



Supplementary material

Natural extracts to augment energy expenditure as a complementary approach to tackle obesity and associated metabolic alterations.

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* Equal contribution

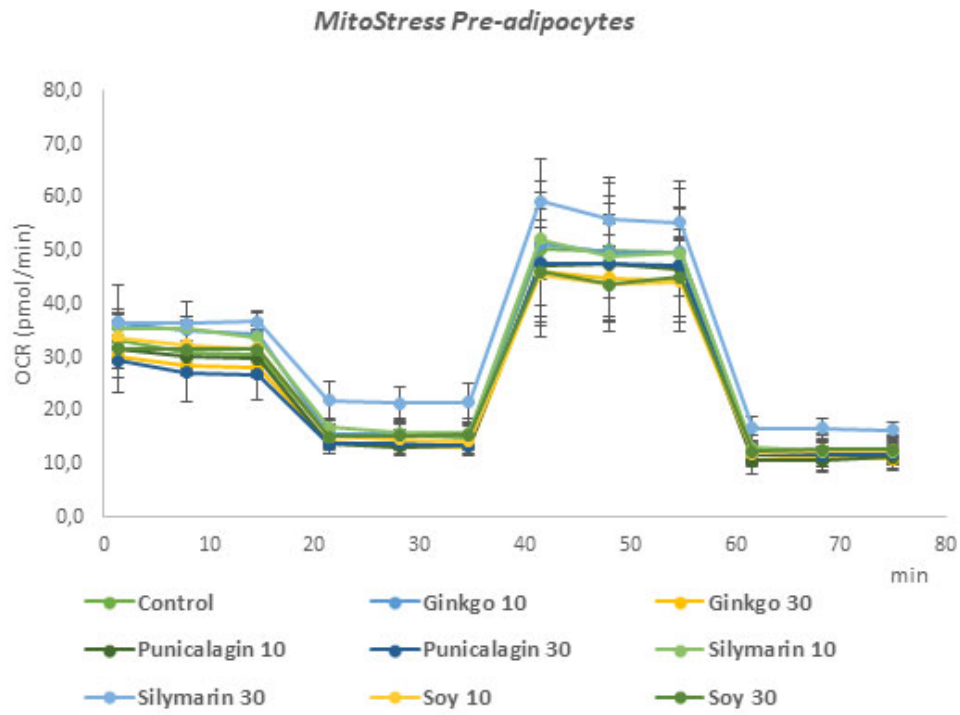


Figure S1. Effect of the extracts on mitochondrial oxidative phosphorylation of humannon-differentiated pre-adipocytes.

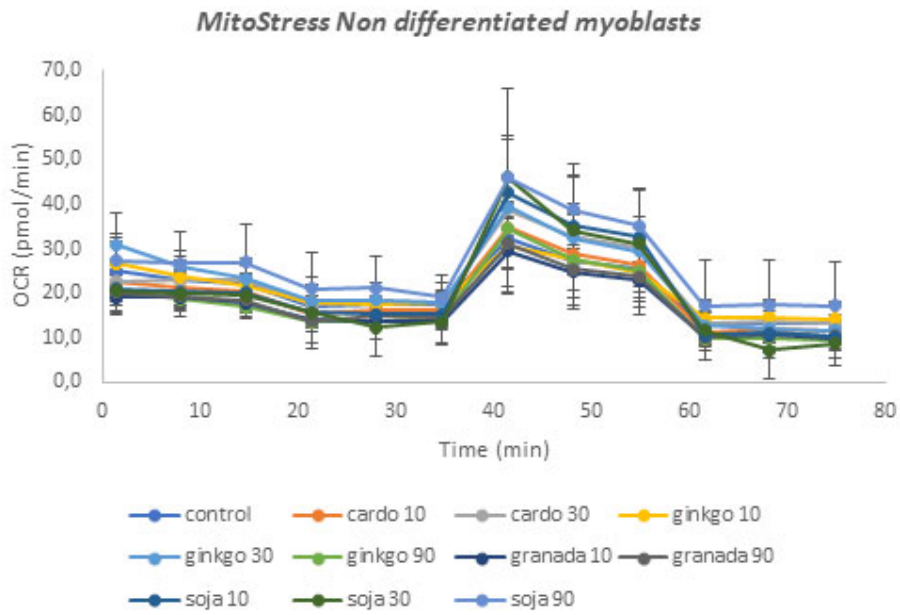


Figure S2. Effect of the extracts on mitochondrial oxidative phosphorylation of human non differentiated myoblasts.

Table S1. List of Taqman probes used in the study.

GENES	Thermofisher probe's code
<i>ADRB1</i>	Hs02330048_s1
<i>AGT</i>	Hs01586213_m1
<i>LEPR</i>	Hs00174497_m1
<i>LPL</i>	Hs00173425_m1
<i>MTHFR</i>	Hs01114487_m1
<i>PPARγ</i>	Hs01115513_m1
<i>TCF7L2</i>	Hs01009044_m1
<i>UCP2</i>	Hs01075227_m1
<i>ACE</i>	Hs00174179_m1
<i>CLOCK</i>	Hs00231857_m1
<i>COL5A1</i>	Hs00609133_m1
<i>HFE</i>	Hs05045803_s1
<i>HIF1A</i>	Hs00153153_m1
<i>PPARA</i>	Hs00947536_m1
<i>PPARGC1A</i>	Hs00173304_m1
<i>AHR</i>	Hs00169233_m1
<i>BNC2</i>	Hs00417700_m1
<i>DLGAP1</i>	Hs00191052_m1
<i>COMT</i>	Hs00241349_m1
<i>IL4-R</i>	Hs00965056_m1
<i>HLA-DQA1</i>	Hs03007426_mH
<i>MCM6</i>	Hs00962418_m1
<i>ABCA1</i>	Hs01059137_m1
<i>FTO</i>	Hs01057139_g1
<i>MC4R</i>	Hs00271877_s1
<i>TMEM18</i>	Hs00894216_m1
<i>GNPDA2</i>	Hs00385682_m1
<i>KCTD15</i>	Hs00225337_m1
<i>ETV5</i>	Hs00927557_m1
<i>NPY</i>	Hs00173470_m1
<i>APOA5</i>	Hs00364830_m1
<i>GCKR</i>	Hs00386984_m1
<i>CD36</i>	Hs00354519_m1
<i>B2M</i>	Hs00187842_m1
<i>GAPDH</i>	Hs02786624_g1

Table S2. List of primers used in the study.

GEN	FORWARD PRIMER	bp	REVERSE PRIMER	bp	SPECIE
<i>MYOG</i>	TCAACCAGGAGGAGCGTGA	19	TCTGTAGGGTCAGCCGTGAG	20	human
<i>MEF2</i>	AACAAAGCCCTCAGCAGGT	19	ACTTGCACAGCTCAGTTCCC	20	human
<i>UCP1</i>	ACAGAAGGGCGGATGAAACT	20	CTTGCTTCTAAACTAGGTGCTG	23	human
<i>TFAM</i>	ATGGCGTTTCTCCGAAGCAT	20	TCCGCCCTATAAGCATCTTGA	21	human
<i>PPARG</i>	ACCAAAGTGCAATCAAAGTGGA	22	ATGAGGGAGTTGGAAGGCTCT	21	human
<i>UCP3</i>	AAGGTCCGATTTCCAGGCCAG	20	GCGATGGTTCTGTAGGCGTC	20	human
<i>C/EBPα</i>	TATAGGCTGGGCTTCCCCTT	20	AGCTTCTGGTGTGACTCGG	20	human
<i>ATP2A1</i>	AAACCACGGAGGAATGTTTGG	21	AGCTCATTGAGGCCGTATTTT	21	human
<i>MYH2</i>	AGAAACTTCGCATGGACCTAGA	22	CCAAGTGCCTGTTTCATCTTCA	21	human
<i>FABP4</i>	ACTGGGCCAGGAATTTGACG	20	CTCGTGAAGTGACGCCCTT	19	human
<i>Cidea</i>	TTATGGGATCACAGACTAAGCGA	23	TGCTCCTGTCATGGTTGGAGA	21	human
<i>Ckmt2</i>	GACCCCGTCATCAAATAAGAC	22	AGCACGTAATGCTCGTCGAAC	21	human
<i>Lep</i>	TGCCTCCAGAAACGTGATCC	21	CTCTGTGGAGTAGCCTGAAGC	21	human
<i>PRDM16</i>	CGATGCCGACTTTTGGGAAG	20	GTGGAGAGGAGTGTCTTCGG	20	human
<i>PGC1α</i>	ATTGGAGCCCATGGATGAA	20	GCGGCTGTTACTCTCTCTCC	20	human
<i>CPT1A</i>	CTCAGTGGGAGCGGATGTTT	20	TGCTGTCTCTCATGTGCTGG	20	human
<i>LDLR</i>	CAGTACCCCTCGAGACAGA	20	CACTGTCCGAAGCCTGTTCT	20	human
<i>IRS1</i>	GCAACCAGAGTGCCAAAGTG	20	GCACTGGGTGTTGAGGAGAA	20	human
<i>FASN</i>	TATGAAGCCATCGTGGACGG	20	GAAGAAGGAGAGCCGTTGG	20	human
<i>UCP2</i>	GACCATTGCCGAGAGGAAG	20	GAAGTGGCAAGGGAGGTCAT	20	human
<i>FNDC5</i>	CGTGGTCTGTTTCATGTGGG	20	GGTCTTGCCCTCACCTTGCT	20	human
<i>BDNF</i>	TAACGGCGGCAGACAAAAGA	21	TGCACTTGGTCTCGTAGAAGTAT	23	human
<i>IL6</i>	CCTGAACCTTCAAAGATGGC	21	TTCACCAGGCAAGTCTCTCA	21	human
<i>SIRT1</i>	TGTGTCATAGGTTAGGTGGTGA	22	AGCCAATTCTTTTGTGTTTCGTG	23	human
<i>B2M</i>	GATGAGTATGCCTGCCGTGT	20	TGCGGCATCTTCAAACCTCC	20	human