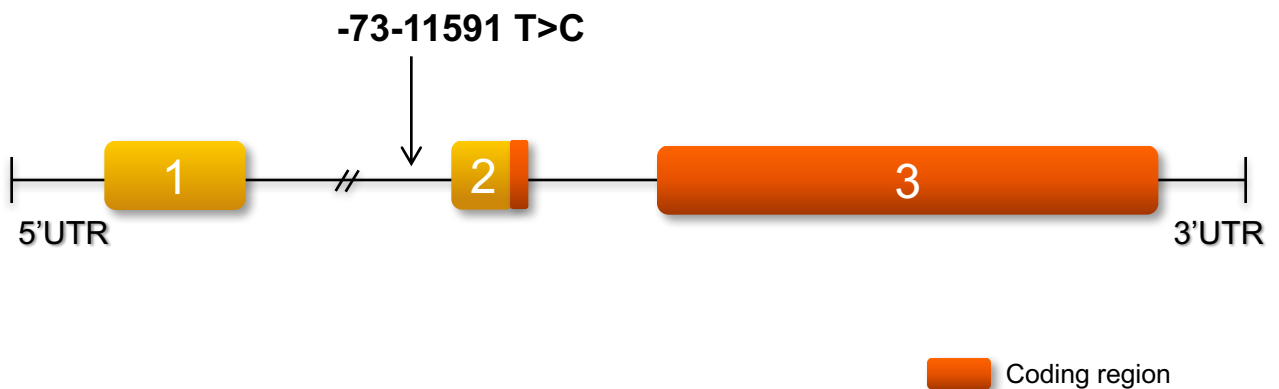


Association of a variant in the gene encoding for ERV1/ChemR23 with reduced inflammation in visceral adipose tissue from morbidly obese individuals

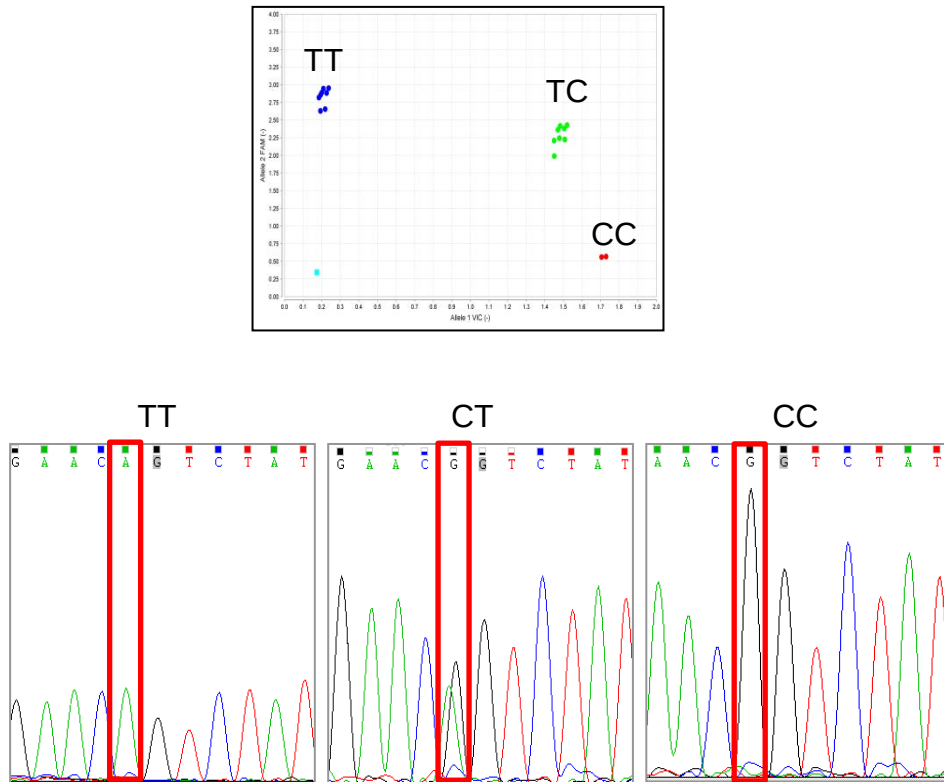
Cristina López-Vicario^{1,3}, Bibiana Rius¹, José Alcaraz-Quiles¹, Ana González-Pérez¹, Ana Isabel Martínez-Puchol¹, Mireia Casulleras¹, Marta Duran-Güell¹, Ainitze Ibarzabal², Ricard Corcelles², Andrés Laguna-Fernández⁵, Magnus Back⁵, Esther Títos^{1,3}, Joan Clària^{1,3,4}

¹Department of Biochemistry and Molecular Genetics and ²Department of Gastrointestinal Surgery, Hospital Clínic, IDIBAPS; ³CIBERehd and ⁴Department of Biomedical Sciences, University of Barcelona, Barcelona, Spain; ⁵Centre for Molecular Medicine, Department of Medicine, Karolinska Institutet and Department of Cardiology, Karolinska University Hospital, Stockholm, Sweden.

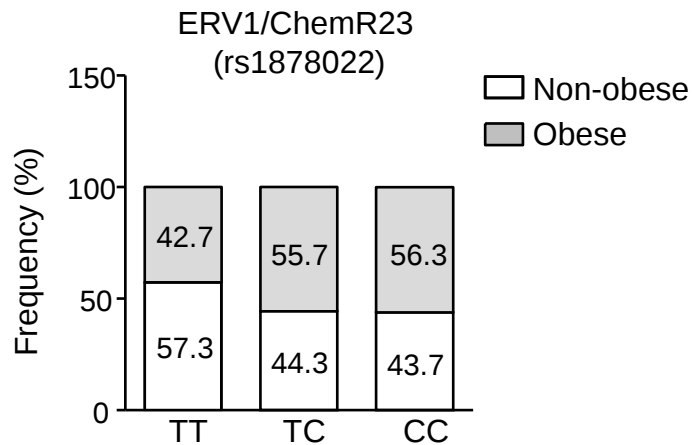


Supplementary Figure 1. Structure of the ERV1/ChemR23 gene. The ERV1/ChemR23 gene contains 3 exons and 2 introns. The SNP rs1878022 (nucleotide change from T to C) is located in the first intron of the gene. Transcription starts in exon 1. Coding region starts in the last codon of exon 2 and covers exon 3.

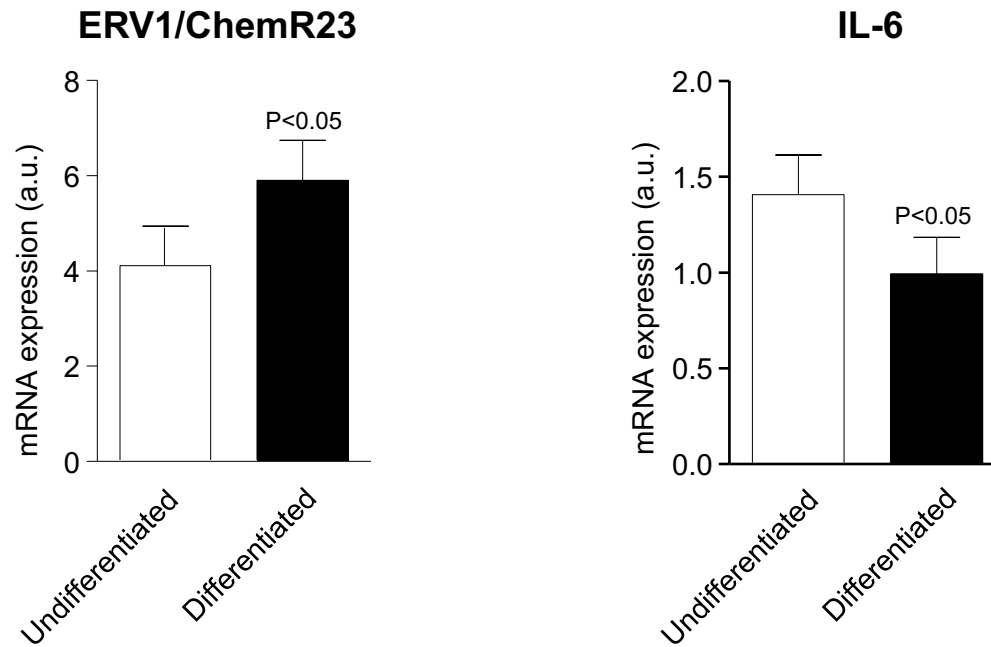
A



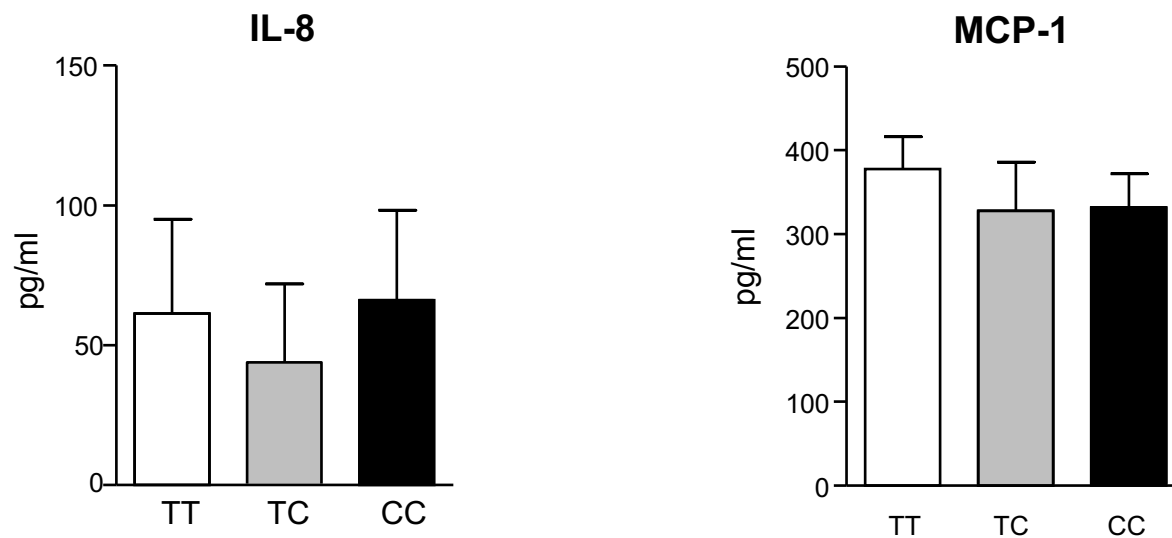
B



Supplementary Figure 2. (A) Representative data of the analysis of the ERV1/ChemR23 rs1878022 SNP by the TaqMan SNP genotyping assay (top) and Sanger sequencing (bottom). (B) Frequency of the three ERV1/ChemR23 rs1878022 TT (n=82), TC (n=70) and CC (n=48) genotypes in obese and non-obese individuals.



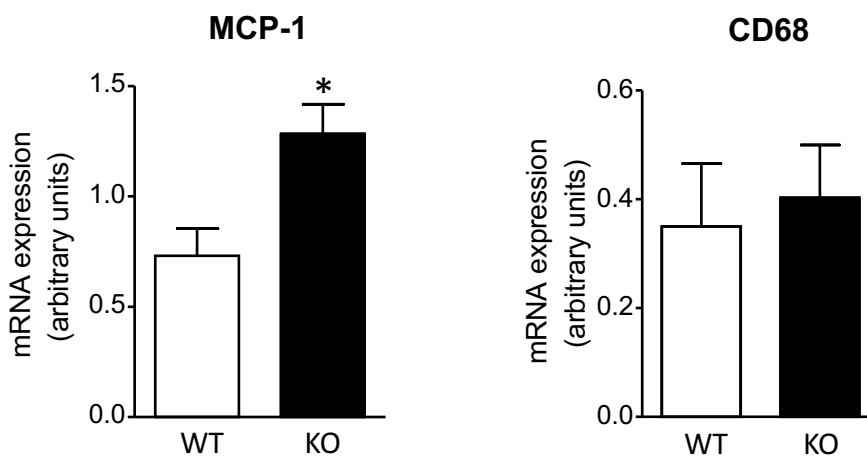
Supplementary Figure 3. ERV1/ChemR23 and IL-6 mRNA expression in undifferentiated and differentiated 3T3-L1 adipocytes. The data represent the mean $\pm$ SEM from 3 different experiments in duplicate.



Supplementary Figure 4. Circulating levels of IL-8 and MCP-1 in obese individuals as determined by Luminex assay. The data represent the mean $\pm$ SEM of the individuals with TT (n=16), TC (n=16) and CC (n=17) genotypes.

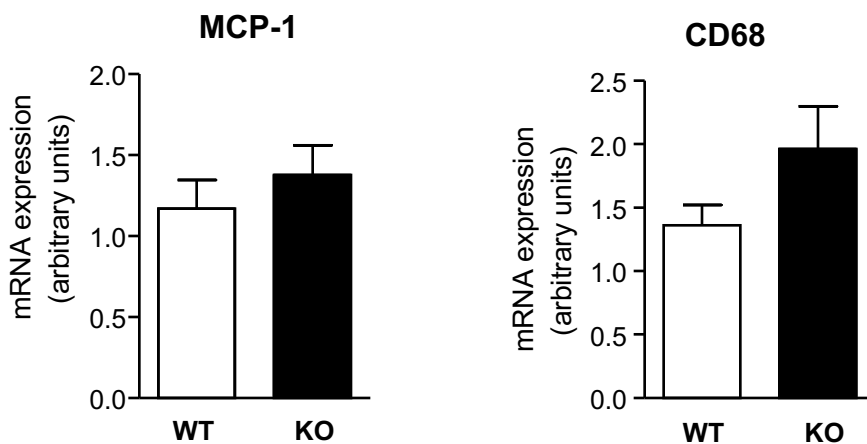
A

Adipose tissue

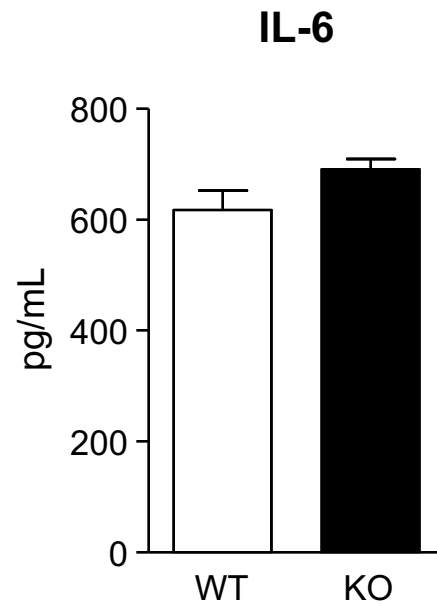


B

Liver



Supplementary Figure 5. RT-PCR results of MCP-1 and CD68 inflammatory markers in (A) adipose tissue and (B) liver from WT and KO mice. The data represent the mean $\pm$ SEM of WT (n=12) and KO (n=12) mice. *, $p < 0.05$ versus WT.



Supplementary Figure 6. IL-6 levels in livers from WT and KO mice. The data represent the mean $\pm$ SEM of WT (n=12) and KO (n=12) mice.

Supplementary Table 1. Identification, alleles, location and references of the SNPs analyzed in the study.

SNP ID.	Gene	Alleles	Chromosome	Position	References
rs1143634	IL-1 β	G/A	2	112832763-112832863	Manica-Cattani MF, et al. <i>Mol Cell Endocrinol</i> 2010. Carter KW et al. <i>Hum Genet</i> 2008.
rs1800795	IL-6	C/G	7	22726976-22727076	Yu Z et al. <i>Obesity</i> 2012. Carulli L et al. <i>Dig. Liver Dis</i> 2009.
rs8069645	STAT3	A/G	17	42342834-42342934	Phillips CM et al. <i>J Nutr.</i> 2009.
rs7849191	JAK2	C/T	9	4988711-4988811	Pneas-Steinhardt A et al. <i>BMC Med Genet</i> 2011.
rs1800871	IL-10	A/G	1	206773239-206773339	Scarpelli D et al. <i>Diabetes</i> 2006.
rs8064821	SOCS3	C/A	17	78361260-78361360	Jamshidi Y et al. <i>Diabetologia</i> 2006. Reid-Lombardo KM. <i>Ann Surg</i> 2013.
rs1878022	ERV1/ChemR23*	C/T	12	108305205-108305305	Wu X et al. <i>J Natl Cancer Inst.</i> 2011. Alcaraz-Quiles et al. <i>Hepatology</i> 2017.

**The ERV1/ChemR23 gene is also known as CMKLRL1.*

Supplementary Table 2: Phenotypic characteristics of ERV1/ChemR23 knockout (ERV1/ChemR23^{-/-}) mice compared to their wild-type (WT) littermates.

	WT (n= 12)	ChemR23^{-/-} (n= 12)
Body weight (g)	22.1 ± 1.2	22.3 ± 1.9
Adipose tissue/body weight	0.01 ± 0.002	0.01 ± 0.001
Liver/body weight	0.04 ± 0.005	0.05 ± 0.002
Total Cholesterol (mg/dL)	76. 6 ± 5.5	74.33 ± 5.8
HDL-c (mg/dL)	22.3 ± 3.7	23.6 ± 3.1
LDL-c (mg/dL)	39.8 ± 4.3	40.9 ± 3.6
Triglycerides (mg/dL)	72.57 ± 9.3	65.0 ± 6.9

*HDL-c: high-density lipoprotein cholesterol; LDL-c: low-density lipoprotein cholesterol.
Results are expressed as mean ± SEM. *P<0.05 versus WT.*