

Paracrine cross-talk between skeletal muscle and macrophages in exercise by PGC-1 α -controlled BNP

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[#]This manuscript is dedicated to the memory of our coworker and friend Markus Beer (deceased on Aug 24, 2015).

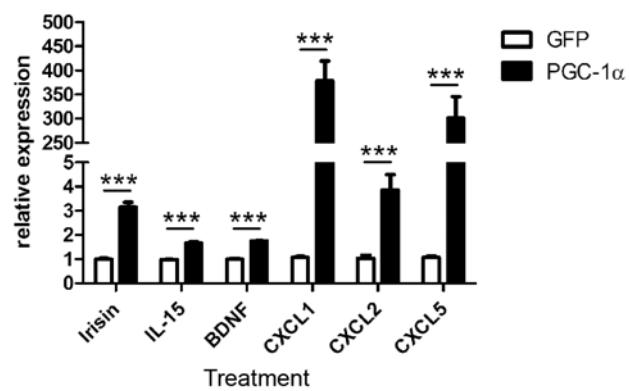
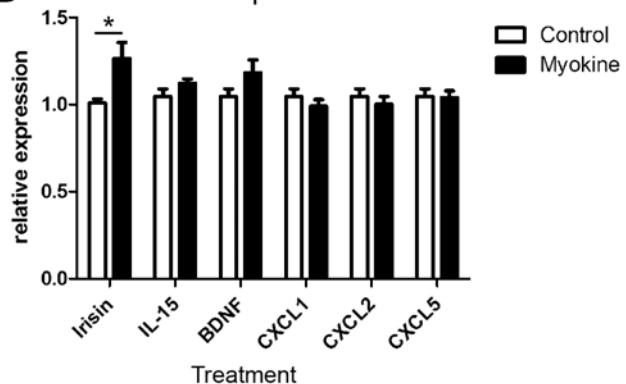
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Supplemental Information

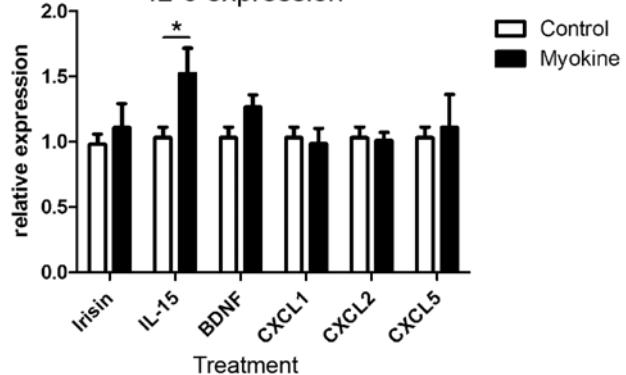
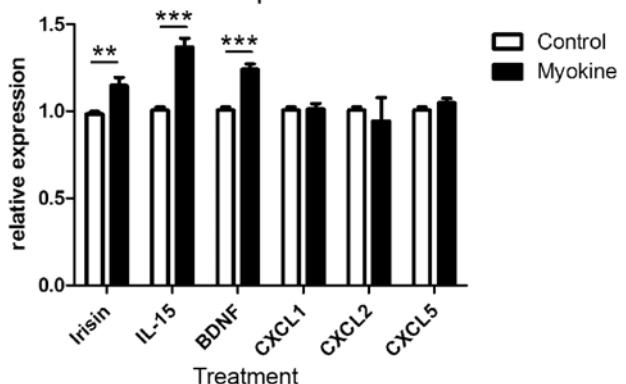
Supplemental Figures

Supplemental Figure S1 related to Figure 2. *Treatment of RAW macrophages with myokines induced by PGC-1 α only mildly increased the expression of pro- and anti-inflammatory cytokines.* **A.** mRNA expression levels of myokines were measured in C2C12 myotubes adenovirally overexpressing PGC-1 α or GFP **B, C.** RAW macrophages were treated for 4h with myokines known to be induced by PGC-1 α . Relative expression of pro- (**B**) and anti-inflammatory (**C**) cytokines was determined by RT-PCR. Values represent the mean of at least 3 independent experiments +SEM. * $P<0.05$; ** $P<0.01$; *** $P<0.001$; myokine versus control.

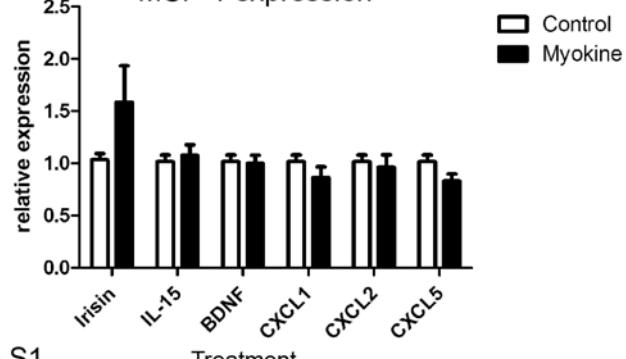
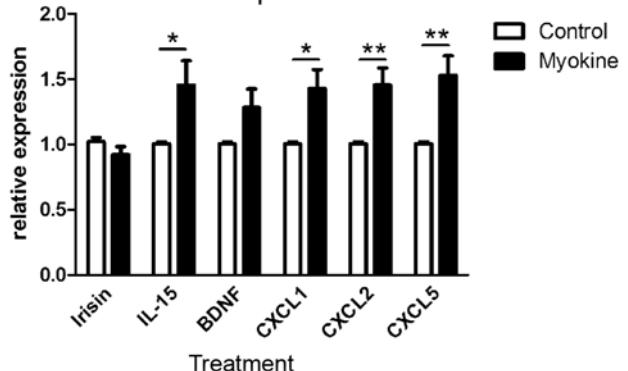
Supplemental Figure S2 related to Figure 4. *Treatment of RAW macrophages with different BNP concentrations increased the expression of pro- and anti-inflammatory cytokines in a dose-dependent manner.* After treating RAW macrophages with PBS, 1, 2, and 4 nM BNP for 4 h, mRNA expression levels of pro- and anti-inflammatory cytokines were determined by RT-PCR. Values represent the mean of at least 3 independent experiments +SEM. * $P<0.05$; ** $P<0.01$; *** $P<0.001$; BNP treated versus control.

A expression in C2C12 myotubes**B** TNF α expression

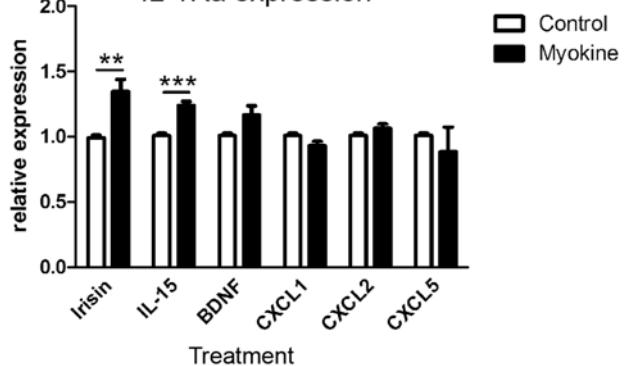
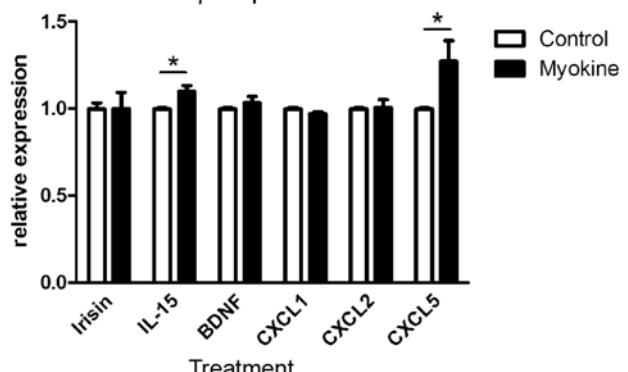
IL-6 expression

MIP-1 α expression

MCP-1 expression

**C** CCL22 expression

IL-1Ra expression

TGF β expression

IL-10 expression

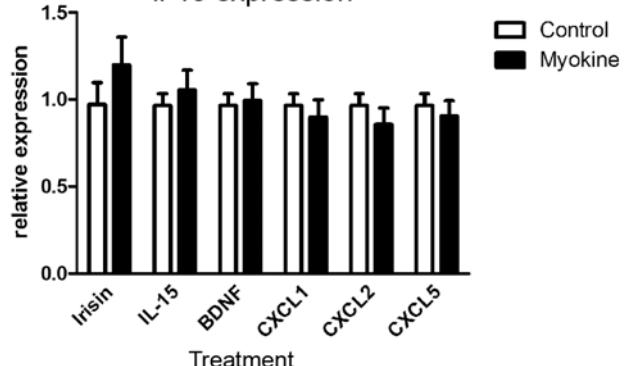


Figure S1

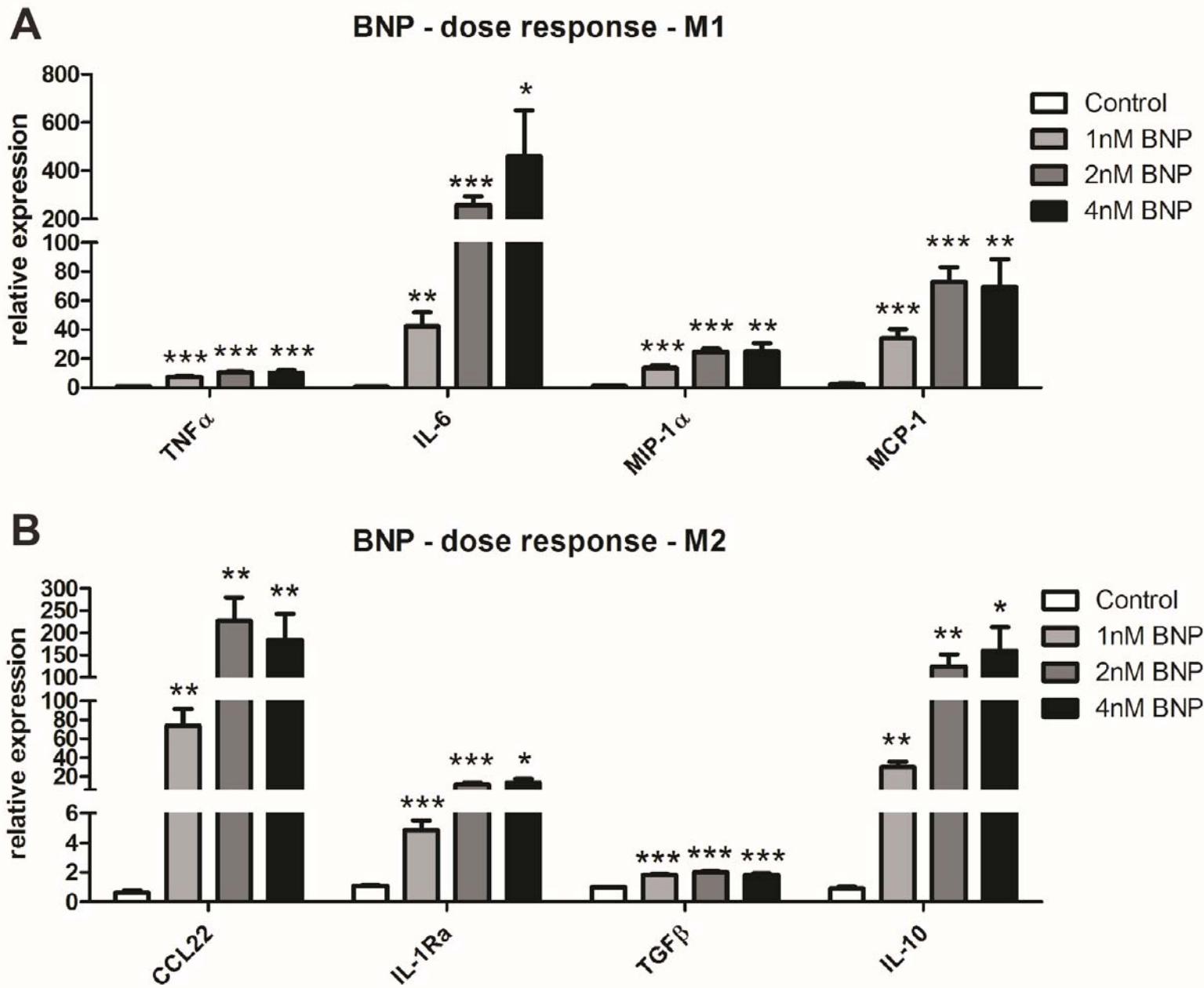


Figure S2

Supplemental Tables

Supplemental Table S1: Primer sequences for RT-qPCR.

Gene	Forward primer	Reverse primer
BDNF	CGGCGCCCATGAAAGAAGTA	AGACCTCTCGAACCTGCCCT
BNP	GGCCTCACAAAAGAACACCC	TGCCCAAAGCAGCTTGAGAT
CCL7	GCTGCTTCAGCATCCAAGTG	CCAGGGACACCGACTACTG
CCL22	TGGAGTAGCTTCTCACCCA	TCTGGACCTCAAAATCCTGC
CXCL1	TGAGCTGCGCTGTCAGTGCCT	AGAAGCCAGCGTTACCCAGA
CXCL2	GAGCTTGAGTGTGACGCCCCCAGG	GTTAGCCTTGCCTTGTTCAGTATC
CXCL5	GCATTCTGTTGCTGTTCACGCTG	CCTCCTCTGGTTTTCAAGTTAGC
CXCL10	TCTGAGTGGGACTCAAGGGA	AGGCTCGCAGGGATGATTTC
IL-1Ra	AAATCTGCTGGGGACCCTAC	TGAGCTGGTTGTTCTCAGG
IL-6	CCACGGCCTTCCCTACTTC	TTGGGAGTGGTATCCTCTGTGA
IL-10	CTGGACAACATACTGCTAACCG	GGGCATCACTTCTACCAGGTAA
IL-12	GCTTCTCCCACAGGAGGTTT	CTAGACAAGGGCATGCTGGT
IL-15	GAGGCCAAGAAGAGTTCTGGAT	TGCCCAGGTAAGAGCTTCAA
Irisin	ATGAAGGAGATGGGGAGGAA	CGGGCAGAAGAGAGCTATAACA
LCN2	TCTGATCCAGTAGCGACAGC	CAGAAGGCAGCTTACGATG
MCP-1	CCCAATGAGTAGGCTGGAGA	TCTGGACCCATTCCCTCTTG
MIP-1α	TCCCAGCCAGGTGTCATT	TTGGAGTCAGCGCAGATCTG
Pcolce	GCTCCATATCGAAGACTCGG	GTTACGTGGCAAGTGAGGGT
PGC-1α	TGATGTGAATGACTGGATACAGACA	GCTCATTGTTGACTGGTTGGATATG
TBP	GGCCTCTCAGAACATCACTA	GCCAAGCCCTGAGCATAA
TGFβ	CAACCCAGGTCTCCTCTAAA	GGAGAGCCCTGGATACCAAC
TNFα	CACAAGATGCTGGGACAGTGA	TCCTTGATGGTGGTGCATGA

Supplemental Table S2: Complete proteomic analysis

See attached Excel file “Proteomics_Supplemental_Table_S1.xlsx”