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Corrigendum to “Genetic interaction in the association between oxidative stress and diabetes in the Spanish population” [Free Radic. Biol. Med. 205 (2023) 62–68]

Rebeca Melero^a, Maria Elena Quiroz-Rodríguez^a, Francisco Lara-Hernández^a, Josep Redón^{b,c}, Guillermo Sáez^{d,e}, Laisa S. Briongos-Figuero^f, Jessica Abadía-Otero^f, Juan Carlos Martín-Escudero^{f,g}, F. Javier Chaves^{a,h,*}, Guillermo Ayala^{i,1}, Ana-Bárbara García-García^{a,h,1}

^a Genomics and Diabetes Unit, INCLIVA Biomedical Research Institute, 46010, Valencia, Spain

^b Cardiometabolic Renal Risk Research Group, INCLIVA Biomedical Research Institute, University of Valencia, 46010, Valencia, Spain

^c CIBEROBN, ISCIII, 28029, Madrid, Spain

^d Department of Biochemistry and Molecular Biology, Faculty of Medicine and Odontology, University of Valencia, 46010, Valencia, Spain

^e Service of Clinical Analysis, University Hospital Dr. Peset-FISABIO, Spain

^f Department of Internal Medicine, Rio Hortega University Hospital, 47012, Valladolid, Spain

^g Department of Medicine, Faculty of Medicine, University of Valladolid, 47002, Valladolid, Spain

^h CIBERDEM, ISCIII, 28029, Madrid, Spain

ⁱ Department of Statistics and Operation Research, University of Valencia, 46100, Burjassot, Valencia, Spain

The authors regret of a typographical error in Table 5, that shows the significant result of three logistic-regression models. This table shows “>” sign for all p-value and adjusted p-value. This sign is an error and it should be “<”. However, showed results do not change because all

considerations have been made taking into account the correct “<” sign. The authors would like to apologise for any inconvenience caused. Here Table 5 with corrected sign is shown:

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* Corresponding author. Genomics and Diabetes Unit, INCLIVA Biomedical Research Institute, Menendez Pelayo 4acc, Valencia, 46010, Spain.

E-mail address: felipe.chaves@uv.es (F.J. Chaves).

¹ Both authors contributed equally to this work.

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Table 5

Significant result of three logistic regression model: MDA, 8-Oxo-dG and GSH/GSSG (for each model) and haplotypes as predictors and T2D as response.

Term	Haplotype	Gene	Coefficient	p-value	Adjusted p-value	
MDA	Geno_d.C.8	SP2	51.8	<10 ⁻¹⁶	<10 ⁻¹⁶	
	Geno_d.C.rare	SP2	0.892	<10 ⁻¹⁶	<10 ⁻¹⁶	
	Geno_d.C.rare	IL18R1	-11.7	<10 ⁻¹⁶	<10 ⁻¹⁶	
	Geno_d.C.rare	EIF2AK2	-14.8	<10 ⁻¹⁶	<10 ⁻¹⁶	
	Geno_d.C.18	TXNRD2	-14.8	<10 ⁻¹⁶	<10 ⁻¹⁶	
	Geno_d.C.5	PPARA	50.6	<10 ⁻¹⁶	<10 ⁻¹⁶	
	Geno_d.C.7	PPARA	776	<10 ⁻¹⁶	<10 ⁻¹⁶	
	Geno_d.C.12	PPARA	-25.3	<10 ⁻¹⁶	<10 ⁻¹⁶	
	Geno_d.C.35	PPARA	-8.64	<10 ⁻¹⁶	<10 ⁻¹⁶	
	Geno_d.C.rare	NDUFS2	8.56	<10 ⁻¹⁶	<10 ⁻¹⁶	
	8-Oxo-dG	Geno_d.C.rare	HIF1A	-391	<10 ⁻¹⁶	<10 ⁻¹⁶
		Geno_d.C.rare	IL18R1	-0.344	<10 ⁻¹⁶	<10 ⁻¹⁶
Geno_d.C.5		PPARA	-17.9	<10 ⁻¹⁶	<10 ⁻¹⁶	
Geno_d.C.8		PPARA	-13.4	<10 ⁻¹⁶	<10 ⁻¹⁶	
Geno_d.C.35		PPARA	-1.77e+03	<10 ⁻¹⁶	<10 ⁻¹⁶	
Geno_d.C.rare		NDUFS2	8.56	<10 ⁻¹⁶	<10 ⁻¹⁶	
Geno_d.C.3		ERN1	0.438	0.00278	0.0183	
GSSG/GSH		Geno_d.C.8	SP2	-0.58	<10 ⁻¹⁶	<10 ⁻¹⁶
		Geno_d.C.rare	SP2	-2.35	<10 ⁻¹⁶	<10 ⁻¹⁶
		Geno_d.C.rare	HIF1A	-0.00846	<10 ⁻¹⁶	<10 ⁻¹⁶
		Geno_d.C.rare	IL18R1	0.977	<10 ⁻¹⁶	<10 ⁻¹⁶
		Geno_d.C.rare	EIF2AK2	139	<10 ⁻¹⁶	<10 ⁻¹⁶
	Geno_d.C.5	PPARA	-1.08	<10 ⁻¹⁶	<10 ⁻¹⁶	
	Geno_d.C.7	PPARA	-4.45	5.88e-11	4.17e-10	
	Geno_d.C.12	PPARA	-0.755	<10 ⁻¹⁶	<10 ⁻¹⁶	