolic	77.7 ± 9.9
MAP	93.4 ± 10.9
Diabetes	
Yes	85 (11.5%)
No	391 (52.6%)
Pre-diabetes	263 (35.4%)
Missing	4 (0.5%)
Antihypertensive medications	
Yes	272 (36.6%)
No	462 (62.2%)
Missing	9 (1.2%)
Statin medications	
Yes	75 (10.1%)
No	668 (89.9%)
Diabetes medications	
Yes	56 (7.5%)
No	687 (92.5%)
Antidepressant medications	
Yes	95 (12.8%)
No	648 (87.2%)
Depressive symptoms	
BDI-II score, median [q1;q3]	9 [5;14]
BDI-II score severity category, n (%)	
Minimal (0–13)	537 (72.27%)
Mild mood disturbance (14–19)	104 (14.00%)
Borderline clinical depression (20-28)	84 (11.31%)
Severe depression (>28)	18 (2.42%)

ALT, alanine aminotransferase; AST, Aspartate aminotransferase; BDI-II, Beck Depression Inventory II; BDNF, brain-derived neurotrophic factor; BMI, body mass index; CRP, C-reactive proteins; GGT, gamma-glutamyltransferase; HbA1C, glycated hemoglobin; HDL-C, high-density lipoprotein cholesterol; HOMA-IR, omeostasis model assessment of insulin resistance; LDL-C, low-density lipoprotein cholesterol; MAP, mean arterial pressure; PCSK9, proprotein convertase subtilisin/kexin type 9; TC, total cholesterol; TG, triglyceride; TSH, thyroid stimulating hormone; WC, waist circumference; QUICKI, quantitative insulin sensitivity check index; Continuous variables are expressed as mean ± standard deviation (SD) or as median [first quartile-third quartile] if not normally distributed; discrete variables are expressed as counts (%).

non-HDL-C 153.8 \pm 40.1 mg/dL). Only 10.1% of subjects were on statin medications. HDL-C and triglyceride (TG) levels were also in the normal range: mean 58.8 \pm 14.9 mg/dL and median 102.2 (75.0, 144) mg/dL, respectively. Mean levels of PCSK9 were 276.3 \pm 102.6 ng/mL.

Complete blood count (CBC), white blood cell formula, and platelets were in the normal range. Mean thyroid-stimulating hormone (TSH) levels were normal: 1.8 (1.2, 2.4) μ U/ml



and no participants were on thyroid substitution/suppression therapies. 16.7% were current smokers and C-reactive protein (CRP) median levels were 0.3 (0.1, 0.6) mg/L. The profile of inflammatory cytokines, namely, IFN- γ (10.4 ± 5.4 pg/mL), IL-8 (9.7 ± 9.9 pg/mL), IL-10 (4.2 [3.1;5.6] pg/mL), IL-18 (276.5 [208.5;356] pg/mL), macrophage inflammatory protein-1 α (MIP-1 α) (87 [58.5;120] pg/mL), macrophage inflammatory protein-1 β (MIP-1 β) (186.5 [147;234] pg/mL), monocyte chemoattractant protein-1 (MCP-1) (174 [142;215.5] pg/mL), and TNF- α (5.7 [3.2;8.5] pg/mL) is reported in **Supplementary Table 1**.

Concerning the glycaemic control, median basal fasting plasma glucose (FPG) was 91.0 (86, 100) mg/dL, median insulin was 12.4 (8.5, 17.8) UI/ml and median glycated hemoglobin (HbA1C) 38.7 (29, 30) mmol/mol; 7.5% were on antidiabetic medications. There were no significant abnormalities in the standard laboratory tests including liver enzymes (AST, ALT, GGT).

Average BDNF plasma concentration was 9.4 \pm 5.6 pg/mL (**Figure 1** and **Table 1**). To dissect the impact of the BDNF Val66Met variant, the entire cohort was genotyped for the rs6265 BDNF SNP. Carriers of BDNF^{Val/Val} were 62.6%, of BDNF^{Val/Met} 32.2%, and of BDNF^{Met/Met} 5.2%. The presence of the Met allele did not affect circulating levels of BDNF. These were 10.5 \pm 7.2 (BDNF^{Met/Met}), 9.1 \pm 5.1 (BDNF^{Val/Met}), and 9.4 \pm 5.7 (BDNF^{Val/Val}) pg/mL (p = 0.3503).

The median BDI-II total score was 9 (5, 14), 1.5% of participants showing severe depression. BDI-II values were higher in women compared to men, namely, 11.9 ± 7.9 and 6.8 ± 5.8 , respectively (p < 0.0001). Less than 13% of the whole cohort was treated with antidepressant medications, namely, selective serotonin or norepinephrine/serotonin reuptake inhibitors. Antidepressant treatment was generally prescribed to subjects with moderate and severe disease, the therapeutic approach being quite homogeneously distributed among the different clusters of depression severity (**Table 1**).

Relation Between BDI-II Score and Selected Study Variables

In univariate negative binomial regression analysis (**Supplementary Table 2**), BMI, fibrinogen, glycated hemoglobin, HOMA-IR score, QUICKI-score, PCSK9, platelets, employment status, use of antidepressant medications, were positively associated with the BDI-II score. A negative association was found for education, lifestyle, blood pressure, pre-diabetes, use of antidepressant medications and diabetes medications (**Supplementary Table 2**).

In the multivariate regression analysis, other variables were positively associated with the severity of depression: the HOMA-IR ($\beta = 0.0293$, SE = 0.125, p = 0.019), PCSK9 ($\beta = 0.0007$, SE = 0.0003, p = 0.0159) and platelet count ($\beta = 0.0017$, SE = 0.0005, p = 0.0018) were positively associated with the severity of depression (BDI-II score) (**Table 2**). In the univariate

TABLE 2 | Negative binomial regression model evaluating the association between BDI-II score and BDNF adjusted for participant's characteristics.

		β	SE	95% CI		Р	
Intercept		-	1.9327	0.4062	1.1366	2.7288	<0.0001
BDNF	-	-0.0195	0.0056	-0.0304	-0.0087	0.0004	
BDNF VAL66	MET/MET	0.0537	0.1343	-0.2096	0.3169	0.6896	0.6078
	VAL/MET	0.0621	0.0646	-0.0645	0.1887	0.3363	
	VAL/VAL	REF					
PCSK9	-	0.0007	0.0003	0.0001	0.0012	0.0159	
Gender	F	0.4424	0.0791	0.2875	0.5974	<0.0001	
	М	REF					
BMI	-	0.0056	0.0062	-0.0066	0.0178	0.3688	
MAP	-	-0.0053	0.0028	-0.0108	0.0001	0.0557	
Education	Primary school or less	0.0478	0.1391	-0.2248	0.3203	0.7312	0.1926
	Secondary school	0.0033	0.0805	-0.1545	0.1611	0.9674	
	High school	-0.1275	0.0771	-0.2786	0.0237	0.0983	
	University or more	REF					
Occupation	Employee	0.0469	0.1145	-0.1775	0.2713	0.6822	0.4386
	Unemployed	0.2072	0.1455	-0.0780	0.4924	0.1544	
	Pensioner	0.0684	0.1225	-0.1717	0.3084	0.5769	
	Housewife	REF					
Lifestyle	Sedentary	0.0929	0.1247	-0.1515	0.3372	0.4563	0.2029
	Active	-0.0242	0.1264	-0.2720	0.2236	0.8482	
	Sporty	REF					
Diabetes medications	Yes	0.2928	0.1317	0.0347	0.5510	0.0262	
	No	REF					
Use of antidepressant medications	Yes	0.3400	0.0871	0.5107	0.1694	<0.0001	
	No	REF					
HOMAIR	-	0.0293	0.0125	0.0048	0.0539	0.0192	
Glycated hemoglobin	-	0.0014	0.0041	-0.0066	0.0095	0.7248	
Fibrinogen	-	-0.0001	0.0005	-0.0012	0.0009	0.8201	
Platelets	-	0.0017	0.0005	0.0006	0.0027	0.0018	
Dispersion		0.3441	0.0285	0.2926	0.4048		

BDNF, brain-derived neurotrophic factor; BMI, body mass index; HOMA-IR, omeostasis model assessment of insulin resistance; MAP, mean arterial pressure; PCSK9, proprotein convertase subtilisin/kexin type 9. Values in bold represent significant changes.

analysis (**Supplementary Table 2**), the severity of depression (BDI-II score) did not significantly associate with plasma BDNF levels ($\beta = -0.009$, SE = 0.005, p = 0.087), whereas this association became significant in multivariate analysis. For every one-unit increase of BDI-II, BDNF levels decreased by 1.93% ($\beta = -0.0195$, SE = 0.0056, p = 0.0004) (**Table 3**). No relationship between BDI-II score and BDNFVal66Met polymorphism was found (**Supplementary Table 2**).

Similarly, the association between BDI-II score and the use of antidepressants and of diabetes medications were robust even after adjusting for the relevant covariates (**Table 2**).

Role of Inflammatory Markers in the Association Between Plasma BDNF Levels and BDI-II Score

Since inflammation has been linked to depression (28), the impact of the inflammatory environment on the previous associations was evaluated. Among the different tested cytokines (**Supplementary Table 1**), IFN-γ significantly modified the association between BDNF and BDI-II (interaction term *p*-value = 0.011). The strength of the association was modified in different IFN-γ strata. Each unit rise in BDNF led to an incidence rate ration (IRR) of 0.963 (95% CI 0.948–0.978), of 0.973 (95% CI 0.962–0.984), of 0.986 (95% CI 0.975–0.0997), and of 0.999 (95% CI 0.983–1.014), respectively, at IFN-γ levels of 0.65, 5, 10.4 and 15.8 pg/mL (**Figure 2**). No significant association between the different tested cytokines and BDI-II score was found (**Supplementary Table 2**).

Association Between BDNF and CV Risk Factors

BDNF levels were positively associated with CV parameters: TC ($\beta = 0.016$, SE 0.005, p = 0.0020), LDL-C ($\beta = 0.014$, SE = 0.006, p = 0.0134), non-HDL-C ($\beta = 0.015$, SE = 0.005, p = 0.0026), and TG ($\beta = 0.007$, SE = 0.005, p = 0.0077). In addition, a positive association between BDNF and platelets ($\beta = 0.032$, SE = 0.003, p < 0.0001), white ($\beta = 0.289$, SE = 0.123, p = 0.0184),

TABLE 3 | Slope coefficients from univariate linear regression models to evaluate association between BDNF and demographics and clinical characteristics of participants.

Disk polymorphion 0.3488 0.3488 0.3288 0.4248 WiMet 0.425 0.444 -1.038 0.477 0.4269 Monifer 1.045 0.886 -0.728 2.877 0.8289 Age: Wards 0.405 0.016 -0.027 0.623 0.6411 Gender 0.128 -0.024 0.038 -0.118 0.021 0.2588 Fornales 0.719 0.477 -0.190 1.834 - 0.2588 To, mg/dL 0.016 0.005 0.006 0.026 0.0290 HDL-C, mg/dL 0.015 0.005 0.006 0.028 0.014 LOL-Phot/C 0.370 0.135 0.008 0.732 0.0449 TO, mg/dL 0.017 0.023 0.012 0.017 0.025 TO, mg/dL 0.017 0.023 0.012 0.012 0.012 TO, mg/dL 0.017 0.023 0.012 0.012 0.012 TO, mg/dL 0.018 0.	Independent variable	β	SE	95% CI		Р	
NahlaPEFNai Mark-0.3280.449-1.0352.4070.4249Agu, Yuan-0.070.05-0.070.230.411Gender0.230.411Famasin0.7190.467-0.1961.834MilesPEFFamasin0.0140.038-0.1180.0310.2487Disk, graft0.0030.014-0.0250.0300.026Disk, graft0.0030.014-0.0250.0300.026Disk, graft0.0070.0360.0260.0020Disk, graft0.0070.0360.0260.0020Disk, graft0.0070.0560.0020.0120.0077Chiller, Graft0.0070.0560.0020.0120.0077Chiller, Graft0.0070.0560.0020.0120.0077Chiller, Graft0.0070.0560.0020.0120.026Disk, graft0.0110.0250.0120.0260.4100Seum creatione, mydt0.0110.0250.0110.0260.4100Seum creatione, mydt0.0110.008-0.0110.0280.014Chiller, Graft0.0110.008-0.0140.0280.014Chiller, Graft0.0110.0240.0240.0240.024Chiller, Graft0.0110.008-0.0140.0280.014Chiller, Graft0.0110.0	BDNF polymorphism						0.3488
NaMAR-0.3280.446-1.2030.5470.4641MavAR-1.0450.303-0.7882.6770.6411Gender0.151-0.1680.671MasBREFMayAR0.0060.0060.0260.026FG,majla0.0160.0060.0260.026TG,mayCl0.0160.0060.0260.026TG,mayCl0.0160.0060.0260.026LD-0,mg/L0.0150.0060.0260.026LD-1,mg/L0.0160.0060.0260.026LD-1,mg/L0.0170.028-0.0320.044LD-1,mg/L0.0170.0280.0280.021LD-1,mg/L0.0170.0280.0260.027CR,mg/L0.0070.0080.0280.028CR,mg/L0.0070.0080.0280.021CR,mg/L0.0070.0080.0280.021CR,mg/L0.0020.014-0.0230.732CR,mg/L0.0200.014-0.0230.021Serum castinke.mg/L0.0240.021-0.0220.021CR, Mu/L0.0240.024-0.0230.0710.3182Homogenitine.umal/L0.044-0.0230.0710.3182Homogenitine.umal/L0.044-0.0230.0710.3182Homogenitine.umal/L0.0240.0240.0230.026CR, Mu/L0.0270.0160.0240.024	Val/Val	REF	-	-	-	-	
NetNet 1.045 0.835 -0.783 2.877 0.283 Age. Nears -0.007 0.015 -0.037 0.023 0.024 Gender - - - - - Marka REF - - - - BM, kgm ² -0.043 0.034 -0.118 0.031 0.2988 TCS, mg/dL 0.016 0.005 0.006 0.026 0.0021 HEL-C, mg/dL 0.014 -0.005 0.030 0.028 0.0032 DLC, Grig/dL 0.015 0.005 0.006 0.732 0.644 DLC-CHUL-C 0.337 0.285 -0.032 0.474 0.081 ULC-CHUL-C 0.337 0.686 -0.292 0.644 0.5814 ULC-CHUL-C 0.307 0.686 -0.292 0.646 0.5814 ULC-CHUL-C 0.307 0.686 -0.292 0.646 0.5814 ULC-CHUL-C 0.307 0.605 0.605 0.657	Val/Met	-0.328	0.446	-1.203	0.547	0.4624	
App. News-0.0070.016-0.0370.0280.6411Gender0.1234MalesREFBolt, Lagrand0.0160.0060.0260.026DEC. mg/dL0.0160.0060.0260.026DEC. mg/dL0.0160.0060.0260.026DEC. mg/dL0.0170.0050.0060.026DEC. mg/dL0.0170.0260.0260.0134non-HOL-C, mg/dL0.0170.0260.0260.026DETC, mg/dL0.0170.0260.0260.026TO/HOL-C0.3700.4850.0060.0260.026TO/HOL-C0.3700.4850.0080.0260.026CHP, mg/L0.0070.0500.0260.0270.026CHP, mg/L0.0070.0500.0260.0260.026CHP, mg/L0.0070.0500.0260.0260.026Serum castinue. mg/dL-0.0180.054-0.0380.026Serum castinue. mg/dL0.0180.0260.0110.0260.026CHT, Lu0.0020.021-0.0220.1490.246Serum castinue. mg/dL0.0110.0260.0210.0280.028CHT, Lu0.0240.0240.0250.0260.026CHT, Lu0.0240.0250.0260.0260.026Serum castinue. mg/dL0.0270.0280.0260.026DEC, mg/dL0.029 </td <td>Met/Met</td> <td>1.045</td> <td>0.935</td> <td>-0.788</td> <td>2.877</td> <td>0.2639</td> <td></td>	Met/Met	1.045	0.935	-0.788	2.877	0.2639	
Örndari 0.1234 Fermales 0.716 0.467 -0.163 0.021 Males REF - - - BML (s/m) ² -0.043 0.083 -0.118 0.021 0.2588 DLC, mg/dL 0.003 0.062 0.003 0.062 0.014 DLC, mg/dL 0.014 0.005 0.003 0.026 0.0014 DLC, mg/dL 0.017 0.283 -0.053 0.026 0.0021 DLC, Mg/dL 0.017 0.283 -0.053 0.026 0.0021 DLC, Mg/dL 0.007 0.036 0.002 0.012 0.0051 0.0051 CMR MQL 0.007 0.003 0.002 0.012 0.002 0.012 0.002 0.012 0.0051 0.0281 Seumonentine.mg/dL 0.018 0.028 0.017 0.028 0.0281 0.0181 Seumonentine.mg/dL 0.019 0.029 0.011 0.028 0.021 0.0281 0.0281 S	Age. Years	-0.007	0.015	-0.037	0.023	0.6411	
Personale 0.479 0.467 -0.96 1.834 Mailes PEF - - - SetM.lsgm ² -0.018 0.031 0.2586 DEC.mg/dL 0.005 0.006 0.026 0.0202 LDL-C.mg/dL 0.016 0.005 0.026 0.026 LDL-C.mg/dL 0.017 0.005 0.005 0.026 0.0026 LDL-C.mg/dL 0.017 0.005 0.006 0.023 0.026 LD-C.mg/dL 0.017 0.237 0.210 0.012 0.012 CR,mg/dL 0.007 0.008 0.006 0.732 0.0490 CR,mg/dL 0.0171 0.288 0832 0.210 0.511 Diendgen mg/dL 0.002 0.144 -0.393 0.210 0.385 Seru mathine.mg/dL -0.018 0.044 -0.221 0.386 0.130 Marce mg/dL 0.011 0.008 0.001 0.028 0.231 GAT, ML 0.029 <th0< td=""><td>Gender</td><td></td><td></td><td></td><td></td><td>0.1234</td><td></td></th0<>	Gender					0.1234	
Mets PEF - - - BML kg/m ² -0.043 0.038 -0.116 0.032 0.026 BUL c, mg/dL 0.063 0.026 0.032 0.042 DL-C, mg/dL 0.015 0.005 0.026 0.026 DL-C, mg/dL 0.015 0.005 0.026 0.026 DL-C, mg/dL 0.017 0.230 -0.653 0.848 0.835 TC/HD-C 0.377 0.165 0.006 0.722 0.049 TC/HD-C 0.377 0.028 0.028 0.251 0.051 TC/HD-C 0.377 0.028 0.021 0.251 0.051 TC/HD-C 0.370 0.281 -0.393 0.251 0.051 Sum oreatino.mg/dL -0.002 0.024 -0.028 0.251 0.251 Sum oreatino.mg/dL -0.012 0.024 -0.028 0.271 0.205 0.231 Guezos.mg/dL 0.024 0.024 -0.022 0.316 0.336 0.49	Females	0.719	0.467	-0.196	1.634		
BM, kg/m² -0.043 0.038 -0.118 0.031 0.2268 TC, mg/dL 0.016 0.005 0.006 0.026 0.0020 LDL-C, mg/dL 0.014 0.006 0.003 0.026 0.0026 LDL-C, mg/dL 0.014 0.006 0.003 0.026 0.0026 LDL-C, mg/dL 0.017 0.230 -0.083 0.848 0.0838 LDL-C, mg/dL 0.007 0.008 0.002 0.012 0.004 CR, mg/dL 0.007 0.008 0.002 0.012 0.012 CR, mg/dL 0.002 0.014 -0.032 0.265 0.012 Romogen, mg/dL -0.032 0.024 -0.032 0.025 0.017 Seture 0.002 0.012 -0.022 0.026 0.027 Alt, UL- 0.026 0.014 -0.022 0.026 0.028 GGT, VL 0.027 0.137 -0.222 0.316 0.738 Mindocysteren, µmo/L 0.047 0.142	Males	REF	-	-	-		
To, mg/dL 0.096 0.006 0.006 0.007 0.007 HDL-C, mg/dL 0.003 0.014 -0.025 0.003 0.014 non-HDL-C, mg/dL 0.015 0.006 0.008 0.028 0.0088 LDL-C, Hold-C 0.370 0.185 0.006 0.732 0.0449 To, mg/dL 0.007 0.003 0.002 0.015 0.067 To, mg/dL 0.017 0.288 -0.032 0.5514 0.038 0.449 Uia add, mg/dL -0.092 0.154 -0.039 0.051 0.5514 Uia add, mg/dL -0.092 0.054 -2.201 0.368 0.4420 Suun metaline.mg/dL -0.018 0.026 -0.077 0.20 0.571 Suun metaline.mg/dL 0.019 0.002 0.010 0.338 0.3796 Strut/L 0.028 0.021 -0.022 0.316 0.3394 Suun metaline.mg/dL 0.024 -0.023 0.711 0.3192 Strut/L <	BMI, kg/m ²	-0.043	0.038	-0.118	0.031	0.2568	
Höl-C, mg/dL 0.003 0.014 -0.025 0.039 0.8472 LD-C, mg/dL 0.014 0.006 0.003 0.026 0.0134 LD-C, mg/dL 0.037 0.230 -0.053 0.848 0.0838 TC/HD-C 0.370 0.230 -0.053 0.848 0.0838 TC, mg/dL 0.077 0.003 0.002 0.012 0.0077 TC, mg/dL -0.002 0.014 -0.333 0.210 0.5514 Holmogen, mg/dL -0.002 0.012 -0.032 0.265 0.1608 Serum reatinine, mg/dL -0.012 0.002 0.025 0.8726 0.8726 ALT, U/L 0.002 0.012 -0.022 0.026 0.8726 GGT, U/L 0.002 0.011 0.038 0.0396 0.410 0.420 Homosysteine, jum/dL 0.644 -0.032 0.140 0.3724 0.3724 GGT, U/L 0.002 0.001 -0.032 0.140 0.3794 GMarcosyst	TC, mg/dL	0.016	0.005	0.006	0.026	0.0020	
LDI-C, mg/dL 0.014 0.006 0.003 0.028 0.0134 Inon-HD-C, mg/dL 0.015 0.005 0.006 0.028 0.0283 TC/HD-C 0.370 0.168 0.002 0.0448 0.0484 TG, mg/dL 0.007 0.003 0.002 0.012 0.0671 TG, mg/dL -0.012 0.014 -0.038 0.151 0.067 TG, mg/dL -0.012 0.0164 -0.038 0.210 0.551 TBrindgen, mg/dL -0.012 0.006 0.006 0.068 0.4880 Strum creatines, mg/dL -0.018 0.654 -2.201 0.365 0.168 AT, U/L 0.019 0.009 0.001 0.028 0.0276 0.225 Homosyntaine, µm/dL 0.026 0.021 -0.032 0.140 0.245 Homosyntaine, µm/dL 0.026 0.021 -0.032 0.141 0.243 Homosyntaine, µm/dL 0.022 0.022 0.021 0.033 0.155	HDL-C, mg/dL	0.003	0.014	-0.025	0.030	0.8472	
nen-HBL-C, mg/dL 0.015 0.005 0.026 0.026 0.0026 LDL-C/HDL-C 0.397 0.230 -0.053 0.44 0.083 TC/, mg/dL 0.007 0.003 0.002 0.012 0.007 CRP, mg/L 0.171 0.288 -0.392 0.735 0.5514 Heine add, mg/dL -0.002 0.014 -0.038 0.205 0.486 Serum creatinine, mg/dL -0.028 0.025 0.205 0.486 Serum creatinine, mg/dL -0.028 0.025 0.020 0.281 0.486 Serum creatinine, mg/dL 0.024 0.024 -0.022 0.281 0.486 AT, U/L 0.020 0.012 -0.022 0.316 0.734 Monocysteine, µmo/L 0.054 0.041 -0.033 0.042 0.935 Honocysteine, µmoL 0.054 0.044 -0.032 0.140 0.271 0.1508 Honocysteine, µmoL 0.011 0.002 -0.033 0.042 0.9352	LDL-C, mg/dL	0.014	0.006	0.003	0.026	0.0134	
LBL-CHOLC 0.397 0.230 -0.053 0.848 0.0838 TC/HDL-C 0.370 0.165 0.006 0.732 0.0449 TC/HDL-C 0.377 0.033 0.002 0.012 0.007 CHP, myL 0.171 0.288 -0.392 0.735 0.5514 Une add, my/dL -0.092 0.04 -0.039 0.055 0.4220 Serum creatinine, my/dL -0.028 0.025 -0.077 0.020 0.2510 AT, UL -0.028 0.026 -0.077 0.020 0.2145 AT, UL -0.028 0.026 -0.071 0.308 0.3382 HOM Coysteine, µm/L 0.044 -0.022 0.140 0.2145 HOM Coysteine, µm/L 0.047 -0.133 0.071 0.3182 HOM Coysteine, µm/L 0.024 0.024 -0.023 0.071 0.308 Glocose, mg/dL 0.021 0.020 0.3305 - 0.011 0.3182 HOMARISCOSE 0.022 <t< td=""><td>non-HDL-C, mg/dL</td><td>0.015</td><td>0.005</td><td>0.005</td><td>0.026</td><td>0.0026</td><td></td></t<>	non-HDL-C, mg/dL	0.015	0.005	0.005	0.026	0.0026	
TC/HDL-C 0.370 0.165 0.008 0.722 0.0449 TG, mg/dL 0.007 0.003 0.002 0.735 0.5514 CRP, mg/L -0.022 0.164 -0.393 0.210 0.5512 Eloinogan, mg/dL -0.022 0.064 -0.009 0.025 0.4270 Serum creditine, mg/dL -0.028 0.025 -0.077 0.020 0.8572 GGT, U/L -0.028 0.025 -0.077 0.020 0.8726 AT, U/L -0.028 0.026 -0.077 0.020 0.8726 GGT, U/L 0.009 0.001 0.038 0.08726 GGT, U/L 0.019 0.024 -0.022 0.140 0.2145 GGT, U/L 0.047 0.137 -0.222 0.316 0.7384 Homospatine, µmo/L 0.041 0.002 0.041 0.3182 GGT, U/L 0.047 0.131 0.223 0.336 GGT, U/L 0.041 0.028 0.021 0.338	LDL-C/HDL-C	0.397	0.230	-0.053	0.848	0.0838	
TG, mg/dL 0.007 0.003 0.002 0.012 0.0077 CRP, mg/L 0.171 0.280 -0.392 0.735 0.5514 Hornogen, mg/dL -0.002 0.004 -0.009 0.005 0.4920 Florinogen, mg/dL -0.018 0.684 -2.201 0.265 0.671 AST, U/L 0.002 0.012 -0.022 0.265 0.8726 GGT, U/L 0.004 0.003 0.038 0.0396 Hornocysteins, µmo/L 0.004 0.024 0.022 0.140 0.244 TSH, U/mL 0.047 0.137 -0.222 0.318 0.7304 GHOROCYSTEINS, µmo/L 0.047 0.137 -0.222 0.318 0.7304 GHOROCYSTEINS, µmo/L 0.047 0.137 -0.222 0.318 0.7304 GHOROCYSTEINS, µmo/L 0.047 0.021 -0.033 0.071 0.3182 Invalin level, U/mL 0.002 0.021 -0.039 0.042 0.3305 U/UKSI score 0.116 0.488 -0.116 0.143 0.220 0.201 <t< td=""><td>TC/HDL-C</td><td>0.370</td><td>0.185</td><td>0.008</td><td>0.732</td><td>0.0449</td><td></td></t<>	TC/HDL-C	0.370	0.185	0.008	0.732	0.0449	
CHP, mg/L 0.171 0.288 -0.392 0.735 0.5514 Unc add, mg/dL -0.092 0.164 -0.393 0.210 0.5512 Entrinogen, mg/dL -0.0918 0.654 -2.201 0.365 0.1608 Sartum creatinine. mg/dL -0.028 0.025 -0.077 0.020 0.2510 AST, U/L -0.028 0.026 -0.077 0.020 0.2510 ALT, U/L -0.028 0.021 -0.022 0.8726 0.8726 GGT, U/L 0.009 0.001 0.038 0.3956 0.413 0.7304 Homosysteine, µmo/L 0.047 0.137 -0.022 0.140 0.2145 GGR, U/L 0.047 0.024 -0.023 0.071 0.3182 HoA1C, mmd/moL 0.011 0.008 -0.014 0.022 0.150 Insulin kwet, U/mL 0.002 0.021 -0.007 0.022 0.3305 Culicki score -19.416 15.842 -50.466 11.833 0.2203	TG, mg/dL	0.007	0.003	0.002	0.012	0.0077	
Life acid, mg/dL -0.092 0.154 -0.393 0.210 0.5512 Fichingen, mg/dL -0.002 0.004 -0.009 0.005 0.4920 AST, U/L -0.028 0.025 -0.077 0.020 0.2510 AST, U/L 0.002 0.012 -0.022 0.025 0.8726 GGT, U/L 0.009 0.001 0.038 0.0366 Homocysteine, µmo/L 0.054 0.024 -0.022 0.316 0.7304 Glucase, mg/dL 0.047 0.137 -0.222 0.316 0.7304 Glucase, mg/dL 0.047 0.137 -0.222 0.316 0.7304 Glucase, mg/dL 0.011 0.008 -0.002 0.021 0.338 HoMAIR score 0.016 0.088 -0.116 0.149 0.8080 OUCKI score -19.165 15.842 -50.466 11.833 0.2203 OUCKI score -9.039 0.022 -0.037 0.025 0.2233 Vihte blood cells 0.29	CRP, mg/L	0.171	0.288	-0.392	0.735	0.5514	
Fibringen -0.002 0.004 -0.009 0.005 0.4420 Serum creatinine. mg/dL -0.918 0.664 -2.201 0.365 0.1608 AST, U/L -0.028 0.026 -0.072 0.020 0.8726 AST, U/L 0.002 0.012 -0.022 0.025 0.8726 GGT, U/L 0.004 -0.032 0.140 0.2145 TSH, U/ML 0.047 0.137 -0.022 0.316 0.7304 Glucose, mg/dL 0.024 0.024 -0.039 0.042 0.3952 HOMAIR score 0.016 0.068 -0.118 0.149 0.8880 HOMAIR score 0.016 0.068 -0.118 0.149 0.8880 QUICK score -19.416 15.842 -50.466 11.633 0.2203 Biod court -0.022 0.033 0.025 0.338 0.004 Pistelets 0.032 0.049 0.530 0.0144 Red blood cells 0.956 0.453 0.	Uric acid, mg/dL	-0.092	0.154	-0.393	0.210	0.5512	
Serun creatione. mg/dL -0.918 0.664 -2.201 0.365 0.1608 AST, U/L -0.028 0.025 -0.077 0.020 0.2510 ALT, U/L 0.002 0.012 -0.022 0.025 0.8726 GGT, U/L 0.019 0.009 0.011 0.038 0.0396 Homocysteine, µmo/L 0.054 0.044 -0.032 0.140 0.2145 GGT, U/L 0.047 0.137 -0.222 0.316 0.7304 Glucose, mg/dL 0.024 0.023 0.071 0.3182 HbA1C, mm0//mol 0.011 0.008 -0.004 0.027 0.1506 Insulin level, U/mL 0.002 0.021 -0.039 0.042 0.9382 UICKI score -19.416 15.842 -50.466 11.833 0.2203 QUICKI score -19.416 15.842 -50.466 1.845 0.0349 VIEK blood cells 0.289 0.123 0.0049 0.530 0.0184 Petales 0.0	Fibrinogen, mg/dL	-0.002	0.004	-0.009	0.005	0.4920	
AST, U/L -0.028 0.025 -0.077 0.020 0.2510 ALT, U/L 0.002 0.012 -0.022 0.025 0.8726 GGT, U/L 0.019 0.009 0.001 0.038 0.0396 Homocysteine, µmol/L 0.054 0.024 -0.032 0.140 0.2145 TSH, U/mL 0.047 0.137 -0.222 0.316 0.7304 Glucose, mg/dL 0.024 0.024 -0.023 0.071 0.3182 HoA1C, mmoL/moL 0.001 0.008 -0.014 0.027 0.1506 Insulin level, U/mL 0.002 0.021 -0.039 0.042 0.9352 HOMAIR score 0.016 0.688 -0.116 0.149 0.8080 QUICKI score -19.416 15.842 -50.466 11.633 0.2203 Biood count -0.002 0.002 -0.007 0.002 0.305 0.0184 Plateits 0.332 0.049 0.530 0.0184 -0.019 0.032 0.293 Hemocysteiodar Volume -0.033 0.025 0.038	Serum creatinine. mg/dL	-0.918	0.654	-2.201	0.365	0.1608	
A.T. U/L 0.002 0.012 -0.022 0.025 0.8726 GGT, U/L 0.019 0.009 0.001 0.038 0.0396 Homcoysteine, µmol/L 0.054 0.044 -0.032 0.140 0.2145 SH, U/mL 0.047 0.137 -0.222 0.316 0.7304 Glucose, mg/dL 0.024 0.024 -0.023 0.071 0.3182 HDA1C, mmoL/moL 0.011 0.008 -0.004 0.027 0.1506 Insulin level, U/mL 0.002 0.021 -0.039 0.042 0.9352 HOMAIR score 0.016 0.068 -0.116 0.149 0.8080 QUICKI score -19.416 15.842 -50.466 11.633 0.2203 Boot count Wite blood cells 0.395 0.049 0.305 0.049 Pisteis 0.032 0.033 0.025 0.038 -0.019 Mean Corpuscolar Volume -0.039 0.032 -0.020 0.026 0.2293 Henoglubin 0.111 0.157 -0.197 0.419 0.4812	AST, U/L	-0.028	0.025	-0.077	0.020	0.2510	
G.T. U/L 0.019 0.009 0.001 0.038 0.0396 Homocysteine, μmol/L 0.054 0.044 -0.032 0.140 0.2145 TSH, U/mL 0.047 0.137 -0.222 0.316 0.7304 BLACC, mmoL/moL 0.024 0.024 -0.023 0.071 0.3182 HDACL, mmoL/moL 0.002 0.021 -0.039 0.042 0.9352 Insulin level, U/mL 0.002 0.021 -0.039 0.042 0.9352 UICKI score 0.916 0.586 -0.116 0.149 0.8080 OUICKI score -9.002 0.002 -0.007 0.002 0.335 Biod count - - - 0.002 0.335 0.049 0.530 0.0184 Red blood cells 0.289 0.123 0.049 0.530 0.0184 Red blood cells 0.289 0.032 -0.102 0.025 0.283 Mean Corpuscolar Volume -0.039 0.032 -0.197 0.419	ALT, U/L	0.002	0.012	-0.022	0.025	0.8726	
Honocysteline, μmol/L 0.054 0.044 -0.032 0.140 0.2145 T5H, U/mL 0.047 0.137 -0.222 0.316 0.7304 Glucese, mg/dL 0.024 -0.023 0.071 0.3182 Hb41C, mmol/MoL 0.002 0.021 -0.039 0.042 0.9352 Insulin level, U/mL 0.002 0.021 -0.039 0.042 0.9362 QUICKI score -19.416 15.842 -50.466 11.833 0.2203 QUICKI score -9.002 0.002 -0.007 0.002 0.3305 Biod court - - -0.007 0.002 0.3305 Biod cells 0.289 0.123 0.049 0.530 0.0144 Red blood cells 0.032 0.003 0.025 0.2293 0.129 Mean Corpuscolar Volume -0.039 0.032 -0.102 0.025 0.2293 Hemoglobin 0.111 0.157 -0.197 0.419 0.4812 Hemoglobin 0.	GGT, U/L	0.019	0.009	0.001	0.038	0.0396	
TSH, U'mL 0.047 0.137 -0.222 0.316 0.7304 Glucose, mg/dL 0.024 0.024 -0.023 0.071 0.3182 HbA1C, mmoL/moL 0.011 0.008 -0.004 0.027 0.1506 Insulin level, U/mL 0.002 0.021 -0.0039 0.042 0.9352 HOMIR score 0.016 0.068 -0.116 0.149 0.8080 QUICKI score -19.416 15.842 -50.466 11.633 0.2203 PCSK9, ng/mL -0.002 0.002 -0.007 0.002 0.3305 Biod count -0.002 0.002 -0.007 0.002 0.3305 Vhite blood cells 0.9956 0.453 0.068 1.845 0.0349 Platelets 0.032 0.003 0.025 0.038 -0.001 Mean Corpuscolar Volume -0.039 0.332 -0.102 0.026 0.2233 Hemoglobin 0.111 0.157 -0.197 0.419 0.4812 Hematorit 0.068 0.062 -0.054 0.191 0.2730 C	Homocysteine, µmol/L	0.054	0.044	-0.032	0.140	0.2145	
Glucose, mg/dL 0.024 0.024 -0.023 0.071 0.3182 HbA1C, mmoL/moL 0.011 0.008 -0.004 0.027 0.1506 Insuli nevel, U/mL 0.002 0.021 -0.039 0.042 0.9352 HOMAIR score 0.016 0.068 -0.116 0.149 0.8080 QUICKI score -19.416 15.842 -50.466 11.633 0.2203 PGSK9, ng/mL -0.002 0.02 -0.007 0.002 0.3305 Biod count - - - 0.032 0.049 0.530 0.0184 Red blood cells 0.956 0.453 0.068 1.845 0.039 Platelets 0.032 0.002 0.025 0.2293 Hemoglobin 0.111 0.157 -0.197 0.419 0.4812 Hematocrit 0.068 0.062 -0.054 0.191 0.2730 Cytoknes - - -0.019 0.008 0.4487 IL-10 -0	TSH, U/mL	0.047	0.137	-0.222	0.316	0.7304	
Hba1C, mmoL/moL 0.011 0.008 -0.004 0.027 0.1506 Insulin level, U/mL 0.002 0.021 -0.039 0.042 0.9352 HOMAIR score 0.016 0.068 -0.116 0.149 0.8080 QUICKI score -19.416 15.842 -50.466 11.633 0.2203 PGSK9, ng/mL -0.002 0.002 -0.007 0.002 0.3305 Biod coult - 0.022 0.002 0.305 - - - - - 0.032 0.032 0.032 0.033 - 0.032 0.022 0.225 0.2293 - - - - - - 0.032 0.032 0.033 - 0.031 0.032 0.041 0.032 0.041 0.041 </td <td>Glucose. ma/dL</td> <td>0.024</td> <td>0.024</td> <td>-0.023</td> <td>0.071</td> <td>0.3182</td> <td></td>	Glucose. ma/dL	0.024	0.024	-0.023	0.071	0.3182	
Insulin level, U/mL 0.002 0.021 -0.039 0.042 0.9352 HOMAIR score 0.016 0.668 -0.116 0.149 0.8080 QUICKI score -19.416 15.842 -50.466 11.633 0.2203 PCSK9, ng/mL -0.002 0.002 -0.007 0.002 0.305 Blood count White blood cells 0.956 0.453 0.068 1.845 0.0349 Platelets 0.032 0.003 0.025 0.2293 0.001 Mean Corpuscolar Volume -0.039 0.032 -0.102 0.025 0.2293 Hemoglobin 0.111 0.157 -0.197 0.419 0.4812 Hemoglobin 0.011 0.062 -0.034 0.019 0.2730 Cytokines II-19 -0.019 0.033 -0.084 0.046 0.5727 IL-18 -0.001 0.001 -0.003 0.002 0.0065 MIP-17 MIP-17 0.001 0.002 0.006 0.015	HbA1C, mmoL/moL	0.011	0.008	-0.004	0.027	0.1506	
HOMAIR score0.0160.068-0.1160.1490.8080QUICKI score-19.41615.842-50.46611.6330.2203PCSK9, ng/mL-0.0020.002-0.0070.0020.3305Blood count </td <td>Insulin level, U/mL</td> <td>0.002</td> <td>0.021</td> <td>-0.039</td> <td>0.042</td> <td>0.9352</td> <td></td>	Insulin level, U/mL	0.002	0.021	-0.039	0.042	0.9352	
QUICKI score -19.416 15.842 -50.466 11.633 0.2203 PCSK9, ng/mL -0.002 0.002 -0.007 0.002 0.3305 Biod count - - - - - - - - - - - 0.002 0.002 0.002 0.003 0.0184 Red blood cells 0.9956 0.453 0.068 1.845 0.0349 Platelets 0.032 0.003 0.025 0.028 -0.001 Mean Corpuscolar Volume -0.039 0.032 -0.102 0.025 0.2293 Hemoglobin 0.111 0.157 -0.197 0.419 0.4812 Hemoglobin 0.111 0.157 -0.019 0.406 0.5727 IL-8 0.038 0.021 -0.003 0.079 0.0677 IL-10 -0.005 0.007 -0.019 0.008 0.4487 IL-18 -0.001 0.002 0.006 0.009 0.6580	HOMAIR score	0.016	0.068	-0.116	0.149	0.8080	
PCSK9, ng/mL -0.002 0.002 -0.007 0.002 0.3305 Blood count White blood cells 0.289 0.123 0.049 0.530 0.0184 Red blood cells 0.956 0.433 0.068 1.845 0.0349 Plate blood cells 0.032 0.003 0.025 0.038 <0.001	QUICKI score	-19.416	15.842	-50.466	11.633	0.2203	
Blood count No. Construction No. Construction White blood cells 0.289 0.123 0.049 0.530 0.0184 Red blood cells 0.956 0.453 0.068 1.845 0.0349 Platelets 0.032 0.003 0.025 0.038 <0.001	PCSK9. ng/mL	-0.002	0.002	-0.007	0.002	0.3305	
White blood cells 0.289 0.123 0.049 0.530 0.0184 Red blood cells 0.956 0.453 0.066 1.845 0.0349 Platelets 0.032 0.003 0.025 0.038 <.0001 Mean Corpuscolar Volume -0.039 0.032 -0.102 0.025 0.2293 Hemoglobin 0.111 0.157 -0.197 0.419 0.4812 Hemoglobin 0.068 0.062 -0.054 0.191 0.2730 Cytokines U U -0.019 0.033 -0.084 0.046 0.5727 IL-8 0.038 0.021 -0.003 0.079 0.0677 IL-10 -0.005 0.007 -0.019 0.008 0.4487 IL-18 -0.001 0.001 -0.003 0.002 0.008 0.0172 MIP-1β 0.004 0.002 0.008 0.015 <0.001 NF-α 0.011 0.002 0.008 0.015 <0.001	Blood count						
Red blood cells 0.956 0.453 0.068 1.845 0.0349 Platelets 0.032 0.003 0.025 0.038 -0.001 Mean Corpuscolar Volume -0.039 0.032 -0.102 0.025 0.2293 Hemoglobin 0.111 0.157 -0.197 0.419 0.4812 Hematocrit 0.068 0.062 -0.054 0.191 0.2730 Cytokines U U -0.019 0.033 -0.084 0.046 0.5727 IL-8 0.038 0.021 -0.003 0.079 0.0677 IL-10 -0.005 0.007 -0.019 0.038 0.4487 IL-18 -0.001 0.001 -0.003 0.022 0.6056 MIP-1β 0.002 0.001 0.008 0.0172 MCP-1 0.011 0.002 0.008 0.015 -5.001 MCP-1 0.011 0.002 0.008 0.015 -5.0168 MCP-1 0.021 0.028 </td <td>White blood cells</td> <td>0.289</td> <td>0.123</td> <td>0.049</td> <td>0.530</td> <td>0.0184</td> <td></td>	White blood cells	0.289	0.123	0.049	0.530	0.0184	
Platelets 0.032 0.003 0.025 0.038 <0.001 Mean Corpuscolar Volume -0.039 0.032 -0.102 0.025 0.2293 Hemoglobin 0.111 0.157 -0.197 0.419 0.4812 Hematocrit 0.068 0.062 -0.054 0.191 0.2730 Cytokines - - - - 0.777 IFN-γ -0.019 0.033 -0.084 0.046 0.5727 IL-8 0.038 0.021 -0.003 0.079 0.0677 IL-10 -0.005 0.007 -0.019 0.008 0.4487 IL-18 -0.001 0.001 -0.003 0.002 0.6656 MIP-1β 0.004 0.002 0.008 0.0172 MCP-1 0.001 0.008 0.0172 Smoking status - - . . Former smoker -0.224 0.456 -1.118 0.670 0.6233 Never smoker <t< td=""><td>Red blood cells</td><td>0.956</td><td>0.453</td><td>0.068</td><td>1.845</td><td>0.0349</td><td></td></t<>	Red blood cells	0.956	0.453	0.068	1.845	0.0349	
Mean Corpuscolar Volume-0.0390.032-0.1020.0250.2293Hemoglobin0.1110.157-0.1970.4190.4812Hematocrit0.0680.062-0.0540.1910.2730Cytokines0.0190.033-0.0840.0460.5727IL-80.0380.021-0.0030.0790.0677IL-10-0.0050.007-0.0190.0080.4487IL-18-0.0010.001-0.0030.0020.6056MIP-1β0.0020.004-0.0060.0090.6580MCP-10.0110.0020.0080.015<0.001	Platelets	0.032	0.003	0.025	0.038	<0.0001	
Hemoglobin 0.111 0.157 -0.197 0.419 0.4812 Hematocrit 0.068 0.062 -0.054 0.191 0.2730 Cytokines - 0.019 0.033 - 0.084 0.046 0.5727 IL-8 0.038 0.021 -0.003 0.079 0.06677 - <td>Mean Corpuscolar Volume</td> <td>-0.039</td> <td>0.032</td> <td>-0.102</td> <td>0.025</td> <td>0.2293</td> <td></td>	Mean Corpuscolar Volume	-0.039	0.032	-0.102	0.025	0.2293	
Hematocit0.0680.062-0.0540.1910.2730CytokinesIFN-γ-0.0190.033-0.0840.0460.5727IL-80.0380.021-0.0030.0790.0677IL-10-0.0050.007-0.0190.0080.4487IL-18-0.0010.001-0.0060.0090.6580MIP-1α0.0020.004-0.0060.0090.6580MIP-1β0.0010.0020.0080.0172MCP-10.0010.0020.0080.0150.5316Smoking status0.111-0.0290.0150.5316Former smoker-0.2240.456-1.1180.6700.62330.1130Never smoker0.9250.584-0.2192.0700.1130	Hemoglobin	0.111	0.157	-0.197	0.419	0.4812	
Cytokines IFN- γ -0.019 0.033 -0.084 0.046 0.5727 IL-8 0.038 0.021 -0.003 0.079 0.0677 IL-10 -0.005 0.007 -0.019 0.008 0.4487 IL-18 -0.001 0.001 -0.003 0.002 0.6056 MIP-1 α 0.002 0.004 -0.006 0.009 0.6580 MIP-1 β 0.004 0.002 0.001 0.008 0.0172 MCP-1 0.011 0.002 0.008 0.015 <0.001 TNF- α -0.007 0.11 -0.029 0.15 0.5316 Smoking status turner	Hematocrit	0.068	0.062	-0.054	0.191	0.2730	
IFN-γ-0.0190.033-0.0840.0460.5727IL-80.0380.021-0.0030.0790.0677IL-10-0.0050.007-0.0190.0080.4487IL-18-0.0010.001-0.0030.0020.6056MIP-1α0.0020.004-0.0060.0090.6580MIP-1β0.0010.0020.0010.0080.0172MCP-10.0110.0020.0080.015<5316Smoking status-0.0070.011-0.0290.0150.5316Former smoker-0.2240.456-1.1180.6700.6233Never smoker0.9250.584-0.2192.0700.1130	Cytokines						
IL-80.0380.021-0.0030.0790.0677IL-10-0.0050.007-0.0190.0080.4487IL-18-0.0010.001-0.0030.0020.6056MIP-1 α 0.0020.004-0.0060.0090.6580MIP-1 β 0.0010.0020.0010.0080.0172MCP-10.0110.0020.0080.015<0.001	IFN-v	-0.019	0.033	-0.084	0.046	0.5727	
IL-10-0.0050.007-0.0190.0080.4487IL-18-0.0010.001-0.0030.0020.6056MIP-1α0.0020.004-0.0060.0090.6580MIP-1β0.0040.0020.0010.0080.0172MCP-10.0110.0020.0080.015<0.001TNF-α-0.0070.011-0.0290.0150.5316Smoking status0.456-1.1180.6700.6233Current smoker0.9250.584-0.2192.0700.1130Never smokerREF	IL-8	0.038	0.021	-0.003	0.079	0.0677	
IL-18 -0.001 0.001 -0.003 0.002 0.6056 MIP-1 α 0.002 0.004 -0.006 0.009 0.6580 MIP-1 β 0.004 0.002 0.001 0.008 0.0172 MCP-1 0.011 0.002 0.008 0.015 <0.0001 TNF- α -0.007 0.011 -0.029 0.015 0.5316 Smoking status0.1608Former smoker -0.224 0.456 -1.118 0.670 0.6233 Never smokerREF $ -$	IL-10	-0.005	0.007	-0.019	0.008	0.4487	
MIP-1α0.0020.004-0.0060.0090.6580MIP-1β0.0040.0020.0010.0080.0172MCP-10.0110.0020.0080.015<0.0001TNF-α-0.0070.011-0.0290.0150.5316Smoking status0.0070.011-0.0290.0150.6233Former smoker-0.2240.456-1.1180.6700.6233Current smoker0.9250.584-0.2192.0700.1130Never smokerREF	IL-18	-0.001	0.001	-0.003	0.002	0.6056	
MIP-1β 0.004 0.002 0.001 0.008 0.0172 MCP-1 0.011 0.002 0.008 0.015 <0.0001 TNF-α -0.007 0.011 -0.029 0.015 0.5316 Smoking status - - 0.1608 0.670 0.6233 Current smoker 0.925 0.584 -0.219 2.070 0.1130 Never smoker REF - - - -	MIP-1a	0.002	0.004	-0.006	0.009	0.6580	
MCP-1 0.011 0.002 0.008 0.015 <0.0001 TNF-α -0.007 0.011 -0.029 0.015 0.5316 Smoking status - - 0.456 -1.118 0.670 0.6233 Former smoker 0.925 0.584 -0.219 2.070 0.1130 Never smoker REF - - - -	MIP-16	0.004	0.002	0.001	0.008	0.0172	
TNF-α -0.007 0.011 -0.029 0.015 0.5316 Smoking status -0.224 0.456 -1.118 0.670 0.6233 Current smoker 0.925 0.584 -0.219 2.070 0.1130 Never smoker REF - - -	MCP-1	0.011	0.002	0.008	0.015	<0.0001	
Smoking status -0.224 0.456 -1.118 0.670 0.6233 Current smoker 0.925 0.584 -0.219 2.070 0.1130 Never smoker REF - - - -	TNF-α	-0.007	0.011	-0.029	0.015	0.5316	
Former smoker -0.224 0.456 -1.118 0.670 0.6233 Current smoker 0.925 0.584 -0.219 2.070 0.1130 Never smoker REF - - - -	Smoking status	5.001	0.011	0.020	0.010	0.0010	0 1608
Current smoker 0.925 0.584 -0.219 2.070 0.1130 Never smoker REF - - - - -	Former smoker	-0 224	0.456	-1.118	0.670	0.6233	21.000
Never smoker REF	Current smoker	0.925	0.584	-0.219	2 070	0 1130	
	Never smoker	REF	-	-	-		

(Continued)

TABLE 3 | Continued

Independent variable	β	SE	95% CI		Р		
Occupation						0.5542	
Unemployed	0.006	0.754	-1.471	1.483	0.9938		
Pensioner	-0.044	0.511	-1.046	0.958	0.9309		
Housewife	-1.091	0.764	-2.589	0.407	0.1535		
Employee	REF	-	-	-			
Education						0.8123	
Secondary school	0.578	0.821	-1.032	2.188	0.4816		
High school	0.526	0.794	-1.029	2.081	0.5075		
University or more	0.810	0.839	-0.835	2.455	0.3346		
Primary school or less	REF	-	-	-			
Lifestyle							
Active	0.047	0.456	-0.846	0.940	0.9173		
Sporty	-0.438	0.865	-2.132	1.257	0.6128	0.8627	
Sedentary	REF	-	-	-			
Heart rate. bpm	0.029	0.020	-0.011	0.068	0.1552		
Blood pressure. mmHg							
Systolic	-0.002	0.013	-0.028	0.023	0.8496		
Diastolic	0.007	0.021	-0.034	0.048	0.7274		
MAP	0.002	0.019	-0.035	0.040	0.9046		
Diabetes						0.1569	
Yes	0.498	0.670	-0.816	1.811	0.4577		
Pre-diabetes	0.854	0.447	-0.021	1.729	0.0558		
No	REF	-	-	-			
Antihypertensive medications					0.3	249	
Yes	-0.424	0.430	-1.267	0.420			
No	REF	-	-	-			
Antidepressant medications					0.79	968	
Yes	-0.159	0.617	-1.368	1.050			
No	REF	-	-	-			
Statin medications					0.34	478	
Yes	-0.642	0.684	-1.982	0.698			
No	REF	-	-	-			
Diabetes medications							
Yes	0.777	0.780	-0.752	2.305	0.3192		
No	REF	-	-	-			

ALT, alanine aminotransferase; AST, Aspartate aminotransferase; BDNF, brain-derived neurotrophic factor; BMI, body mass index; CRP, C-reactive proteins; GGT, gammaglutamyltransferase; HbA1C, glycated hemoglobin; HDL-C, high-density lipoprotein cholesterol; HOMA-IR, omeostasis model assessment of insulin resistance; IFN-γ, interferon gamma; IL, interleukin; MCP-1, monocyte chemoattractant protein-1; MIP-1α, macrophage inflammatory protein-1 alpha; MIP-1β, macrophage inflammatory protein-1 beta; LDL-C, low-density lipoprotein cholesterol; PCSK9, proprotein convertase subtilisin/kexin type 9; TC, total cholesterol; TG, triglyceride; TNF-α, tumor necrosis factor-α. Values in bold represent significant changes.

and red ($\beta = 0.956$, SE = 0.453, p = 0.0349) blood cells were found. Among the tested proinflammatory cytokines, MIP-1 β ($\beta = 0.004$, SE = 0.002, p = 0.0172) and MCP-1 ($\beta = 0.011$, SE = 0.002, p < 0.0001) were positively associated with plasma BDNF levels (**Table 3**).

EVs-Derived miRNA, Atherothrombosis, and Depression

To further dissect the possible involvement of BDNF in CV disease, we analyzed EV-derived miRNAs isolated from the same individuals. After data cleaning, 508 miRNAs were expressed in

at least one subject. In a model adjusted for age, gender, BMI, and smoking habit, BDNF levels were associated with a significant increase of 80 miRNAs (the up-regulated group) and a significant decrease of 59 miRNAs (the down-regulated group), after FDR adjustment for multiple comparisons (FDR p < 0.05; **Figure 3** and **Supplementary Table 3**).

By using the DisGeNET database, we identified the genes targeted by these 139 miRNAs specifically related to thrombosis, atherosclerosis, atherothrombosis and CV disease. Among the group of up-regulated miRNAs targeted genes (11,890), 1,091 were associated with CV disease,





log10(FDR P-value) 0 -15 -10 10 15 -20 -5 0 5 20 Δ% FIGURE 3 | Volcano plot reporting the association of BDNF levels with all measured miRNAs in extracellular vesicles. Red and green dots represent miRNAs, respectively, with negative and positive statistically significant association with BDNF. BDNF, Brain-Derived Neurotrophic Factor.

1,253 with atherosclerosis, 60 with thrombosis, and 69 with atherothrombosis (**Supplementary Figure 1**). In particular, they represented more than 50% of total genes included in the databases of atherothrombosis and more than 60% of those included in the databases of thrombosis, atherosclerosis and CV disease.

Among the group of down-regulated miRNAs targeted genes (11,121), 1,008 were associated with CV disease, 1,180 with atherosclerosis, 50 with thrombosis, and 63 with atherothrombosis (Supplementary Figure 2). In particular, they represented between 50 and 57% of total genes included in the four datasets. Among all the identified target genes, 18 were reported in all four (thrombosis, atherosclerosis, atherothrombosis, datasets and cardiovascular disease). Specifically, 15 target genes were associated with the group of up-regulated miRNAs and 13 target genes with the down-regulated group. The following 10 target genes CYP2C19, F3, IL6, MTHFR, PLAT, PTGS 2, SERPINE 1, TFP1, THBD, and TNF were shared between the up- and down-regulated groups (Supplementary Table 4).

In silico multiple-protein interaction analysis (STRING) revealed that all 18 target genes (AC2, PTGS2, CYP2C19, IL18, IL6, TNF, PTGS1, MTHFR, LPA, PLAT, SERPINE1, SELP, F3, VWF, THBD, CPB2, TFPI, and PROC) directly or indirectly interacted with BDNF (**Figure 4**). Overall, these data support the important relationship between circulating BDNF levels and increased CV risk in this population.

Finally, in order to dissect the potential relationship among BDNF, obesity, atherothrombosis and depression, we performed a further analysis by using specific datasets of the DisGeNET pertaining to depression, i.e., chronic depression, clinical depression, mild depression, recurrent depression and severe depression. We found that 51 out of 11,890 genes targeted by up-regulated miRNAs were associated with depression, of which 8 were reported also in the



atherothrombosis database (**Supplementary Table 5**). Regarding the genes targeted by down-regulated miRNAs, 52 out of 11,121 were related to depression, 7 of which were also found in the atherothrombosis dataset (**Supplementary Table 6**). Interestingly, BDNF emerged among the genes shared by depression and atherothrombosis, and it was well-connected to all the genes identified in both databases, namely, IL-1 β , IL-6, IL-18, TNF, APOE, ACE, MTR, MTHFR genes (**Figure 5**).

DISCUSSION

In the present study by using data from an Italian cohort of obese individuals, we identified a significant negative association between depression scores and BDNF plasma levels, a finding that was modified by elevated inflammatory levels of IFN- γ . In addition, we provided evidence that the amount of circulating BDNF can reflect the EV-derived miRNAs related to CV disease and depression, supporting the hypothesis that BDNF may be the link between mood disorders and cardiometabolic diseases. Depression development is linked to several factors among which obesity most likely has a causal role. Mendelian randomization studies have provided clear evidence that high BMI is causally linked to the likelihood to develop depression (31).

Among central nervous system (CNS) determinants of obesity, BDNF could exert a potentially significant role. The relationship of BDNF with obesity and depression has been the subject of many investigations which collectively described a



reduction of brain BDNF levels in both obesity and depression. Indeed, by contributing to the regulation of both synaptic plasticity and energy metabolism, including feeding behavior, BDNF has been recognized as a key target in the relationship between metabolic syndrome and psychiatric diseases. BDNF by activation of tyrosine kinase receptor-B (Trk-B) promotes neuronal survival, neuroplasticity, synaptogenesis, and survival and differentiation of serotonergic neurons through different signaling, such as PLC/DAG/IP3, PI3K/Akt, and Ras/MAPK-ERK pathways. BDNF can modulate the release of several neurotransmitters by interacting with other neuropeptides. Specifically, BDNF may regulate serotonin release (32), and modulates dopamine release via PLC-y/IP3 pathway (33). Additionally, it may facilitate glutamate release through the MAPK/SynI activation (34), and controls GABAergic synaptic transmission influencing neuronal excitability. Reduction of BDNF levels affects formation of synaptic spine density, and decreases excitatory neurons leading to depression (35). BDNF can also interact with melanocortin, leptin, corticotrophinreleasing hormone (CRH), and thyrotropin-releasing hormone (TRH) playing critical physiopathological roles. BDNF and TrkB play a significant role in the neural circuitry triggering food intake and body weight, regulating appetite through the modulation of melanocortin signaling and hypothalamic serotonergic mechanisms. Pre-clinical data have shown that central alterations of BDNF signaling in mice results in severe

hyperphagia and obesity (14, 36). On the other hand, high-fat diet promotes obesity, depressive-like phenotypes and reduction of BDNF in the hippocampus, a brain region important in controlling the mood. Moreover, hippocampal reductions of BDNF have been also reported in animal models of depression and in the postmortem brain samples of depressed patients (35). Interestingly, BDNF is potentially associated with obesity, depression and coronary syndromes (22). Being a versatile regulator of food intake as well as of inflammatory adipocyte maturation, it has been considered a critical mediator of obesity (14). In addition, compared to healthy subjects, reduced levels of BDNF have been found in patients with coronary syndromes (11, 12), and elevated BDNF appears to exert a protective effect (15). However, an open question remains as to whether BDNF has a fundamental role in CV complications, or it is just a bystander.

Waiting for the understanding of whether "came first the chicken or the egg," several mechanisms have been proposed, starting from the knowledge that BDNF plays a cardio-protective role thanks to its antioxidant, pro-survivor and pro-angiogenic properties. It promotes the activation of reactive oxygen species detoxifying enzymes, increases cardiomyocyte survival under hypoxic conditions, stimulates new vessel assembly, and affects the structure, polymerization and viscoelastic properties of the fibrin clot, thus influencing the anticoagulation process (12, 37), without changing circulating fibrinogen levels (12, 29). According to these data (12, 29), in the SPHERE population no association between BDNF and fibrinogen levels was found. However, the ability of BDNF to modify fibrin clot structure suggests its clinical relevance as a potential biomarker of thrombosis including acute coronary syndrome (ACS). Indeed, as emerged from the PLATO study, fibrin clot properties independently predict adverse clinical outcomes after ACS (38). Future studies in depressed obese subjects should be performed to dissect out the potential relationship between BDNF and fibrin clot properties.

Interestingly, in a human study, Wu et al. found a positive correlation between severity of coronary artery stenosis and depression scores, and reported that lower serum BDNF levels were closely related to the development of depression in patients with stable coronary artery disease (39). Since it is wellestablished that reduced BDNF levels in the brain are linked to obesity, BDNF may represent the bridge among obesity, depression, and CV diseases. However, whether obesity itself may influence circulating BDNF levels is not well-defined as yet (22). In the present study, plasma BDNF levels did not correlate with BMI, whereas an evident relationship with BDI-II emerged, overall suggesting the strong impact of BDNF on depression. The lack of follow-up of the SPHERE study did not allow to assess whether antidepressant medications were able to raise plasma BDNF levels. We did not find any interaction between the antidepressant medications and BDNF levels (p = 0.9013) as well as between antidepressant medications and BDNF rs6265 (p = 0.0603) on BDI-II score.

Alterations in GABAergic, catecholaminergic and serotonergic systems as well as an aberrant cytokine production related to dysregulation of both innate and adaptive immune systems have been reported in people suffering from depression

when compared to healthy individuals (40, 41). However, no significant correlations between catecholamine metabolites and BDNF in the blood of subjects with major depression have been reported (42), and platelet serotonin levels do not seem to go hand in hand with BDNF levels. Platelet serotonin levels do not differ between depressed and control subjects, whereas both reduced platelet BDNF content and plasma BDNF levels were found in subjects with depression (20, 30, 43). Then, it has been hypothesized that the altered BDNF levels observed in patients with depression are due to a reduced release from platelets which are considered one of the major circulation sources of BDNF. This latter is released by platelets upon their activation, raising to 100-fold higher levels in serum vs plasma (44) and correlating positively with platelets number (45). However, in people with depression, the reduction in serum and plasma BDNF levels seem to be independent of platelet reactivity (20), suggesting that other cells are involved in this modulation.

Recently, awareness has been raised regarding the relationship among obesity, depression and diet-related inflammation (46, 47). Gialluisi et al. (46) showed that the low-grade inflammation (INFLA)-score, based on four circulating inflammatory biomarkers (C-reactive protein, granulocyte-tolymphocyte ratio, platelet and white blood cell counts), could explain the association between depressive symptoms and the dietary inflammatory index.

Considering that the studied cohort predominantly comprises overweight individuals, our findings are two fold: depression is directly related to a higher number of platelets, as in normal weight people (48); and the positive association between platelet number and depression score was mitigated by BDNF (Supplementary Figure 3). Altogether, this suggests that other cell populations (49, 50) could actively affect circulating BDNF levels. On this matter, we found that BDNF was positively correlated with white and red blood cells. This correlation can be explained, at least in part, by the ability of peripheral blood mononuclear (PBMC) to produce/secret BDNF (51, 52) that in turn can promote cell proliferation and survival. Indeed, BDNF induces the maturation, activation, proliferation and survival of T and B lymphocytes (49, 50). Since TrkB receptor was found in several bone marrow cells including osteoblasts, eosinophils, mastocytes, megakaryocytes and erythroblasts (49), it is possible to hypothesize that BDNF can affect also the behavior of these cell types.

Accumulating evidence shows that chronic inflammation is associated with depressive symptoms, and pre-clinical evidence supports the critical involvement of IFN- γ in the pathogenesis of this disease (53). In the SPHERE population, we did not measure levels of catecholamines, acetylcholine, serotonin, or of other neurotransmitters, but several cytokines have been evaluated. Interestingly, we did not find evidence of an association between BDI-II score and the measured cytokines, although IFN- γ affected the association BDNF-depression. The evidence that modulation of the BDNF-depression association occurs primarily at low IFN- γ levels suggests that under elevated inflammatory conditions, as found, e.g., in severe obesity (54), the ability of BDNF to impact on depressive symptoms is potentially negligible. *In vitro* data showed that T cells co-secreted larger

BDNF and EV-miRNAs

amounts of BDNF and IFN- γ upon CD40 activation, whereas activated monocytes produced BDNF but not IFN- γ (55). Interestingly, obese patients displayed high levels of CD40 Ligand (56) suggesting that in this clinical condition activation of PBMC, rather than platelets, could be an important source of circulating BDNF.

The increased circulating levels of BDNF paralleled by the enhanced expression of the inflammatory markers MCP-1 and MIP1^β, might be a consequence of ongoing pathological events. Abundant BDNF is present in the atherosclerotic lesions localizing in macrophages, vascular smooth muscle cells, as well as in perivascular adipose tissue (13, 37, 57). It may be secreted by activated monocytes in response to inflammatory cytokines (58) to exert anti-inflammatory and anti-apoptotic effects. However, when BDNF is chronically expressed in the presence of severe inflammatory conditions, it could have deleterious effects. BDNF enhances oxidative stress and promotes tissue factor (TF) activity in monocyte/macrophages (13, 51) and down regulates TrkB (57). Reduction in TrkB expression can contribute to atherosclerosis progression by affecting endothelial and smooth muscle cells survival and function, as well as heart development (37). Interestingly, our group showed that despite circulating levels of BDNF being lower in CV patients compared to healthy controls, a positive relationship between plasma BDNF and coronary plaque vulnerability, also including macrophage infiltration, appears to be present (11).

The relationship between the BDNFVal66Met polymorphism and depression is currently under debate. Specifically, the systematic review and meta-analyses performed by Kishi et al. (59) did not found any association between this polymorphism and depression, while noting that plasma BDNF levels were reduced in depressed patient. On the contrary, a positive association between this polymorphism and depression only in men, but not in women, was found by Verhagen (60). Similarly, the genetic association studies of the BDNFVal66Met polymorphism in obesity and cardiometabolic diseases provided inconsistent results. BDNF polymorphism could not differentiate carriers of Val vs. Met based on BMI (61, 62), or with CNS BDNF levels in suicidal depressed patients (63). However, this polymorphism may increase the risk of developing obesity when associated with the other polymorphisms related to the metabolic pathway (62). By contrast, other reports showed both a positive and an inverse association between obesity and the Met allele (64, 65). Finally, a greater adiposity based on BMI percentiles or BMIfor-age z-scores has been also described in Met allele carriers (61). In agreement with these studies, in the SPHERE cohort we did not find differences in BDNF allele frequency in relation with BMI, depression and circulating BDNF levels. Specific additional studies, including also subjects with a normal weight, may be needed in order to provide more definite conclusions.

Finally, by analyzing the EV-miRNA profiles we found that 139 miRNAs, known to target genes involved in atherothrombotic and CV processes, were associated with BDNF levels. Even if evidence on the ability of BDNF to directly modulate miRNAs expression is scanty, BDNF alters the expression of miR-433, miR-19b-1, miR-432, and miR-181c in human endothelial progenitor cells (66), and upregulates miR-214 in mouse embryonic stem cells (67). These findings provide evidence that some of the miRNAs here identified may be affected by BDNF.

Interestingly, the 139 miRNAs here identified target 10 genes (CYP2C19, F3, IL6, MTHFR, PLAT, PTGS2, SERPINE1, TFPI, THBD, TNF) present in all datasets analyzed. Remarkably, the protein-protein interaction analyses showed that BDNF could represent a critical hub among all these genes.

In addition, when data were analyzed focusing on depression state, 9 genes (ACE, APOE, BDNF, IL18, IL1B, IL6, MTHFR, MTR, and TNF) were shared with the atherothrombotic dataset. All these 9 genes were closely interlinked, and BDNF was directly related to 6 of them. Altogether these observations support the hypothesis that BDNF may be a common link among obesity, CV diseases and depression.

Remarkably, ACE has been identified as the target gene for both atherothrombotic complications (68) and management of depression (69). Overexpression of ACE2 in a transgenic animal model attenuated plaque vulnerability and inflammation (70) and reduced the propensity to develop an anxiety phenotype (29). Recently, it has been demonstrated that down regulation of ACE2 by SARS-CoV-2 exacerbates the thrombotic COVID-19 severity and predisposes to depression (71). Changes in the levels of serum BDNF were observed in post-infected-COVID-19 symptomatic patients (72), predicting an intensive care admission and a worse prognosis (73).

In this scenario, Lorkiewicz et al. (74) proposed to include BDNF among biomarkers of COVID-19-derived depression, whereas De Sousa et al. (75) underlined how the rise of BDNF levels, induced by physical activity, was crucial for maintaining good mental health after COVID-19 syndrome.

Of note, alterations in the ACE signaling-pathway affected the release of modulators of fibrinolytic processes, such as Serpin1 and Tissue-type plasminogen activator (PLAT). Although Serpin1 and PLAT were not co-shared between the depression and atherothrombotic phenotypes, it has been hypothesized that PLAT and Serpin1, by modulating the proteolytic ability of plasmin, may regulate the cleavage of pro-BDNF precursor to the mature form (76). All these findings clearly summarize the strong connections of these genes in obesity, depression, and CV disease.

Our analysis also identified the possible involvement of prostaglandin-endoperoxide synthase 2 (PTGS2), a key enzyme in the production of prostanoids. Increased PTGS2 levels have been detected in atherosclerotic plaques (77) as well as in the rat brain showing depression-like phenotypes (78). The imbalance in PTGS2-BDNF signaling pathway has been suggested to be one of the pathogenetic mechanisms of depression (79).

Although P-selectin, thromboglobulin, and MTHFR, a key modulator of purine and thymidylate biosynthesis, play a critical role in thrombosis (80, 81), and their alterations have been observed in subjects with depression (82, 83), definitive potential relationship with BDNF is actually missing.

The present study presents some limitations. Firstly, this is a monocentric study, and the questionnaire was self-administered, without final psychiatric interview diagnosing the depressive

state. Moreover, the BDI-II questionnaire was administered only once at the recruitment of patients. BDI-II has been widely applied including earlier studies in cardiac patients (84); having been successfully administered both to adolescents and adults, as a tool of both medical assistance and research, it represents an established self-rating scale. A large portion of subjects classified with depression has subclinical or mild symptomatology, only a fraction of those taking psychotropic medications. Although we collected extensive information on habitual physical activity and education, many other factors may influence BDNF levels, and not all potential confounders could be taken into consideration here. The follow-up of these subjects is not available at present, limiting the potential prognostic value of BDNF for cardiovascular events at this moment. No other traditional markers of predisposition to thrombosis have been measured. Finally, the molecular mechanisms underlying the relation between BDNF and the emerged miRNAs affecting obesity, depression and CVD have not been investigated as yet.

CONCLUSIONS

The described link among BDNF and obesity, classical and non-classical CV risk factors in individuals with obesity and different degrees of depression provide one more explanatory variable (i.e., BDNF) as a culprit. In the present report, there was clear evidence of a negative association between BDNF and major clinical depression scores, with no impact of BDNF genetic polymorphisms. Although the association between BDNF and depression risk factors was lost in the presence of elevated levels of a major inflammatory marker (IFN- γ), the detection of this link is novel as was the association with miRNAs related to atherothrombosis and depression. This last analysis detected a large number of up-regulated and down-regulated genes, eventually allowing to create a network analysis of potential use in the molecular characterization of both CV and depression in obese individuals.

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DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding authors.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by Ethics Committee of Fondazione IRCCS Cà Granda OspedaleMaggiore Policlinico (approval 1425).

AUTHOR CONTRIBUTIONS

PA, CM, MZ, and MG performed the experiments. CF performed the statistical analyses. LD isolated the EV-derived miRNAs. GS performed the bioinformatic analyses of miRNAs. LV was responsible for the recruitment of SPHERE cohort. MB was responsible for the evaluation of BDI-II score. LS, AI, AP, and CS critically reviewed the manuscript. SB, MR, and VB conceived the study and wrote the first draft of the manuscript. All authors contributed to the article and approved the submitted version.

FUNDING

This work was partially supported by Fondazione Cariplo (2018-0511 to MR and 2018-0525 to SB), Banca di Credito Cooperativa di Milano (donation to MR), European Research Council (ERC-2011-StG 282,413 to VB), and Italian Ministry of Health, Rome, Italy (Ricerca Corrente RC 2021, and 5×1000 2018).

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fcvm. 2022.906483/full#supplementary-material

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