










PON1 polymorphisms can predict generalized anxiety and depressed mood in patients with multiple chemical sensitivity

Alessio Mosca^{‡,1} , Antonio Del Casale^{*,‡,2,3} , Marina Borro^{1,4} , Giovanna Gentile^{1,4} , Leda Marina Pomes^{1,4}, Alessio Padovano^{1,3}, Federica Fiaschè^{1,3}, Vito Pinzone^{1,3}, Chiara Rapinesi¹, Teodolinda Zoppi^{1,3}, Roberto Brugnoli^{1,3}, Gabriele Sani⁵ , Georgios Demetrios Kotzalidis¹ , Paolo Girardi^{1,3} , Stefano Ferracuti⁶ , Maurizio Simmaco^{1,4} & Maurizio Pompili^{1,3} 

¹ Department of Neuroscience, Mental Health, & Sensory Organs (NESMOS), Faculty of Medicine & Psychology, Sapienza University, Rome, Italy

² Department of Clinical & Dynamic Psychology, Faculty of Medicine & Psychology, Sapienza University, Rome, Italy

³ Unit of Psychiatry, 'Sant'Andrea' University Hospital, Rome, Italy

⁴ Unit of Laboratory & Advanced Molecular Diagnostics, 'Sant'Andrea' University Hospital, Rome, Italy

⁵ Institute of Psychiatry & Psychology, Department of Geriatrics, Neuroscience & Orthopedics, Fondazione Policlinico Universitario IRCCS "A. Gemelli", Università Cattolica del Sacro Cuore, Rome, Italy

⁶ Department of Human Neuroscience, Sapienza University; Unit of Risk Management, 'Sant'Andrea' University Hospital, Rome, Italy

*Author for correspondence: Tel.: +390633777774; antonio.delcasale@uniroma1.it

‡Authors contributed equally

Background: Multiple chemical sensitivity (MCS) is a chronic condition with somatic, cognitive and affective symptoms that follow contact with chemical agents at usually non toxic concentrations. We aimed to assess the role of genetic polymorphisms involved in oxidative stress on anxiety and depression in MCS. **Materials & methods:** Our study investigated the *CAT* rs1001179, *MPO* rs2333227, *PON1* rs662 and *PON1* rs705379 polymorphisms in MCS. **Results:** The AG genotype of the *PON1* rs662 and the TT and CT genotypes of the *PON1* rs705379 were involved in anxiety and depression. **Discussion:** These results are in line with existing evidence of *PON1* involvement in MCS and suggest a further role of this gene in the exhibition of anxiety and depression in this disease.

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Multiple chemical sensitivity (MCS) is a chronic syndrome characterized by different somatic, cognitive and affective symptoms occurring after exposure to chemical agents at concentrations that are not usually associated with toxic reactions in the general population [1,2]. MCS etiology and pathogenesis are not clarified, with no laboratory abnormality being consistently associated with it. It mainly affects women between 40 and 50 years [3,4], without differences in ethnicity, education and economic status [5]; its prevalence is not well defined since it is not a universally accepted diagnosis [6–8]. Common symptoms include weakness, lethargy, sore throat, hyperosmia, dyspnea, dizziness, headache, confusion and difficulty concentrating [1].

The Cullen Criteria were extended by Lacour [9], according to whom MCS is a chronic condition that causes significant lifestyle or functional impairments for at least 6 months (a); CNS recurrently symptoms are related to self-reported odor hypersensitivity (b); at least one CNS symptom is associated to at least one symptom of another organ/system (c); symptoms are related to the exposure to low levels (d) of multiple unrelated chemicals (e) and improve or are resolved when exposure is avoided (f).

Table 1. Genes possibly involved in the manifestation of psychological symptoms in multiple chemical sensitivity.

Gene	Name	Function	Mechanisms involved	Ref.
<i>PON1</i>	Paraoxonase-1	Antioxidant activity of HDL and protection of LDL from oxidative modification Biodegradation of organophosphates and oxidized phospholipids. Hydrolysis of pesticides and nerve gases	Increase of lipid peroxidation, oxidative stress and hydroperoxide lipids. Differences in the regulation of enzymatic activity, expression and polymorphisms have been reported in most psychiatric disorders	[16–20]
<i>CAT</i>	Catalase	Converts hydrogen peroxide into water and oxygen. Clearance of reactive oxygen species (ROS)	Increased ROS and oxidative stress. Differences in the regulation of enzymatic activity have been reported in patients with GAD and depression	[21–23]
<i>MPO</i>	Myeloperoxidase	Peroxidase enzyme expressed in neutrophil granulocytes produces hypohalous acids to carry out their antimicrobial activity. Involved in inflammation regulation and oxidative stress.	The increase in activity is associated with a rise of oxidative stress, inflammation and mood disorders	[24–29]
<i>SOD2</i>	Manganese-dependent superoxide dismutase	Clear mitochondrial ROS	Increased ROS and oxidative stress. Differences in the regulation of enzymatic activity have been reported in patients with anxiety and depression	[21–23]
<i>GSTM1</i> , <i>GSTT1</i> , <i>GSTP1</i>	Glutathione S-transferase	A family of isozymes involved in the detoxification of xenobiotics by catalyzing the nucleophilic attack by GSH	Increased oxidative stress and inflammatory processes. Differences in the regulation of enzymatic activity and polymorphisms have been reported in mood disorders	[30]

Anxiety and depression have been attributed to the social and work hardships related to the disease, and frustration of being affected by medically unexplained conditions [10]. Besides this, some authors argued that MCS diagnosis is a misdiagnosis of an unrecognized psychiatric disorder [11,12] or a ‘psychosomatic’ illness [13,14]. These theses are supported by a remarkable overlap between the diagnostic criteria of somatoform disorders and MCS symptoms [15]. Furthermore, anxiety and depression may also constitute a principal component between MCS symptoms, even relatively to genomic correlates (Table 1), although no somatic origins of these symptoms have been demonstrated.

Individual responses to chemicals are so varied that over time a very wide range of nosological terms and entities have been used to describe sensitivity to environmental factors and chemical intolerances, including ‘chemical intolerance’, ‘environmental sensitivities’ [10], ‘ecological illness’, ‘environmental illness’ and ‘idiopathic environmental intolerances’ (IEI), a term coined by the WHO in 1996, which also refers to hypersensitivity to electromagnetic fields, radio signals, loud noise and intolerance to odors [14]. Today, considering that symptoms do not always occur in response to a chemical, MCS is viewed as an aspect of the IEI, and in the scientific literature, the terms MCS and IEI are mostly interchangeable [13,14].

Patients with MCS have an increased likelihood of having a history of a psychiatric disorder [31] and a very high rate of psychiatric comorbidity. MCS patients significantly showed higher rates of depression and anxiety than the general population, with more frequent lifetime diagnoses of major depressive disorder and/or generalized anxiety disorder [12,32,33]. Other reported frequent psychiatric comorbidities were histrionic personality disorder [15,34], panic attacks [32], as well as trauma and childhood abuse [13,35].

MCS has different biological correlates. Alterations of the nitric oxide/peroxynitrite (NO/ONOO⁻) cycle were related to the formation of peroxynitrite, and subsequent neuronal sensitization and neuroinflammation involved in MCS [36]. Oxidative stress has been implicated in the etiopathogenesis of both MCS and other psychiatric disorders [16,17,37], in which the SOD2 [21–23] and GSTM1 [30] genes were involved. Furthermore, oxidative stress has been related to mitochondrial dysfunction [38,39] and changes in neuroplasticity [40,41].

Genes with a role in oxidative stress, including paraoxonase1 (*PON1*) [42,43], catalase (*CAT*) [21–23,40,41,44,45] and myeloperoxidase (*MPO*) [46,47] were consistently involved in the pathophysiology of anxiety and depression. Changes in the activity of paraoxonase have been considered possible pathophysiological correlates of MCS [48].

The *PON1* gene is located on the chromosome 7 long arm (7q21.3-22.1) [49] and encodes paraoxonase, a calcium-dependent glycoprotein synthesized in the liver and secreted in the blood. It bounds the HDL, and is responsible for most of their antioxidant activity, protecting the LDL from oxidative modification, and counteracting the formation of atherosclerotic plaques. *PON1* plays a key role in the biodegradation of various organophosphates (OPs) and oxidized phospholipids and participates in hydrolysis of pesticides and nerve gases [50]. *PON1* A-575G polymorphism has been related to the magnitude of paraoxonase activity [51], while the C-108T polymorphism to variation in the amount of its activity [52]. Different antidepressants (i.e. mirtazapine and escitalopram), displayed a potential inhibitory effect on *PON1* activity *in vitro* [53], as well as on 6-phosphogluconate dehydrogenase and

glucose-6-phosphate dehydrogenase enzymes [54]. Also, other drugs, including some sulfonamides, antihypertensives, antineoplastics, some quinones, benzenesulfonamide derivatives and usnic and carnolic acids showed potential inhibitory effects on PON1 [55–60].

PON1 A-575G heterozygotes are more likely to present neurological symptoms, which were more common in Gulf War Syndrome veterans [61–63]; a lower mean activity of the *PON1* enzyme has been reported in post-traumatic stress and attention deficit and hyperactivity disorders [64]. Some biological correlates of post-traumatic stress could be shared with MCS, fibromyalgia and chronic fatigue syndrome [65,66].

Patients with bipolar disorder showed diminished *PON1* activity correlated with A-575G polymorphism, which suggested that these polymorphisms may play a role in the pathophysiology of bipolar disorder, type I, due to the decreased ability to cope with stress oxidative [18]. Furthermore, A-575G polymorphism was significantly associated with depressive symptoms in women [38]. *PON1* polymorphisms, activity and expression have been correlated with the diagnosis of schizophrenia [67–69], first-episode psychosis [19] and mood disorders [20,42,70].

MPO (myeloperoxidase) is a heme enzyme expressed in brain cells [71] and immune cells; it is involved in inflammation regulation and oxidative stress [72]. Its activation is correlated to hypochlorous acid and other toxic oxidants production, with subsequent antimicrobial function [24,25,73]. Low *MPO* levels correlated with reduced risk of inflammatory states [26] and depression [74–77]. *MPO* deficiency also attenuates cytokine production [27]. *MPO* has been associated with oxidative stress and inflammation and involved in recurrent MDD and related cognitive dysfunction [28,29], and in bipolar I disorder [78].

CAT (Catalase) is one of the primary antioxidant enzymes against superoxide. It catalyzes the decomposition of hydrogen peroxide to oxygen and water and plays an important role in reactive oxygen clearance [79]. Different studies have described *CAT* polymorphisms and activities with various pathophysiological states and diseases [80], including major depressive [44] and bipolar [81] disorders, and schizophrenia [82].

Our main hypothesis was that *PON1*, *CAT* and *MPO* polymorphisms might have a role in the pathophysiology of anxiety and depressive symptoms in patients with MCS. The main objective of this study was to identify gene polymorphisms that constitute predictors of anxiety and depressive symptoms in patients with MCS.

Materials & methods

This was a single-center observational study conducted at the Centre of Personalised Medicine, "Sant'Andrea" University Hospital, Rome, Italy, during the years 2014–2015. All admitted patients required a medical consult and a pharmacogenomics assessment for previous and proven drug treatment resistance. All those who manifested their interest in participating were considered possible candidates.

Inclusion criteria were a manifestation of MCS syndrome according to the Cullen [1] and Nethercott *et al.* [83] criteria for MCS extended by Lacour (see description above) [9]; age ≥ 18 years; and ability to provide valid informed consent. Exclusion criteria included pregnancy, recent brain injury, severe medical illness, DSM-5 [84] diagnoses of schizophrenia-spectrum disorder, substance use disorder, other acute psychiatric disorders, and inability to provide informed consent. Based on our inclusion and exclusion criteria, the final sample included 129 of 173 assessed patients.

During the medical consult, the physician investigated the presence of MCS symptoms through the compilation of a checklist of symptoms as described in another study of our group [85]. Participants underwent venous blood sampling for functional biochemical analyses. All patients were fully informed about the observational nature of the study, and each provided written informed consent.

Genetic analyses

Genomic DNA was isolated from peripheral blood samples using the X-tractor Gene system (Corbett Life Science, Australia), according to the manufacturer's protocol. We studied the following polymorphisms: *CAT* C-262T (rs1001179), *MPO* G-463A (rs2333227), *PON1* A-575G (rs662) and *PON* C-108T (rs705379). We performed genotyping with the MassARRAY system, in other words, a MALDI-TOF mass spectrometry platform by the iPLEX chemistry (Agena Bioscience, CA, USA). We used the Agena Bioscience Assay Design 4.0 software to design the locus-specific amplification and extension SNP primers (Table 2), considering a success rate of 97.9–100% for the genotype call of each SNP.

Table 2. Amplification and extension primers used in the multiplex genotyping assay.

Gene name	Polymorphism		Oligonucleotides (F: forward; R: reverse; E: extension)
CAT C-262T	rs1001179	F	ACGTTGGATGAGCAATTGGAGAGCCTCGC
		R	ACGTTGGATGAGGATGCTGATAACCGGGAG
		E	AGCCCCGCCCTGGGTTCCGGCTAT
MPO G-463A	rs2333227	F	ACGTTGGATGCTTAGCCACATCATCAAT
		R	ACGTTGGATGGGCTGGTAGTGCTAAATTC
		E	CTTTGGGAGGCTGAGGC
PON1 A575G	rs662	F	ACGTTGGATGTAGACAACATACGACCACGC
		R	ACGTTGGATGGATCACTATTTTCTTGACCC
		E	TTTCTTGACCCCTACTTAC
PON C-108T	rs705379	F	ACGTTGGATGCTTCTGTGCACCTGGTCGG
		R	ACGTTGGATGTGCTGGGGCAGCGCCGATT
		E	CGCCGATTGGCCCGCCC

Statistical analyses

We used the IBM SPSS[®] Statistics 24.0 (Armonk, NY, USA: IBM Corp. 1989, 2016) for all analyses, except for the Hardy–Weinberg equilibrium deviation test, for which we used an online calculator based on the methods of Rodriguez *et al.* [86,87].

We performed descriptive statistics with a one-way analysis of variance for the continuous variables and the *Chi-square* (χ^2) test for categorical variables. We performed a binary logistic regression with the 'Enter' method, using the presence/absence of both generalized anxiety and depressed mood as the dependent variable, and the polymorphisms of genes involved in cellular oxidative stress (*PON1* A575G, *PON1* C-108T, *CAT* C-262T and *MPO* G-463A) as the independent, categorical variables. We set the cut off for statistical significance at (two-tailed) $p < .05$.

Results

Study participants were 129 MCS patients (112 women and 17 men) with a mean age of 51.58 years (SD = 11.34). The group of women did not differ significantly from the group of men in age ($F = 0.603$; $p = 0.439$). The sample showed significant prevalence of women ($\chi^2 = 69.961$; $p < .001$). Patients most frequently showed hyperosmia (96.8%), asthenia (82.4%) and dyspnea (81.6%). We reported the main clinical characteristics of this sample in another study on MCS [85]. Considering the whole sample, 70 patients (54.26%) had comorbid anxiety and depression (63 women, 7 men; mean age = 51.75 years, SD = 10.07); 7 patients showed generalized anxiety but not depressed mood; one patient had depressed mood but not anxiety; the other 51 patients did not show generalized anxiety nor depressed mood. We summarized the sociodemographic and clinical characteristics of the study sample in Table 3.

The analyzed SNPs were in Hardy–Weinberg equilibrium. Logistic regression showed that our model was significant for a good predictability of comorbid anxiety and depressive symptoms (Omnibus Tests of Model Coefficients $\chi^2 = 22.263$; $p = 0.004$) and explained over 25% of the variance (Nagelkerke $R^2 = 0.253$). Overall, this logistic regression model proved to be useful, with good positive predictive value (66.4%). *PON1* A575G and *PON1* C-108T polymorphisms were significant predictors of comorbid anxiety and depressed mood. The odds ratio (OR) for *PON1* A575G AG versus GG was 10.403 (unstandardized coefficient $B = 2.342$; $p = 0.011$; 95% CI: 1.71–63.43), while that of AG versus AA was 4.04 ($B = 1.397$; $p = 0.008$; 95% CI: 1.44–11.36). The overall effect of *PON1* A-575G polymorphism on anxiety and depressive symptoms was significant (Wald = 9.646; df: 2; $p = 0.008$). The OR for *PON1* C-108T CT versus CC was 5.559 ($B = 1.715$; $p = 0.006$; 95% CI: 1.63–19), and for TT versus CC was 4.814 ($B = 1.571$; $p = 0.019$; 95% CI: 1.3–17.86). The overall effect of *PON1* C-108T polymorphism on anxiety and depressive symptoms was significant (Wald = 7.925; df: 2; $p = 0.019$). The logistic regression showed that *CAT* C-262T and *MPO* G-463A polymorphisms did not predict comorbid anxiety and depressed mood (Table 4).

Discussion

Our results showed that two polymorphisms of the *PON1* gene, which is involved in cellular detoxification, predict comorbid anxiety and depressed mood in patients with MCS. Our sample was mainly composed of women, in line with many reports of the high prevalence of MCS in middle-aged women in the USA, Europe and Japan [88–90].

Table 3. Demographic and clinical characteristics of the study sample.

Sample	Men	Women	Test	p-value
Number	17	112	$\chi^2 = 69.961$	<0.001
Age (standard deviation)	49.59 (10.67)	51.89 (11.46)	1-way ANOVA $F = 0.603$	0.439
Symptom	Absent (%)	Present (%)	χ^2	p-value
Hyperosmia	3.2	96.8	109.512	<0.001
Asthenia	17.6	82.4	52.488	<0.001
Dyspoea	18.4	81.6	49.928	<0.001
Cough	26.4	73.6	27.848	<0.001
Cephalalgia (headache)	28	72	24.2	<0.001
Tachypnea	28.8	71.2	22.472	<0.001
Attention deficit	30.4	69.6	19.208	<0.001
Sense of confusion	33.6	66.4	13.448	<0.001
Nausea	35.2	64.8	10.952	0.001
Sense of obnubilation	36.8	63.2	8.712	0.003
Sense of suffocation/choking	41.6	58.4	3.528	0.06
Dyspepsia	41.6	58.4	3.528	0.06
Sleep disturbance	42.4	57.6	2.888	0.089
Paresthesia	44	56	1.8	0.18
Generalized anxiety	44.8	55.2	1.352	0.245
Decision-making deficit	46.4	53.6	0.648	0.421
Arthromyalgia	47.2	52.8	0.392	0.531
Gastric pyrosis	48.8	51.2	0.072	0.788
Pruritus	49.6	50.4	0.008	0.929
Depressed mood	49.6	50.4	0.008	0.929
Dizziness	53.6	46.4	0.648	0.421
Rash	54.4	45.6	0.968	0.325
Fibromyalgia	55.2	44.8	1.352	0.245
Working memory deficit	55.2	44.8	1.352	0.245
Erythema	55.2	44.8	1.352	0.245
Diarrhea	57.6	42.4	2.888	0.089
Meteorism	57.6	42.4	2.888	0.089
Motor incoordination	58.4	41.6	3.528	0.06
Palpitations	58.4	41.6	3.528	0.06
Chest tightness	64	36	9.8	0.002
Vomiting	64	36	9.8	0.002
Hyporexia	66.4	33.6	13.448	<0.001
Trembling	70.4	29.6	20.808	<0.001
Pressure peaks	70.4	29.6	20.808	<0.001
Gastro-esophageal reflux	72	28	24.2	<0.001
Cystitis	73.6	26.4	27.848	<0.001
Recurrent fever	76	24	33.8	<0.001

The AG genotype of the A-575G polymorphism of the *PON1* gene has about tenfold probability of manifestation of anxiety and depression compared with the GG genotype. The TT genotype of the C-108T polymorphism of the *PON1* gene also showed higher (4.5) probability of depressed mood and anxiety compared with the CC genotype, and the CT versus CC also showed a risk higher than 5.5-fold.

This is in-line with the existing evidence of involvement of *PON1* in the manifestation of MCS [48], although there are some inconsistent findings in this regard [91–93]. However, our results suggest an additional and specific role for both the *PON1* A-575G and C-108T polymorphisms in the exhibition of anxiety comorbid with depressed mood in MCS.

Table 4. Logistic regression model for predictors of comorbid generalized anxiety and depressed mood.												
Variables in the equation		B	SE	Wald	df	p-value	Exp(B)	95% CI for EXP(B)				
								Lower	Upper			
Step 1	CAT C262T (CC)			2.736	2.000	0.255						
Contrast 1												
	CAT C262T (CT)	0.304	0.464	0.429	1.000	0.512	1.355	0.546	3.366			
	CAT C262T (TT)	-1.926	1.345	2.051	1.000	0.152	0.146	0.010	2.034			
	PON1 A575G (AA)			9.646	2.000	0.008						
	PON1 A575G (AG)	1.397	0.527	7.027	1.000	0.008	4.043	1.439	11.356			
	PON1 A575G (GG)	-0.945	0.861	1.204	1.000	0.272	0.389	0.072	2.102			
	PON1 C108T (CC)			7.925	2.000	0.019						
	PON1 C108T (CT)	1.715	0.627	7.487	1.000	0.006	5.559	1.627	18.996			
	PON1 C108T (TT)	1.571	0.669	5.516	1.000	0.019	4.814	1.297	17.864			
	MPO G463A (AA)			2.876	2.000	0.237						
	MPO G463A (AG)	-0.381	1.340	0.081	1.000	0.776	0.683	0.049	9.448			
	MPO G463A (GG)	0.412	1.336	0.095	1.000	0.758	1.509	0.110	20.703			
	Constant	-1.530	1.462	1.096	1.000	0.295	0.217					
Step 1	CAT C262T (TT)			2.736	2.000	0.255						
Contrast 2												
	CAT C262T (CT)	2.230	1.370	2.647	1.000	0.104	9.298	0.634	136.423			
	PON1 A575G (GG)			9.646	2.000	0.008						
	PON1 A575G (AG)	2.342	0.922	6.447	1.000	0.011	10.403	1.706	63.427			
	PON1 C108T (TT)			7.925	2.000	0.019						
	PON1 C108T (CT)	0.144	0.519	0.077	1.000	0.781	1.155	0.418	3.193			
	MPO G463A (GG)			2.876	2.000	0.237						
	MPO G463A (AG)	-0.793	0.468	2.875	1.000	0.090	0.453	0.181	1.132			
	Constant	-2.418	1.597	2.294	1.000	0.130	0.089					
			Omnibus tests of model coefficients			Model summary			Hosmer and Lemeshow Test			
			Chi-square	df	p-value	- 2 Log likelihood	Cox & Snell R Square	Nagelkerke R Square	Chi-square	df	p-value	
Step 1			22.263	8	0.004	123.358	0.188	0.253	1.193	8	0.997	
Classification												
Observed		Predicted, selected cases										
		Comorbid AD				Percentage correct						
		No		Yes								
Step 1	Comorbid AD	No	25	20	55.6							
		Yes	16	46	74.2							
Overall percentage						66.4						
Dependent variable: Comorbid generalized anxiety and depressed mood. Independent variables entered on Step 1: CAT C262T, MPO G463A, PON1 A575G and PON1 C108T polymorphisms. Step 1 contrast 1: Contrast reference categories are: CAT C262T (CT vs CC; TT vs CC); PON1 A575G (AG vs AA; GG vs AA); PON1 C108T (CT vs CC; TT vs CC) and MPO G463A (AG vs AA; GG vs AA). Step 1 Contrast 2: Contrast reference categories are: CAT C262T (CT vs TT); PON1 A575G (AG vs GG); PON1 C108T (CT vs TT) and MPO G463A (AG vs GG). Bold italic font denotes p-values significant for less than 0.05. AD: Anxiety and depressed mood; B: Unstandardized coefficient; df: Degree of freedom; Exp(B): Odds ratio; SE.: Standard error.												

In particular, the condition of heterozygosity of the A-575G polymorphism of *PON1* (rs622) and the presence of the T allele in the genotype of the C-108T polymorphism (rs705379) are more frequently correlated with the manifestation of anxiety and depression in subjects with MCS. Why the heterozygous condition of the rs622 can be more at risk for anxiety/depression in MCS should be the subject of further studies. A possible hypothesis is that phenomena due to linkage disequilibrium may be involved. Another aspect concerns the protective role of the C allele in the rs705379 genotype on the manifestation of anxiety and depression in MCS, which is also worthy of further scientific studies.

A deficit in detoxifying OPs has been hypothesized in the etiopathogenesis of MCS; this has been confirmed by data showing an association between *PON1* polymorphisms, MCS [48], and outcome of OP poisoning [94].

Low *PON1* activity in patients with anxiety correlated with high levels of lipid peroxidation, oxidative stress and hydroperoxide lipids, and has been involved in the etiopathogenesis of generalized anxiety [43]. Cholinergic hyperactivation has been implicated in the pathophysiology of anxiety and depression, and response to antidepressant treatment [95]. OPs act as cholinesterase inhibitors and subacute intoxication with OP insecticides induce anxiogenic effects [96]. *PON1* protects from oxidative stress by detoxifying OPs that are potential cholinesterase inhibitors in the peripheral and CNS [97,98]. When the enzyme is inhibited, it may not hydrolyze acetylcholine, leading to the accumulation of the neurotransmitter, which would, in turn, increase macrophage release of proinflammatory cytokines involved in anxious mechanisms [99]. The toxic effects involve parasympathetic, sympathetic, motor and CNSs [100]. These data are in-line with inverse correlations between acetylcholinesterase and anxiety, as well as with *PON1* activity, whereby subjects with low *PON1* activity may be at greater risk of severe state anxiety [43]. Furthermore, polymorphisms of *PON1* and *ACHE* genes, which are adjacent in the locus 7q21-22 and influence reciprocal expression in an allelic dependent manner [97] may play a role in susceptibility to chronic organophosphorus intoxication [100].

Recent studies showed the role of neuroinflammation in the pathogenesis of mood and anxiety disorders [101–105]. Neuroinflammation and oxidative stress have also been correlated with anxiety disorders, and sleep deprivation and anxiety [106–108]. Oxidative stress and diminished capacity in detoxifying organophosphates have been involved in the pathophysiology of MCS [36]. This evidence is in line with the involvement of oxidative stress in the pathophysiology of mental disorders, based on the vulnerability of the CNS to free radicals. These could damage the structure of the neuronal cell, reacting with the membrane lipids, proteins and nucleic acids [106,109–111].

Furthermore, limbic hyperactivation has been correlated to sensitization caused by neurogenic inflammation [112]. Another important aspect regards the odor processing in MCS patients. Some neuroimaging studies showed that exposure to odorants in MCS correlated to dysfunctions in the hippocampus, amygdala and thalamus [113]. MCS patients also showed hyperactivation of the anterior cingulate cortex and cuneus-precuneus related to exposure to odors, but without signs of neuronal sensitization [114].

The functionality of the frontal cortex in MCS subjects must be further investigated, as there is evidence of both increased and reduced metabolism during exposure to odors. MCS patients showed increased metabolism than controls in the prefrontal cortex during olfactory stimulation, and for about 20–30 s in the orbitofrontal cortex after stimulation, at the recognition threshold or usually perceived level [115]. These functional neural changes have been related to cognitive and memory processing adaptations during past exposure to unsafe chemicals [115]. Another study showed frontal cortex hypoactivation, with possible active recruitment of the left inferior temporal cortex during olfactory stimulation [116].

These data suggest that patients with MCS process odors differently from the healthy population, pointing to the need for further investigation on the relationships between anxiety-depressive symptoms, oxidative stress and odor processing in MCS.

Further investigations are needed with the aim of clarifying the factors related to the expression levels and activity of the paraoxonase enzyme, which is influenced not only by its related genetic polymorphisms but also by environmental factors and/or substances that can increase or decrease its expression and consequently its activity. In fact, it has been shown that cigarette smoking, a high-fat diet and several drugs, including antidepressants, anti-hypertensives, antiepileptics, cardiovascular, antineoplastics, some quinones, benzenesulfonamide derivatives, usnic and carnolic acids, and sulfonamides can decrease the activity of paraoxonase [52–60], which in turn can influence the levels of atherosclerosis and risk of the onset of cardiovascular, neurological and psychiatric diseases [20,51,56].

This evidence showed that the regulation of paraoxonase is very complex, and it is still necessary to shed light on how these factors interact with each other to define its activity levels. This points to the need of further studies aimed to research other mechanisms underlying the *PON1* activity and the pathophysiological correlates that can be associated with different organic and psychiatric conditions, including atherosclerosis, hypertension, other cardiovascular illnesses and anxiety, and depression.

Limitations

The results of this study must be considered with caution because of their preliminary nature and considering the following limitations. First, the sample size we obtained was not large enough to yield definite results, especially in the light of the fact that peculiarity of the population. Second, results are not corrected for multiple comparisons; however, this was related to the exploratory nature of the study. Third, the lack of use of assessment scales for anxiety and depression constitutes a significant limitation, for which the results of this study should be taken with

caution and need to be replicated. However, the compilation of a checklist by a physician allowed an adequate categorical assessment of symptoms. Furthermore, the lack of a control sample from the general population made our results not applicable to other than this population, the existence of which has still to gain consensus.

Future perspective

Due to the serious state of suffering of some affected subjects, understanding the etiopathogenesis of MCS and finding optimal therapy represents an invariable challenge for medicine. In the next few years, new studies that include neuroimaging data and correlations with psychological and psychopathological dimensions are necessary to understand the nature of this pathology. A growing body of evidence seems to demonstrate the close link between cellular detoxification, oxidative stress, psychiatric disorders and MCS; then new therapies could be developed to prevent brain damage from free radicals, such as the use of antioxidant agents, anti-inflammatory drugs, dietary recommendations and nutritional interventions [117,118]. Oxidation-reduction mechanisms could be a novel target for pharmacological intervention in MCS, also considering that pharmacogenomics will support the choice of targeted therapies in the field of precision medicine [119]. The recognition of specific polymorphisms that may be related to the efficacy of the pharmacological response may direct the clinician in the best choice of treatments that can be undertaken in this category of patients, also for the treatment of anxiety-depressive symptoms.

Conclusion

In this study, we showed *PONI* polymorphisms to predict anxiety and depressive symptoms in an MCS population. The *PONI* AG versus GG genotype of the A575G polymorphism increased the likelihood of reporting depressive or anxiety symptoms by more than ten-times, and *PONI* C-108T also appears to be significantly involved in the manifestation of comorbid generalized anxiety and depressed mood in MCS. These results apply to an MCS sample and have to be tested in the general and other populations.

Author contributions

A Mosca, A Del Casale, M Borro and GD Kotzalidis wrote the first draft of the manuscript and performed the statistical analyses. M Simmaco, M Borro, G Gentile and LM Pomes performed laboratory analyses, sample inclusion, and created the database. A Del Casale, A Mosca, A Padovano, F Fiaschè, V Pinzone, C Rapinesi, T Zoppi, R Brugnoli and G Sani did the literature search. GD Kotzalidis, P Girardi, S Ferracuti, M Simmaco and M Pompili supervised the study. Each author participated in manuscript writing and approved the final version.

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Ethical conduct of research

The authors state that they have obtained appropriate institutional review board approval or have followed the principles outlined in the Declaration of Helsinki for all human or animal experimental investigations. In addition, for investigations involving human subjects, informed consent has been obtained from the participants involved.

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