

Table S1. Primer sequences used for gene identification, synthesis of complete coding sequence and for shRNA gene silencing

Mouse gene	Gene ID	Primer Sequence	Product size (bp)
Primer sequences used for gene identification in the Quantitative-PCR method			
<i>Chemerin</i>	NM_027852	Fw TACAGGTGGCTCTGGAGGAGTTC Rv CTTCTCCCGTTTGGTTGATTG	195 bp
<i>CMKLR1</i>	NM_008153	Fw CGAGTTCTCAAACCCTGAAGTCGC Rv CAAGTCCACAAAGTAGCCAAAGCC	222 bp
<i>PPARγ</i>	NM_011146	Fw TCGCTGATGCACTGCCTATG Rv GAGAGGTCCACAGAGCTGATT	102 bp
<i>CD36</i>	NM_007643	Fw GAACCACTGCTTCAAAAAGTGG Rv GTCCTGAGTTATATTTTCCTTGG	178 bp
<i>Adiponectin</i>	NM_009605	Fw AGCCGCTTATATGTATCGCTCA Rv TGCCGTCATAATGATTCTGTTGG	118 bp
<i>ccna2</i>	NM_009828.2	Fw ACATTCACACGTACCTTAGGGA Rv CATAGCAGCCGTGCCTACA	242 bp
<i>ccnb2</i>	NM_007630.2	Fw GCCAAGAGCCATGTGACTATC Rv CAGAGCTGGTACTTTGGTGTTT	114 bp
<i>ccnd2</i>	NM_009829.3	Fw TGTGGATTGTCTCAAAGCCTG Rv CAACATCCCGCACGTCTGTA	149 bp
<i>CyclophilinA</i>	X52803	Fw GAGCTGTTTGACAGACAAAGTTC Rv CCCTGGCACATGAATCCTGG	124 bp
Primer sequences used in the synthesis of cDNA for the generation of expression vector			
<i>Chemerin</i>	NM_027852	Fw AAAAGGATCCGACTGAGGTGAAGCCATGAAGTGC Rv AAAAGAATTCTTATTTGGTTCTCAGGGCCCTGGAGAA	
<i>CMKLR1</i>	NM_008153	Fw AAAAGGATCCGAGATGGAGTACGACGCTTACAAC Rv AAAATCTAGATTCCCAGGTGAGGTTTCAGAGGG	
<i>PPARγ</i>	NM_011146	Fw AAAAGGTACCTATGCTGTTATGGGTGAAACTCTGG Rv AAAACTCGAGCCTGCTAATACAAGTCCTTGTAGATC	
Oligonucleotide sequences used for shRNA gene silencing			
<i>Chemerin</i>	NM_027852	GGAGTTGCAATGCATTAAGAT	
<i>CMKLR1</i>	NM_008153	GGAAGATAACCTGCTTCAACA	
<i>PPARγ</i>	NM_011146	GCCCTTACCACAGTTGATTT	

Fw, Forward; Rv, reverse; Restriction sites used for cloning are underlined.

Table S2. Primer sequences used for chemerin promoter analysis and chromatin immunoprecipitation (ChIP)

Primer	Position	Primer Sequence
Primer sequences used for generating chemerin promoter constructs, -4459/+38 and -600/+38		
Chem(-4459/+38)-Fw	-4459/-4435	AAAA <u>ACGCGT</u> ACATGAGTAAACCACCATTGCTTGC
Chem(-600/+38)-Fw	-600/-578	AAAA <u>ACGCGT</u> GACTGATGCCAGCCAGTCAAAGC
Chem-Rv	+16/+38	AAAA <u>CTCGAG</u> CTCCTCTCTCTTGGTCCCCAAAG
Primer sequences used for generating direct repeat 1 (DR1) constructs, pDR1 (-61/-49) and dDR1 (-472/-460)		
pDR1-Fw	-125/-104	AAAA <u>CTCGAG</u> GTCTCATTGGTGCTCCAGGC
pDR1-Rv	-12/+9	AAAA <u>AGATCT</u> CCTTTTCCCACTGACCTTCTC
dDR1-Fw	-566/-546	AAAA <u>CTCGAG</u> GGCTGTGAACAAACCCGAGAC
dDR1-Rv	-394/-371	AAAA <u>AGATCT</u> TGACCAGGTGATCTTCCACTAGAA
Primer sequences used for generating promoter constructs with mutations in the DR1 elements		
ΔpDR1-Fw		GCTCAATGCAAGAGGAG <u>GTAC</u> CGGGAACCTTGGGAAACAGAAACTC
ΔpDR1-Rv		GAGTTTTCTGTTTCCCAAAGTTCCCG <u>GTAC</u> CTCCTCTTGCATTGAGC
ΔdDR1-Fw		GGAGCTCTCTCACTGGACACAC <u>GGTAC</u> CAACTGAAGCAACAGACC
ΔdDR1-Rv		GGTCTGTTGCTTCAGTT <u>GGTAC</u> CGTGTGTCCAGTGAGAGAGCTCC
Primer sequences used for identifying the specific binding of chemerin promoter target sequences to PPARγ in the CHIP assay		
CHIP-target region-Fw	-138/-115	TATTCAGCCTGCAGTCTCATTGG
CHIP-target region-Rv	-35/-13	CCAGAGTCCGGAGTTTGGAGTT
CHIP-nontarget region-Fw	-3693/-3670	AGAAGAACAGAGACCACGGGTATG
CHIP-nontarget region-Rv	-3529/-3505	GGACTTGAAGTGAAGAGAAATGGCAG

Fw, Forward; Rv, reverse; Restriction sites used for cloning are underlined.