

Supplementary materials

Table S1 Sources of diet components

Component (product name)	Source (Company, City, Country)
Maize starch (Maisita – Native Maisstärke)	Agrana Beteiligungs-AG, Gmünd, Austria
Casein (Nährkasein, Säurekasein 110 Mesh)	Meggle, Wasserburg, Germany
Saccharose (Puderzucker)	Suedzucker, Mannheim, Germany
Soybean oil (Heidenreich – Soja 3-Plus)	Karl Heidenreich GmbH, Mannheim, Germany
Cellulose (Arbocel – Rohfaserkonzentrat, Typ R)	Rettenmaier & Söhne GmbH + Co KG, Rosenberg, Germany
Mineral mix ¹	Self-made
Vitamin mix ²	Self-made
L-Cystein	Sigma Aldrich, St. Louis, USA
Supplemental NA	Lonza, Basel, Switzerland

¹Components mineral mix

MgSO ₄ x 7H ₂ O	Roth, Karlsruhe, Germany
CaCO ₃	Bergmann, Kasendorf, Germany
NaCl	Roth, Karlsruhe, Germany
FeSO ₄ x 7H ₂ O	Merck, Darmstadt, Germany
ZnSO ₄ x 7H ₂ O	Grüssing, Vilsum, Germany
MnSO ₄ x H ₂ O	Roth, Karlsruhe, Germany
CuSO ₄ x 5H ₂ O	Roth, Karlsruhe, Germany
KI	Sigma Aldrich, St. Louis, USA
NaFl	Merck, Darmstadt, Germany
CrK(SO ₄) ₂ x 12H ₂ O	Merck, Darmstadt, Germany
LiCl	Sigma Aldrich, St. Louis, USA
Na ₂ MoO ₄ x 2H ₂ O	Acros organics, New Jersey, USA
Na ₂ SeO ₃ x 5H ₂ O	Merck, Darmstadt, Germany

KH ₂ PO ₄	Roth, Karlsruhe, Germany
² Components vitamin mix	
Choline chloride (50% S)	Bergophor, Kulmbach, Germany
Microvit A Promix 1000, 1.000.000 I.E./g	Bergophor, Kulmbach, Germany
Microvit B1 Promix Thiamine Mono, 100%	Bergophor, Kulmbach, Germany
Microvit B2 Supra 80, 80%	Bergophor, Kulmbach, Germany
Microvit B3 Promix Niacin 99.0%	Bergophor, Kulmbach, Germany
Microvit B5 Promix D-Calpan, 100%	Bergophor, Kulmbach, Germany
Microvit B6 Promix Pyridoxine, 98%	Bergophor, Kulmbach, Germany
Microvit H Promix Biotin, 2%	Bergophor, Kulmbach, Germany
Microvit B9 Supra 100, 100%	Bergophor, Kulmbach, Germany
Microvit B12 Promix 1000, 1.0%	Bergophor, Kulmbach, Germany
Microvit D3 Promix 500, 500.000 I.E./g	Bergophor, Kulmbach, Germany
Microvit E Promix 50, 500 I.E./g	Bergophor, Kulmbach, Germany
Microvit K3 Promix MNB 51%	Bergophor, Kulmbach, Germany

Table S2 Characteristics of gene specific primers used for qPCR

Gene symbol	Primer sequence (5'-3', forward, reverse)	Product size (bp)	NCBI GenBank accession number
<i>Reference genes</i>			
<i>Atp5b</i>	GCCAGAGACTATGCGGCGCA, CCCCCAAATGCTGGGCCACC	187	NM_016774
<i>Canx</i>	GTCCCCGGGAGGCTCGAGATAGAT, ACCTCCCCTGTTGGAAGTGGAGC	234	NM_007597
<i>Cycl</i>	CGCAATGGAAGCTGCCGGGA, CGCAATGGAAGCTGCCGGGA	99	NM_025567
<i>Target genes</i>			
<i>Acadl</i>	CACAGGCCCTGGCTTCAGCC, ATGCCATGGGCAGGCGATCG	303	NM_007381
<i>Acadm</i>	AGGTTTCAAGATCGCAATGG, CTCCTTGGTGCTCCACTAGC	152	NM_007382
<i>Atf4</i>	TTCCTCGAATCCAGCAAAGCCC, CTCCAACATCCAATCTGTCCGG	286	NM_009716
<i>Casp12</i>	GATGGAGAAGGAGGGACGAACAC, CACCAGGAATGTGCTGTCTGAGG	235	NM_009808
<i>Clpp</i>	CACACCAAGCAGAGCCTACA, TCCAAGATGCCAAACTCTTG		NM_017393
<i>Cox4i1</i>	CTTCGAGCACATGGGAGTGT, ATCAGAACGAGCGCAGTGAA	296	NM_009941
<i>Ddit3</i>	AGCCAGAATAACAGCCGGAACC, GGGACTCAGCTGCCATGACTG	117	NM_007837
<i>Dnajc3</i>	CTGCATGGACACAGGTCCCAG, AGGCTGGACACCCCTACCTC	217	NM_008929
<i>Gadd34</i>	GCGGCTCAGATTGTTCAAAGCC, AGATGGGTTTCTAAGGCGTGCC	271	NM_008654
<i>Gapdh</i>	CAACGACCCCTTCATTGAC, AGACTCCACGACATACTCAG	196	NM_001289726
<i>Gpx1</i>	GTCCACCGTGTATGCCTTCT, TCTGCAGATCGTTCATCTCG	152	NM_008160
<i>Gys1</i>	AGGGGATGAATGGGGTGACA, CATTTCGTGGAAGTGGGCAAC	384	NM_030678
<i>Herpud1</i>	CGCAGTTGGAGTGTGAGTCGC, ACCTTTGTGCTGGTTTCTGGC	216	NM_022331
<i>Hmox1</i>	GATTTGTCTGAGGCCTTGAAG, CTAAAGCCTTCTCTGGACAC	111	NM_010442

<i>Hspa5</i>	TCATCGGACGCACTTGAATGAC, CGCTGGGCATCATTGAAGTAAGC	250	NM_001163434
<i>Hsp10</i>	CTGACAGGCTCAATCTCTCCAC, AGGTGGCATTATGCTTCCAG	108	NM_008303
<i>Hsp90b1</i>	CCTGCTGACCTTCGGGTTTCG, CTGTCCTTGAGCCTTCTCGGC	98	NM_011631
<i>Ldha</i>	CTTGGCGCTCTACTTGCTGT, CTCATCCGCCAAGTCCTTCA	334	NM_001136069
<i>Pdia4</i>	CCGTTGACTATGATGGCTCCAGG, GGTCTGTCTGTTCGGTGGCG	285	NM_009787
<i>Pfkm</i>	CCAGTGCATCTTAACCGACCAT, TCGGGCACTTCCAATCACTG	309	NM_021514
<i>Phka1</i>	TGAGTGGAGACGGGGTCTTT, TCACGGTCAGCATTTTTGCG	404	NM_008832
<i>Ppargc1a</i>	GACCCTCCTCACACCAAAC, GCGACTGCGGTTGTGTATG	91	XM_006503779
<i>Pygm</i>	CTCCAGGACATCATCCGACG, TTCACTGTTACGTCCCAGGC	200	NM_011224
<i>Slc25A20</i>	TGGACACTGTTGCTGAGA, TTGGCCAAAGGTATCGAG	225	NM_020520
<i>Sod1</i>	GAGACCTGGGCAATGTGACT, GTTTACTGCGCAATCCCAAT	220	NM_011434
<i>Txnrd1</i>	GACTGGCAGCAGCTAAGGAGG, GCTTGTCCGAGCAAAGCTGC	166	NM_001042523
