Clinical significance of hyperhomocysteinemia in infective endocarditis

A case-control study

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Abstract

Blood coagulation plays a key role in the pathogenesis of infective endocarditis (IE). Conditions associated with thrombophilia could enhance IE vegetation formation and promote embolic complications.

In this study, we assessed prevalence, correlates, and clinical consequences of hyper-homocysteinemia (h-Hcy) in IE.

Homocysteine (Hcy) plasma levels were studied in 246 IE patients and 258 valvular heart disease (VHD) patients, as well as in 106 healthy controls.

IE patients showed Hcy levels comparable to VHD patients (14.9 [3–81] vs 16 [5–50] μ mol/L, respectively; P=0.08). H-Hcy was observed in 48.8% of IE patients and 55.8% of VHD (P=0.13). Vegetation size and major embolic complications were not related to Hcy levels. IE patients with h-Hcy had a higher prevalence of chronic kidney disease and a higher 1-year mortality (19.6% vs 9.9% in those without h-Hcy; OR 2.21 [1.00–4.89], P=0.05). However, at logistic regression analysis, h-Hcy was not an independent predictor of 1-year mortality (OR 1.87 [95% CI 0.8–4.2]; P=0.13).

Our data suggest h-Hcy in IE is common, is related to a worse renal function, and may be a marker of cardiac dysfunction rather than infection. H-Hcy does not appear to favor IE vegetation formation or its symptomatic embolic complications.

Abbreviations: BD = blood donors, CI = confidence interval, CIED = cardiac implantable electronic device, <math>CIEDE = cardiac implantable electronic device endocarditis, <math>CRP = C-reactive protein, Hcy = homocysteine, h-Hcy = hyper-homocysteinemia, IE = infective endocarditis, INR = international normalized ratio, IQR = interquartile range, MTHFR = 5,10 methylene tetrahydrofolate reductase, NVE = native valve IE, OR = odds ratio, PVE = prosthetic valve IE, SD = mean \pm standard deviation, VHD = valvular heart disease.

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1. Introduction

Infective endocarditis (IE) remains a challenging clinical condition showing increasing incidence and an unchanged, high

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mortality rate.^[1] The endocardial vegetation is the diseasedefining lesion of IE, generating the common embolic complications of the disease.^[2]

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IE vegetation originates at the site of endocardial lining damage or after an intracardiac device has been implanted as a consequence of hemostasis system activation and subsequent bacterial seeding on an initially sterile clot.^[3,4] In light of the central role played by the hemostasis system in IE vegetation formation, we hypothesized that a prothrombotic state might increase the risk of IE onset or trigger the early IE vegetation development.^[4] Moreover, a thrombophilic condition could also promote the embolic complications of the disease. Recently, we observed a significant association between the G20210A mutation of the prothrombin gene, a major inherited thrombophilic polymorphism, and prosthetic valve endocarditis.^[5] Several other conditions affecting inflammation and hemostasis could play a role in IE pathogenesis.^[4,6,7] Homocysteine (Hcy), a sulfur-containing amino acid, absent in natural diets, is a metabolic intermediary in transmethylation and transsulfuration reactions. Such reactions are essential to normal cellular growth, differentiation, and function. Excess homocysteine is associated with vascular disease and related disorders.^[8] Hyperhomocysteinemia (h-Hcy) commonly results from inherited or acquired metabolic defects of Hcy metabolism.^[9] It is associated with an increased risk of cardiovascular diseases mediated by endothelial damage and induction of both arterial and venous thrombosis.^[10-13] H-Hcv due to the C677T transition in the methylene-tetra-hydro-folate reductase (MTHFR) gene is currently listed among the major genetically determined thrombophilias.^[14] Thus far, no

study has evaluated the potential role of this condition in the pathogenesis of IE.

Hence, we performed this study with the aim to assess the prevalence, correlates, and possible clinical significance of h-Hcy in patients with IE. Our hypothesis was that IE patients could have an inherently higher rate of h-Hcy.

2. Methods

2.1. Study design

In this case-control study, we included patients with a diagnosis of "definite" IE according to the modified Duke University criteria, admitted to our unit between 2004 and 2014. Additional eligibility criteria were age >18 years, Caucasian ethnicity, hospitalization in our center because of IE, and ability to provide the informed consent. As a result, the enrolled group did not include consecutive IE cases, as those in worse conditions or who went to cardiac surgery emergently were less likely to undergo additional blood sampling for research purposes. As controls, we recruited in our hospital valvular heart disease (VHD) patients with documented absence of IE, including uncomplicated native and prosthetic VHD patients, patients with heart valve disease undergoing cardiac surgical repair/replacement, and carriers of cardiac implantable electronic devices (CIED). Routine blood donors (BD) from our Hospital Transfusion Service were also studied as a further control group made of apparently healthy subjects. They were age- and sex-matched with IE patients in a 2:1 ratio and were selected between 2008 and 2009 among routine BD based on willingness to undergo additional blood sampling and a negative clinical history of heart disease and no prior thromboembolic event. Our hypothesis was that IE patients had an Hcy level that was 25% higher than healthy controls. Assuming a mean Hcy level of 9.2 µmol/L in healthy BD,^[8] a power of 0.8, and an alpha error of 5%, the sample size needed was of about 55 IE patients.

A written informed consent to participation in this study and the collection and storage of DNA for genetic analyses was obtained from each participant. Our Department's Review Board approved the study protocol and procedures, which complied with the Declaration of Helsinki.

2.2. Type of and rationale for the analyses

Hcy plasma levels and the C677T polymorphism in the MTHFR gene were studied because of their association with a prothrombotic state as well as their measurable prevalence in the general population.^[12,13] H-Hcy was defined as a plasma Hcy concentration >15 μ mol/L.^[13] 5,10-MTHFR is a key enzyme in the conversion pathway of Hcy into methionine.^[15] The C677T transition in the MTHFR gene impairs its enzyme activity causing Hcy accumulation, that in turn exerts oxidant, atherogenic, and procoagulant effects.^[8] The phenotypic effects of this polymorphism fully exploit when it occurs in the homozygous form. As a measure of C677T prevalence, allelic frequency was used, that is, the number of mutated alleles divided by twice the number of subjects studied.

To assess whether h-Hcy was possibly associated with the risk of developing IE, we compared IE patients (cases) with BDs (controls). In order to exclude that h-Hcy was associated to heart disease rather than IE, we compared IE patients with VHD patients. Subsequently, to verify the possible association between Hcy levels and a particular form of IE (NVE, PVE, CIEDE), subgroup analyses were carried out. Further, we evaluated whether Hcy levels affected IE vegetation size or the occurrence of embolic events. Finally, we looked at the possible effect of h-Hcy on the major outcomes of IE, including complications, need for surgery, and mortality.

2.3. Clinical procedures

Upon admission to our unit, patients signed their informed consent and underwent a detailed baseline clinical evaluation, including history, physical examination, chest x-ray and abdominal ultrasound scan. During the in-hospital stay, patients were carefully assessed for signs or symptoms suggestive of symptomatic embolic complications, such as sudden neurologic or visual dysfunction, abdominal pain, hematuria, chest discomfort, abrupt dyspnea, or limb pain.^[16] No active search for silent embolic events was systematically done. Contrastenhanced computed tomography scanning was performed to confirm the clinical suspicion of major or symptomatic cerebral, pulmonary, or intra-abdominal embolism.^[17]

Also, 2 to 4 sets of blood cultures were obtained from each patient on the day of admission. Molecular microbiology on whole blood (Septifast) was also performed in cases with recent prior antimicrobial therapy, and valve/device cultures were always obtained in surgical patients. No routine serology for Coxiella and Bartonella or molecular biology on valve samples was performed.

All patients underwent trans-thoracic and, whenever needed and feasible, trans-esophageal echocardiography, measuring number, length (greatest diameter), and location of endocardial vegetations.

A standard panel of laboratory investigations was performed in both patients and controls by our Hospital Central Laboratory, including full blood count and measurement of prothrombin time, expressed as International Normalized Ratio (INR), activated partial thromboplastin time, plasma fibrinogen, and serum creatinine. In IE patients, C-reactive protein (CRP) serum levels were also assessed.

Patients were followed up until discharge from hospital or death. Patients discharged alive were invited to come back within 4 weeks at our outpatient IE clinic. All subjects, irrespective of whether or not returning to visit, were called by phone in order to obtain 1 year after discharge status.

2.4. Analytical procedures

Whole blood, serum, and plasma samples were obtained on the first day of admission from all patients included in the study and from VHD and BD controls. Samples were rapidly processed (centrifuged or refrigerated at 4 °C) and eventually stored in our laboratory at -80 °C. After centrifugation at 3500 rpm for 10 minutes, plasma was separated from EDTA blood and stored. Subsequently, plasma Hcy levels were measured on the single sample obtained on admission by means of a semi-automatic chemiluminescent microparticle immunoassay run on an Architect i2000SR platform (Abbott Diagnostics, Rome, Italy). By means of a rapid spin-column method (QIAquick PCR purification, Qiagen, Milan, Italy), whole blood was used to extract genomic DNA, which in turn was stored at -80 °C. DNA samples were then subjected to polymerase chain reaction assays (Hot Master Mix Tag, 5 Prime, Milan, Italy) with primers (Biofab Research, Milan, Italy) specific for the genetic regions of interest, as previously described.^[17] To detect the C677T SNP in the MTHFR gene, PCR products were digested with restriction endonucleases (Biofab Research, Milan, Italy) and subsequently run on 2% agarose gel electrophoresis (Carlo Erba Reagenti, Milan, Italy).

2.5. Data analysis

Multiple IE patient data, including demographics, predisposing conditions and pre-existing illnesses, causative pathogens, medications taken, presenting signs/symptoms, echocardiographic findings, treatment strategies, disease complications and outcome, were recorded by means of a data collection form and entered in a computerized datasheet for subsequent analysis.

For the purpose of this study, the following variables were considered: age, sex; causative microorganism; site of infection; creatinine and coagulation parameters; vegetation size. One goal of the study was to evaluate whether h-Hcy was associated with major IE outcomes, including intracardiac complications of IE, outcome of hospitalization, and 1-year follow-up status after discharge.

Differences between groups in numerical variables were assessed through Mann–Whitney or Kruskal–Wallis tests for 2 or multiple independent samples, respectively. Categorical variables were compared with the Fisher's exact test. Odds ratios for the differences with 95% confidence intervals were calculated. Correlation between numerical variables was assessed by Spearman's coefficient. Logistic regression analysis of independent predictors of 1-year mortality was performed by block entering in the model all variables associated with this outcome on the univariate analysis (P < 0.1).

Genotype data were analyzed for deviation from the Hardy— Weinberg equilibrium by calculation of Pearson's chi-square.

Numerical data are presented as mean \pm standard deviation (SD) or median and interquartile range (IQR). Categorical data are presented as the number and the percent. Statistical analyses were carried out with the aid of the SPSS 20.0 software. The significance level was set at 5% and all tests were 2-tailed.

3. Results

Included in the analysis were 246 IE patients, showing either left heart or right heart infection (149 and 97 patients, respectively), and including 116 patients with native valve IE (NVE), 55 with prosthetic valve IE (PVE), and 75 with cardiac implantable electronic device (CIED)-related IE. As controls, we studied 2 groups: (a) 106 healthy blood donors (BD); (b) 258 VHD patients without IE, including 154 uncomplicated native and prosthetic VHD patients, 30 patients with heart valve disease undergoing cardiac surgical repair/replacement, and 74 carriers of a cardiac implantable electronic device (CIED). BD were significantly younger than IE patients; therefore, only a subgroup of the latter, comprising the youngest IE patients, was compared with BDs.

3.1. Homocystein levels in Infective Endocarditis and controls

As shown in Table 1, IE patients showed higher levels of plasma Hcy compared to an age- and sex-matched group of healthy BDs. A condition of hyper-Hcy was more common among IE patients than their healthy counterparts (Table 1). Only a small subgroup of IE patients was compared to BD as the overall IE group had a much higher age (see below and Table 2). To assess whether h-Hcy was associated with heart disease rather than IE itself, we

Table 1

Hcy levels in IE patients compared to an age and sex-matched group of healthy BDs.

	Infective endocarditis (n = 53)	Healthy subjects (n=106)	Р
Age, y	39 (15-61)	36.5 (19-63)	0.624
Male gender	39 (73.6)	83 (78.3)	0.553
Body mass index, kg/m ²	23.5 (17.3-42)	24.7 (18.7–32.4)	0.929
Homocysteine, µmol/L	15 (3–56)	12.6 (5.6-47.3)	0.014
Hyper-homocysteinemia	26 (49.1)	27 (26.7)	0.007
MTHFR, T allele frequency	0.52	0.43	0.107

Data are median (range) or number (percent).

Hcy = homocysteine, IE = infective endocarditis, MTHFR = 5,10 methylene tetrahydrofolate reductase.

compared IE patients with a cohort of VHD patients. Characteristics of these 2 groups are comparatively shown in Table 2. Hcy levels were not significantly different between IE and VHD patients.

3.2. Correlates of hyperhomocysteinemia in IE

In each of the 3 groups studied, TT homozygotes at position 677 of the MTHFR gene showed significantly higher Hcy levels (data not shown). The allelic frequency of this polymorphism was not significantly different when we compared IE patients and BD (Table 1). Similarly, there were no differences in allele frequency between IE and VHD patients (Table 2). Despite of this, h-Hcy was observed significantly more often in IE and VHD patients, suggesting the effect of factors other than the MTHFR genotype. We, therefore, studied clinical variables associated with h-Hcy.

Table 2

Clinical and biochemical features of IE and VHD patients.

	Infective endocarditis	Valvular heart disease	Р
N	246	258	NA
Age, y	62 (13-90)	60 (23–93)	0.5
Male gender	181 (73.5)	176 (68.2)	0.2
Chronic heart failure	107 (43.5)	204 (79.1)	0.01
Diabetes	48 (19.5)	69 (26.7)	0.16
Chronic obstructive Pulmonary disease	48 (19.5)	108 (41.9)	0.001
Liver disease	55 (22.4)	56 (21.7)	0.830
Chronic kidney disease	45 (18.3)	51 (19.7)	0.734
Creatinine, mg/dL	1 (0.6-9.1)	1.1 (0.7-2.5)	0.06
Microbial etiology			
Staphylococcus aureus	32 (13)	NA	
Coagulase-neg. staphylococci	55 (22.4)	NA	
Streptococci	66 (26.8)	NA	
Enterococci	25 (10.2)	NA	
Other	21 (8.5)	NA	
Negative cultures	47 (19.1)	NA	
Vegetation size, mm	13 (3-40)	NA	
Symptomatic embolic complication	99 (40.2)	NA	
Homocysteine µmol/L	14.9 (3-81)	16 (5-50)	0.08
Hyper homocysteine	120 (48.8)	144 (55.8)	0.13
MTHFR, T allele frequency	0.451	0.402	0.21

Data are median (range) or number (percent).

IE = infective endocarditis, INR = international normalized ratio, MTHFR = 5,10 methylene tetrahydrofolate reductase, NA = not applicable, VHD = valvular heart disease.

Table 3

Clinical features of IE patients with or without hyperhomocysteinemia.

	Hyperhomocysteinemia			
	Yes (n=120)	No (n=126)	OR (95% CI)	Р
Age, y	65 (19-86)	58 (13-87)		0.06
Male gender	91 (75.8)	90 (71.4)	0.79 (0.45-1.4)	0.47
Vegetation location				0.39
Aortic valve	44 (38.3)	55 (46.6)		
Mitral valve	24 (20.9)	18 (15.3)		
Tricuspid/pulmonary valve	6 (5.2)	9 (7.6)		
CIED	41 (35.7)	36 (30.5)		
Chronic heart failure	51 (43.6)	56 (47.1)	0.86 (0.5-1.45)	0.60
Diabetes	23 (19.7)	25 (21)	0.92 (0.4-1.73)	0.87
Liver disease	15 (18.5)	15 (17.9)	1.04 (0.4–2.3)	0.999
Chronic kidney disease	31 (27.7)	14 (13.1)	2.54 (1.2-5.1)	0.01
Creatinine, mg/dL	1.1 (0.1-8.3)	0.9 (0.1–9.1)		0.08
Platelet count, $\times 10^{3}/\mu$ L	180 (0.16-640)	209 (47-453)		0.30
Prothrombin time, INR	1.3 (1-4)	1.3 (1–5)		0.88
Vegetation size, mm	12.7 (0.5–36)	14.7 (0.3–40)		0.45
Symptomatic embolic event	48 (41.4)	51 (42.9)	0.94 (0.5-1.5)	0.895
Length of hospitalization (days)	23	24		0.99
Cardiac complications*	39 (48.1)	45 (54.7)	0.80 (0.43-1.4)	0.534
Indication for surgery	89 (79.5)	88 (83)	0.79 (0.4–1.5)	0.60
Hospital mortality	9 (7.5)	6 (4.8)	1.62 (0.5-4.7)	0.431
1-year mortality	20 (19.6)	11 (9.9)	2.21 (1.00-4.89)	0.05

Data are median (range) or number (percent).

Indicates a composite endpoint including intracardiac abscess/fistula, heart failure, and arrhythmias.

CI=confidence interval, CIED=cardiac implantable electronic device, INR=international normalized ratio, OR=odds ratio.

As shown in Table 3, IE patients with h-Hcy were older and had a higher prevalence of chronic kidney disease. In IE patients, Hcy levels were directly correlated to those of creatinine (r=0.337; P=0.0001) and age (r=0.177; P=0.006). Hcy did not correlate with C-reactive protein (r=-0.053; P=0.443).

A trans-esophageal echocardiography was performed in 162 (66%) patients, including 58% of NVE, 78% of PVE, and 69% of CIED-IE cases. IE patients with h-Hcy had neither larger vegetations nor a higher rate of symptomatic embolic complications (Table 3). In particular, there were no differences in median Hcy values according to vegetation size (15.5 μ mol/L for vegetation ≤ 10 mm and 16.0 μ mol/L for vegetation >10 mm; *P* = 0.71). Likewise, there was no difference in the prevalence of patients with h-Hcy between small and large vegetation subgroups (53% vs 52.8%, respectively). Similarly, IE patients suffering an embolic complication did not have a higher prevalence of h-Hcy compared to those who did not (48.5% vs 50.0%; *P*=0.89). In particular, median Hcy levels were 14.3 μ mol/L in patients with and 15.2 μ mol/L in those without symptomatic embolism (*P*=0.75).

Hcy levels were not significantly higher in left-sided IE cases (14.7 vs $15.9 \,\mu$ mol/L in right-sided IE; P=0.67). Likewise, no

differences in Hcy levels were observed between patients with NVE, PVE, or CIED-related IE (data not shown). IE patients who were deemed clinically to have heart failure as a consequence of IE did not show higher Hcy levels (14.1 vs 15.3 μ mol/L in those without *de novo* heart failure; *P*=0.93).

Table 4 presents data on Hcy levels, MTHFR polymorphism, vegetation size, and symptomatic embolic complications according to IE microbial etiology. No association was found between the causative pathogen and Hcy levels. In particular, there were no differences between cases due to *Staphylococcus aureus* and other etiologies. Significantly larger vegetations were observed among *S aureus* IE cases (Table 4). Embolic complications were also significantly associated with the microbial etiology of IE, with higher rates in cases due to *S aureus* and lower rates in enterococcal IE (Table 4).

3.3. Association of hyper-homocysteinemia with major IE outcomes

The major outcomes of IE according to the presence of h-Hcy are presented in Table 3. There were no significant differences in hospitalization length, rate of major complications, and need for

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	Staphylococcus aureus (N = 32)	Coagulase-negative <i>Staphylococci</i> (N=55)	Streptococci (N = 66)	<i>Enterococci</i> (N = 25)	Negative Cultures (N=47)	Other (N = 21)	Р
Hcy levels median (range)	14 (6-32)	16.8 (4-40)	15.7 (8–51)	14.2 (5–29)	15.3 (8–51)	13.8 (3–81)	0.51
Hyper-Hcy $>$ 15 μ mol/L, (%)	12 (38.7)	29 (53.7)	34 (52.3)	11 (44)	19 (51.3)	9 (42.8)	0.74
MTHFR C677T AF, %	47	40	47 ^a	52	46.7	33.3 ^a	0.024
Vegetation size >10 mm, %	69.5	57.1	38.6 ^{a,b}	40 ^{c,d}	73.9 ^{a,d}	92 ^{b,c}	0.002
Symptomatic embolic event, %	67.7 ^{a,b,c,d}	46.3 ^e	40 ^b	20 ^{c,e}	40.5 ^a	30 ^d	0.01

Each superscript letter denotes of microbial etiology categories whose column proportions do differ significantly from each other at the 0.05 level.

AF = allele frequency, Hcy = homocysteine, MTHFR = 5,10 methylene tetrahydrofolate reductase.

Table 5

Multivariate analysis of 1-year mortality predictors in the 246 IE patients.

	1-year outcome		Univariate analysis			Logistic regression		
	Alive	Dead	Odds ratio	(95% CI)	*Р	Odds ratio	(95% CI)	* P
Age			3.09	(1.4–6.77)	0.005	2.63	(1.17–5.89)	0.02
>65 y	61	19						
VS								
≤65 y	119	12						
Нсу			2.21	(1.00-4.89)	0.05	1.87	(0.83-4.23)	0.13
$>15\mu$ mol/L	82	20						
VS								
\leq 15 μ mol/L	100	11						
Type of IE			0.49	(0.18–1.31)	0.19			
Native	92	10						
VS								
Prosthetic	41	9						
Type of IE			0.87	(0.33-2.29)	0.81			
Prosthetic	41	9						
VS								
Device	49	12						
Vegetation size			1.11	(0.39-3.14)	0.99			
>10 mm	64	10						
VS								
\leq 10 mm	50	7						
Surgery			0.47	(0.21-1.02)	0.06	0.57	(0.25-1.27)	0.17
Yes	131	17						
VS								
No	51	14						
Causative pathogen			1.17	(0.37-3.7)	0.99			
Staphylococcus aureus	22	4						
VS								
Other pathogens	155	24						
creatinine			1.54	(0.68-3.50)	0.31			
>1.0 mg/dL	68	16						
VS								
\leq 1.0 mg/dL	79	12						
Heart failure			1.10	(0.49-2.45)	0.84			
Yes	85	14						
VS								
No	94	14						

* P was generated by Fisher's exact test.

CI = confidence interval, IE = infective endocarditis.

cardiac surgery between patients grouped according to the presence of h-Hcy. Hospital mortality was not associated with h-Hcy (Table 3). In contrast, 1-year mortality was higher in IE patients with h-Hcy (19.6% vs 9.9%; OR 2.21 [95% CI 1.0–4.9]; P=0.05). Median Hcy was higher in subjects not surviving at 1 year (17.3 [5–81] vs 14 [3–56] in those surviving; P=0.09). In order to further assess the relationship between h-Hcy and 1-year mortality, we analyzed factors possibly associated with this outcome. As shown in Table 5, a significant association was found between 1-year mortality and patient age (23.7% in patients >65 years vs 9.1% in those ≤ 65 years; OR 3.1 [95% CI 1.4–6.77]; P=0.005) as well as h-Hcy. At logistic regression analysis, age, but not h-Hcy, was an independent predictor of 1-year mortality (Table 5).

4. Discussion

The IE vegetation is made of bacteria embedded in a plateletfibrin mesh.^[3,18,19] We hypothesized that h-Hcy, by inducing a prothrombotic state, could favor IE vegetation formation, influencing incidence and clinical course of the disease. In this study, Hcy levels and the prevalence of h-Hcy were higher in IE compared with healthy BD. This finding pointed to a possible role of h-Hcy in the pathogenesis of IE. Previous clinical observations had indeed led us to hypothesize a potential role for h-Hcy with MTHFR C677T homozygous mutation in nonbacterial thrombotic endocarditis^[20] and in pace-maker lead IE.^[21] However, the equally higher levels observed in VHD as well as the correlation of Hcy with creatinine levels suggest h-Hcy could be a marker of cardiac dysfunction related to IE rather than a condition predisposing to IE. This interpretation is also consistent with the absence of differences in MTHFR C677T genotype across the 3 study groups. However, we lack data showing increased Hcy levels in IE with symptomatic heart failure.

H-hcy could not be a mere marker of cardiac dysfunction in IE. Interestingly, we observed a significant association of h-Hcy with 1-year mortality. Hcy levels also tended to be higher in IE patients dying in hospital. Further studies are needed to ascertain whether h-Hcy is solely a marker of risk or whether it contributes to a worse prognosis due to its potential atherogenic, proinflammatory, and prothrombotic effects.

At variance with what we expected, Hcy levels did not affect vegetation size or major embolic complications. This finding further suggests that embolic events in IE are the result of a complex interaction between host and microbial factors. An investigation of the risk factors for embolism in IE was, however, beyond the aims of the present study.

This work has some important limitations. We compared with age-matched healthy BD only a selected subgroup IE patients made of the youngest subjects, and we lacked a control group made of patients with bacteremia without IE. Only symptomatic embolic events were diagnosed and analyzed. The overall mortality which we observed was low, likely as we were less successful to obtain samples of blood for Hcy and genetics from the most severely ill patients or those undergoing emergency cardiac surgery. Despite of these lower rates of mortality, differences were observed in patient outcome at 1 year.

In conclusion, we screened a cohort of 246 IE patients for h-Hcy and its major genetic ground. Based on our current data, we are unable to support a role for this condition in the susceptibility to IE. However, IE patients have a high rate of h-Hcy and this is associated with late mortality, but not vegetation size or symptomatic embolic events. Further studies are surely warranted to better understand the role of h-Hcy and other inherited or acquired homeostasis defects in the pathogenesis and clinical course of IE.

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