

SUPPLEMENT

Table S1. Primers used for real-time PCR. Primers used for gene expression analysis, sequence designed for human ZnTs and ZiPs using Primer 3.0 software

Sequence name	Primer	Supplement references [SR]
ZnT1F	TGTGAACCTGCCTGCAGAAC	this study
ZnT1R	TGGGGTCTTTCTGCATCCT	this study
ZnT2F	TGTCCTAGTGGCAGCCTATATT	[1]
ZnT2R	TCACATCTCTCAGGATGGTCAA	[1]
ZnT3F	CACCCTCCGAGACGTTCTTC	[2]
ZnT3R	GGCACCGACAACAGCGTAT	[2]
ZnT4F	TGACCTAAGCGCCATCATACT	[1]
ZnT4R	AGCTGACAAAACCTCTAACGCG	[1]
ZnT5F	GGAGGCATGAATGCTAACATGAGG	[3]
ZnT5R	GTGGATACGATCACACCAATGCTG	[3]
ZnT6F	CCTAACATACAGGCCAAGC	this study
ZnT6R	TGGTGTAGGCTGTGGTTCA	this study
ZnT7F	TTTCTCCTGTGCCTGAACCTCTC	[3]
ZnT7R	GAGTCGGAAATCAAGCCTAACGAG	[3]
ZnT8F	GATCCAGGCGACTGTGATGAT	[1]
ZnT8R	TGGCTTGTACTTCCTTGATTG	[1]
ZnT9F	AGCCTAGGTTCTTGGGTGT	this study
ZnT9R	ACTTGTCTGGCTGGATGGA	this study
ZnT10F	GCTGTGCCTGGAATTAGCAGT	[1]
ZnT10R	ATGTGCAGGGTGGCAATAATC	[1]
ZIP1F	TAAGGACAGGCTCACATGGG	this study
ZIP1R	GCTAGGACCAACAGGACACT	this study
ZIP2F	CTCACGATGGCAGTGTCTC	[4]
ZIP2R	ATGAAGGCAAAACCAGCGGC	[4]
ZIP3F	CTGCTGGCTTCTCATGAC	this study
ZIP3R	GTTGAAGGTCTCCAGGTCGA	this study
ZIP4F	ATGTCAGGAGCGGGTCTTGC	[5]

ZIP4R	GCTGCTGTGCTGCTGGAAC	[5]
ZIP5F	GGGGCTGTCAGTGCTCGGAG	[6]
ZIP5R	TCCGGATCCAAGTTGCGTGTT	[6]
ZIP6F	AGTTCCGTCTCTGCTGGG	this study
ZIP6R	CACCACTCAAAGTCCCAACG	this study
ZIP7F	AAGATTAGTGTCCCAGGGCC	this study
ZIP7R	CCGCTCTCCCATTCCCTAA	this study
ZIP8F	TGCTACCCAAATAACCAGCTCC	[6]
ZIP8R	ACAGGAATCCATATCCCCAAACT	[6]
ZIP9F	TCAACTTGCTGCTGAACCC	this study
ZIP9R	AGTCCCAGCAAACCTCTCCTC	this study
ZIP10F	CACAGTCACCAACATGCACA	this study
ZIP10R	TGCCTCCTAGAGCAACAA	this study
ZIP11F	CGGCATCTGCTACCTTGAGAG	[7]
ZIP11R	ATGATGTCGTCCATGACCACG	[7]
ZIP12F	TTTCCTGGGATCAGACCTGCT	[6]
ZIP12R	GTTGGTCCTGGTAAGTGGC	[6]
ZIP13F	TCTGAAAACATCCACCGTGA	this study
ZIP13R	AGCCATAATCCCCAACCAT	this study
ZIP14F	TTTGACCTCCTCTTCCCCAC	this study
ZIP14R	GCAGTCTCCCTCTAACACA	this study
MT-F	ATGGACCCCAACTGCTCCTGC	this study
MT-R	GGCACAGCAGCTGCACTCTC	this study
GAPDH-F	AATCCCATCACCATCTTCCA	this study
GAPDH-R	TGGACTCCACGACGTACTCA	this study
B-actin-F	TCCCTGGAGAAGAGCTACGA	this study
B-actin-R	AGCACTGTGTTGGCGTACAG	this study

Table S2. Zinc levels in cells were estimated using zinquin dye, zinc specific fluorochrome, comparing with zinc-zinquin standard curve. Cells were incubated with 10 µM zinquin for 30 min and washed with PBS to remove extracellular dye. Cells (5×10^6 /ml) were resuspended in PBS, fluorescence was measured at an excitation wavelength of 370 nm and emission wavelength 490 nm. The values represent mean \pm SE ($n = 3$). Values with different superscripts indicate significance of differences at $p < 0.05$. Significance of differences between treated and control samples was analysed using Analysis of variance (ANOVA) with post hoc multiple comparison by Dunnett's *t*-test

S. No	Specific cells	Control	Zinc	Zinc depletion
1	THP-1 monocytes	1.97 ± 0.103 nM	7.92 ± 0.10 nM	0.94 ± 0.06 nM
2	RD muscle	5.78 ± 0.3 nM	13.5 ± 0.44 nM	3.98 ± 0.45 nM

Table S3. Zinc transporter expression levels in response to exogenous zinc and zinc depletion

Gene	Zinc supplementation		Zinc depletion	
	THP-1	RD	THP-1	RD
<i>MT</i>	24↑	25↑	1.8↓	0.58↓
<i>ZnT1</i>	4↑	5↑	0.1↓	0.3↓
<i>ZnT5</i>			0.5↓	1.9↑
<i>ZnT6</i>			0.4↓	
<i>ZIP6</i>				1.5↑
<i>ZIP10</i>		0.3↓	4.7↑	2.5↑
<i>ZIP13</i>			0.3↓	1.7↑
<i>ZIP14</i>			4.9↑	

Notes. ↑ – Upregulated; ↓ – down regulated.

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