

Supplementary Material

Supplementary Table T1

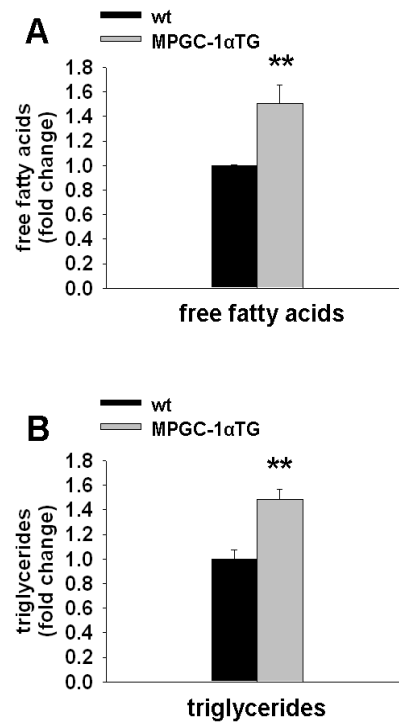
	FORWARD	REVERSE
<i>mTBP</i>	ATATAATCCCAAGCGATTTGC	GTCCGTGGCTCTCTTATTCTC
<i>hTBP</i>	GCCATAAGGCATCATTGGAC	AACAACAGCCTGCCACCTTA
<i>mGLUT1</i>	CGAGGGACAGCCGATGTG	TGCCGACCCTCTTCTTTCAT
<i>mGLUT4</i>	GATGAGAAACGGAAGTTGGAGAGA	GCACCACTGCGATGATCAGA
<i>mHK</i>	CCCTGCCACCAGACGAAA	GACTTGAAACCCTTAGTCCATGA
<i>mG6PDH</i>	AGGTGACCCTAAGCCGGAC	AGGTTTCTTTGGGTAGAAGACCA
<i>mRiboRed</i>	CTGCTCAGATCACCATGAAAGT	TGCTTCGTGGTCAAGGTCG
<i>mFAS</i>	CGGAACTTCAGGAAATGTCC	TCAGAGACGTGTCACTCCTGG
<i>hFAS</i>	TGTGGACATGGTCACGGAC	GGCATCAAACCTAGACAGGTC
<i>mFAT/CD36</i>	TGGCCTTACTTGGGATTGG	CCAGTGTATATGTAGGCTCATCCA
<i>mFABPpm</i>	AGCGGCTGACCAAGGAGTT	GACCCCTGCCACGGAGAT
<i>mFATP1</i>	GGCTCCTGGAGCAGGAACA	ACGGAAGTCCCAGAAACCAA
<i>mFATP3</i>	CAGCTCTACAGCCATGTTTCTGA	CAAAGATTCTGGAGCCTGAGA
<i>mFATP4</i>	GGCTTCCCTGGTGTACTATGGAT	ACGATGTTTCTGCTGAGTGGTA
<i>mFATP6</i>	GGCTTGAGGATGCCGCTTA	GTAATCTGGGCTCATGCTATGAAG
<i>mLPL</i>	GGGAGTTTGGCTCCAGAGTTT	TGTGTCTTCAGGGGTCCTTAG
<i>mDGAT</i>	TCCGTCCAGGGTGGTAGTG	TGAACAAGAATCTGCAGACGA
<i>mtGPAT</i>	ACAGTTGGCACAATAGACGTTT	CCTTCCATTTCACTGTTGCAGA
<i>mACS</i>	CTCACATTATATTGCTGCCTGT	TCTCTTTGCCATAGCGTTTTTCT
<i>hACS</i>	AAAAGCTAAGCCCACTTCAGAC	ACTTGGATACAGCATGGTCAAAT
<i>mLXR alpha</i>	GGGAGGAGTGTGTGCTGTGCTCAG	GAGCGCCTGTTACACTGTTGC
<i>hLXR alpha</i>	CATGGCACCAGATCCCCATAG	GGGTAGCTGTTAGCAAAGTCAA
<i>mLXR beta</i>	GGCCTGGACGATGCAGAGT	CGATCGGCTGAGAAGATGTTG
<i>mRXR alpha</i>	AACCCAGCTCACCAAATGACC	AACAGGACAATGGCTCGCAGG
<i>hRXR alpha</i>	TTCGCTAAGCTCTTGCTC	ATAAGGAAGGTGTCAATGGG
<i>mRXR beta</i>	GCCAAGCTGCTGTTACGTCTT	ACAGGTGCTCCAGACACTTGA
<i>hRXR beta</i>	GAAGCTCAGGAAACACTAC	TGCAGTCTTTGTTGTCCTC
<i>mRXR gamma</i>	CCGCTGCCAGTACTGTCG	ACCTGGTCTCCAAGGTGAG
<i>mSREBP1c</i>	GCATGCCATGGGCAAGTAC	CCACATAGATCTCTGCCAGTGTG
<i>mLipin</i>	CTCCGCTCCCGAGAGAAAAG	TCATGTGCAAATCCACGGACT
<i>hGLUT4</i>	CTCAGCAGCGAGTACTGG	CCCCAATGTTGACCCAAACTG
<i>hHK</i>	GAGCCACCCTACCCCTACT	ACCCAAAGCACACGGAAAGTT
<i>hG6PDH</i>	CAAACAGAGTGAGCCCTTCTTC	CTCATGCAGGACTCGTGAATG

Supplementary Table T1

Sequences of primers used for RT-PCR.

m, mouse-specific primers; h, human-specific primers.

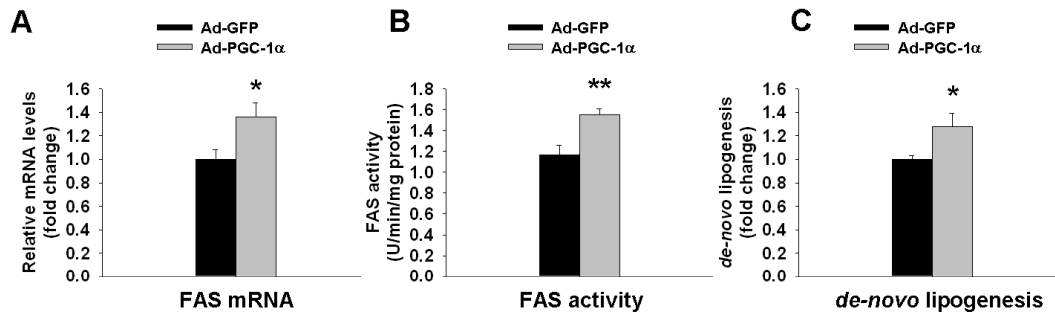
Supplementary Figure S1



Supplementary Figure S1. Incorporation of *de-novo* synthesized lipids into different lipid components.

(A and B) Tritium labeled free fatty acids (A) and triglycerides (B) in glycolytic muscle of MPGC-1α TG mice vs. controls. All values are means \pm SE. (n=5 per group). ** p<0.01.

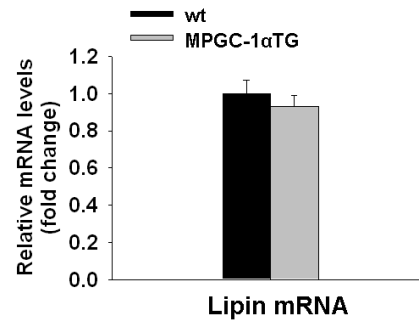
Supplementary Figure S2



Supplementary Figure S2. Enhanced lipid anabolism in isolated muscle cells overexpressing PGC-1 α .

(A-C) FAS gene expression (A), activity (B) and *de-novo* lipogenesis (C) in isolated muscle cells overexpressing either GFP (control), or PGC-1 α . All values are means \pm SE. (n=8 per group). * p<0.05; ** p<0.01.

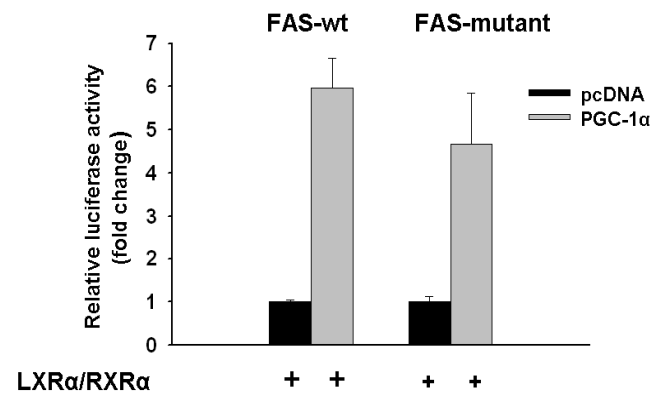
Supplementary Figure S3



Supplementary Figure S3. Lipin gene expression

Relative gene expression of lipin in EDL as measured by RT-PCR and expressed as fold change over controls.

Supplementary Figure S4

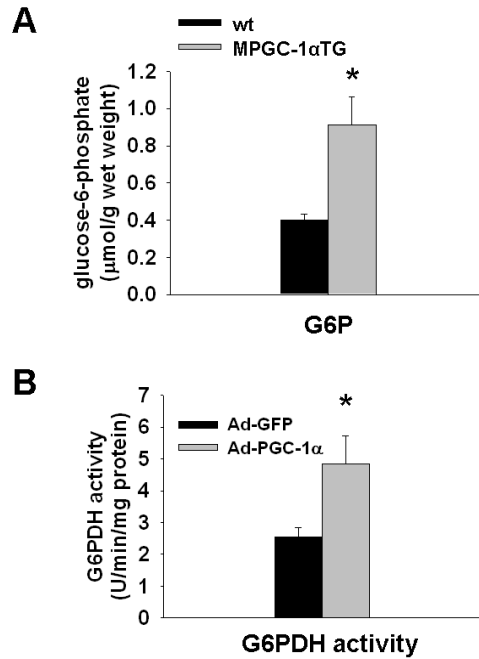


Supplementary Figure S4. Luciferase assay

Promoter activity of the FAS wild-type promoter (FAS-wt) and of a SREBP₁-binding mutant (FAS-mutant) following transfection of myoblasts with pcDNA (empty vector) (black bars) or PGC-1α (grey bars). LXRα/RXRα expression plasmids were cotransfected for all assays.

Values are expressed as means ±SE (n =6 per group)

Supplementary Figure S5

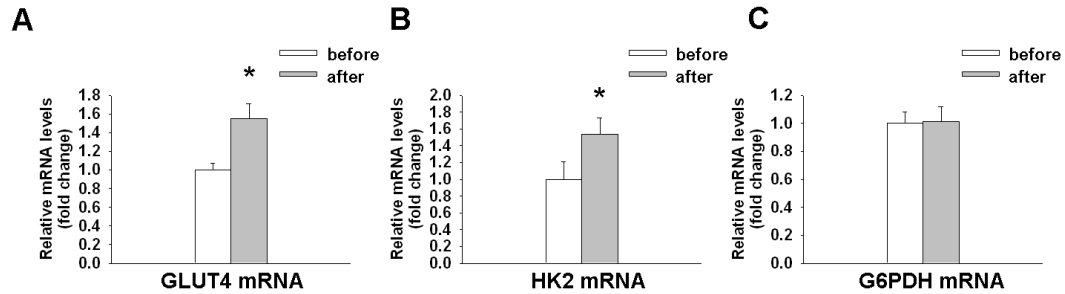


Supplementary Figure S5. Increased Glucose-6-phosphate levels in muscle and elevated G6PDH activity in isolated muscle cells overexpressing PGC-1 α .

(A) Glucose-6-phosphate (G6P) levels in glycolytic muscles of MPGC-1 α TG mice vs. controls. All values are means \pm SE. (n=5-6 per group). * p<0.05.

(B) G6PDH activity in isolated muscle cells overexpressing either GFP (control), or PGC-1 α . All values are means \pm SE. (n=6 per group). * p<0.05.

Supplementary Figure S6



Supplementary Figure S6. GLUT4, HK2 and G6PDH mRNA levels in endurance athletes pre- and post-exercise.

(A-C) Relative expression of GLUT4 (A), HK2 (B) and G6PDH (C) in human muscle biopsies pre- and post-exercise. Mean values are expressed as bars with SE (n=6). * p<0.05 as assessed by paired *t*-test.