

## Supplemental Materials for

### **Immunometabolic modulation of retinal inflammation by CD36 ligand**

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Fig. S1. MPE-001 downregulates subretinal inflammation and photoreceptor degeneration in light-exposed CX3CR1<sup>-/-</sup> and CX3CR1<sup>-/-</sup>/CD36<sup>-/-</sup> mice.

Fig. S2. Effect of MPE-001 on R-FSL1-stimulated release of cytokines in human macrophages.

Fig. S3. Colocalization of CD36 and TLR2 in lipid rafts and MPE-001 effect on TLR2-signaling

Fig. S4. MPE-001 exerts no effect on macrophage phagocytosis.

Fig. S5. Azapeptide MPE-001 decreases caspase-1 expression in outer retina.

Fig. S6. Immuno-metabolic effects of MPE-001 on macrophages.

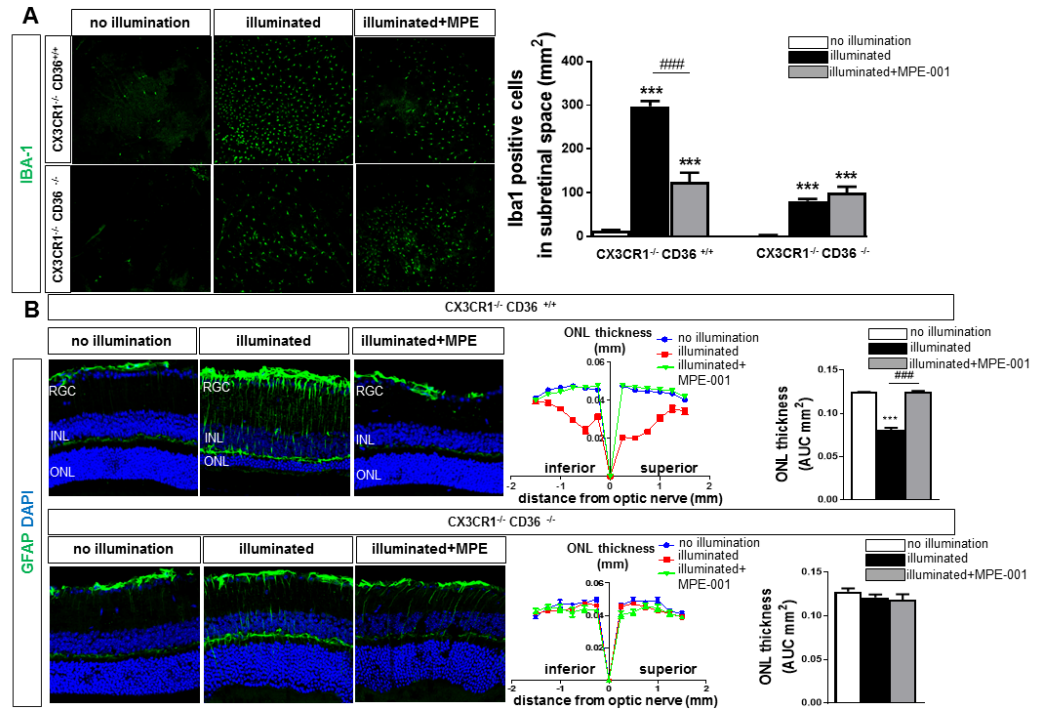
Fig. S7. Subretinal MPs in CD36<sup>+/+</sup>/TLR2<sup>+/+</sup> in mice and aging human retina.

Table S1. Percentage of pro-inflammatory cytokine/chemokine reduction in response to MPE-001-treatment in TLR2-stimulated peritoneal macrophages.

Table S2. Percentage of IL-1 $\beta$  reduction in response to MPE-001-treatment on R-FSL1-stimulated peritoneal macrophages.

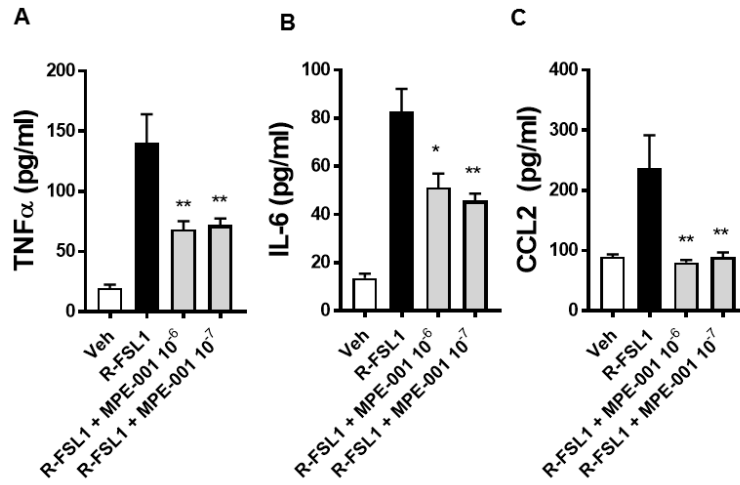
Table S3. Primers ID and sequences used for quantitative RT-PCR

## Supplemental figures



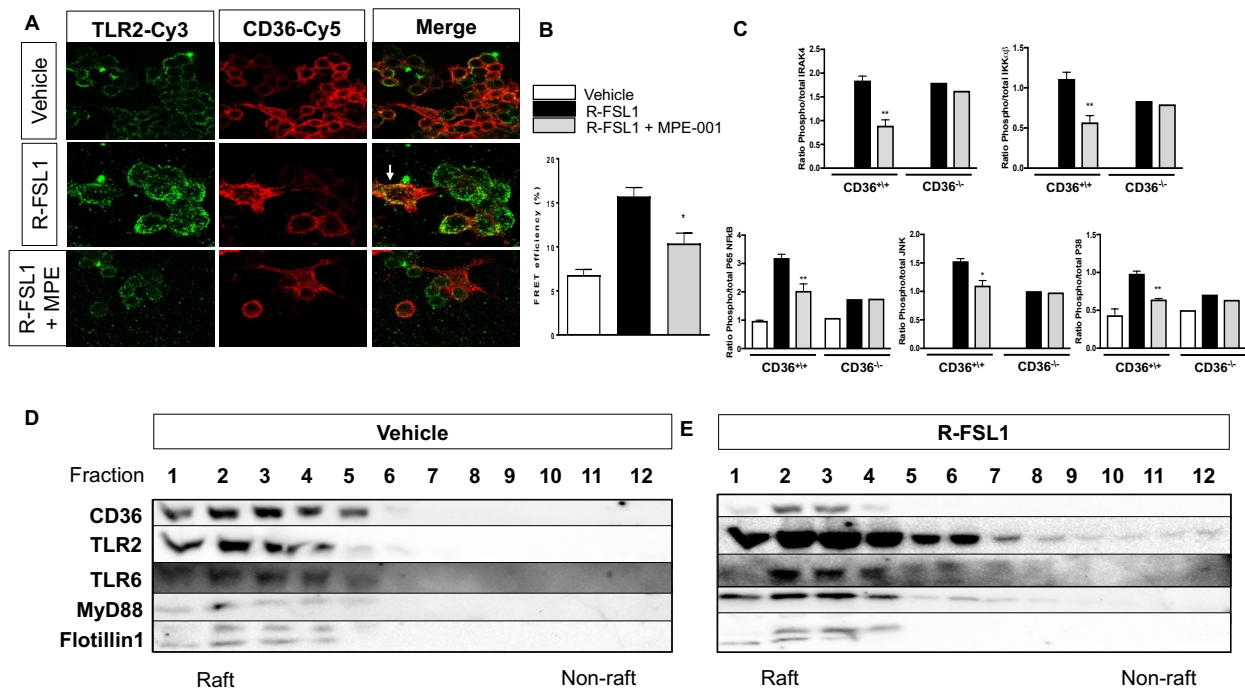
### Figure S1. MPE-001 downregulates subretinal inflammation and photoreceptor degeneration in light-exposed CX3CR1<sup>-/-</sup> and CX3CR1<sup>-/-</sup>/CD36<sup>-/-</sup> mice.

(A) Evidence of CD36 involvement in light-induced accumulation of subretinal macrophages. Representative RPE/Chr/Scl flat mounts, illustrating the accumulation of IBA-1+ cells (green) in the subretinal space of CX3CR1<sup>-/-</sup> or CX3CR1<sup>-/-</sup>/CD36<sup>-/-</sup> mice illuminated with blue light for 5 days and treated or not with 289 nmol/kg per day of MPE-001 from day 1 to day 7. Quantification of IBA-1+ cells in the subretinal space of mice unexposed, illuminated with blue light, and treated or not with MPE-001 (n = 3-4 mice per group). (B) Evidence of CD36 involvement in light-induced degeneration of ONL. GFAP staining (green) in retinal cryosections from CX3CR1<sup>-/-</sup> or CX3CR1<sup>-/-</sup>/CD36<sup>-/-</sup> mice exposed or not to blue light and treated or not with MPE-001. ONL thickness measurements and spider graph representations showing both side of the optic nerve from CX3CR1<sup>-/-</sup>/CD36<sup>+/+</sup> or CX3CR1<sup>-/-</sup>/CD36<sup>-/-</sup> mice unexposed or exposed to blue light for 5 days and treated or not with 289 nmol/kg per day of MPE-001 from day 1 to day 7. Area under the curves from ONL thickness measurement. One-way ANOVA with Newman-Keuls for multiple comparisons was performed. \*\*\*  $P < 0.001$  vs no illumination, ###  $P < 0.001$  vs illuminated group. Data are shown as mean  $\pm$  S.E.M. ONL: Outer Nuclear Layer, INL: Inner Nuclear Layer, GCL: Ganglion Cell Layer.



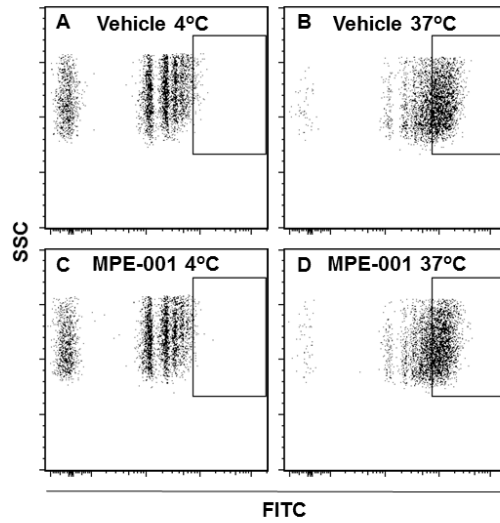
**Figure S2. Effect of MPE-001 on R-FSL1-stimulated release of cytokines in human macrophages.**

(A - C) TNF $\alpha$ , IL-6 and CCL2 in supernatants of human macrophages after 4 h stimulation with 300 ng/ml R-FSL1 in the presence of MPE-001 or vehicle. One-way ANOVA with Newman-Keuls post-test for multiple comparisons was performed. Data are representative of 2 independent experiments (n = 4 per group). \*  $P < 0.05$ , \*\*  $P < 0.01$  vs R-FSL1. Data are shown as mean  $\pm$  S.E.M.



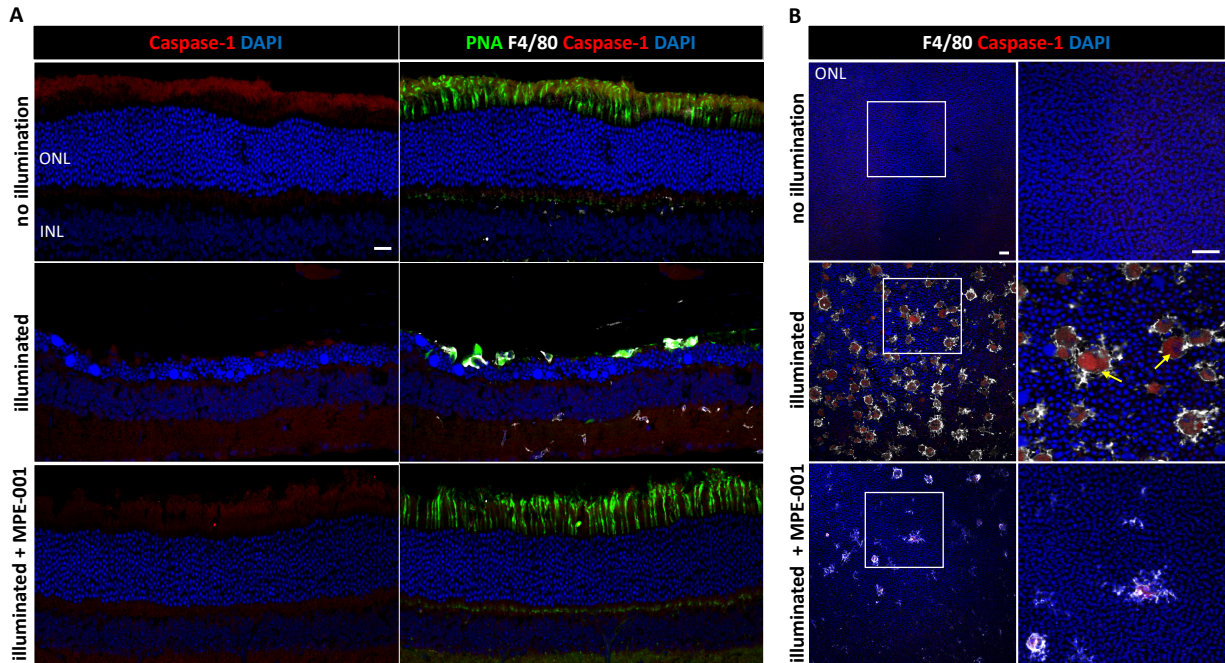
**Figure S3. Colocalization of CD36 and TLR2 in lipid rafts and MPE-001 effect on TLR2-signaling.**

(A) RAW macrophages were stimulated with 300 ng/ml R-FSL1 and treated with  $10^{-7}$  M MPE-001 for 10 min. Azapeptide MPE-001 disrupted the interaction between CD36 labeled with Cy5 (red), TLR2 labeled with Cy3 (green) and merge shown in yellow as assessed by fluorescence resonance energy transfer (FRET), measured using LSM-700 (Zeiss) confocal microscope. (B) Quantification of FRET efficiency expressed as the percentage of energy transfer. Data are representative of 3 independent experiments ( $n = 10-30$  per group). (C) Densitometric analysis data of western blots shown in Fig. 4E. (D, E) CD36 and TLR2 colocalization in lipid rafts after R-FSL1 stimulation. Peritoneal macrophages were treated with vehicle (D) or stimulated with 300 ng/ml R-FSL1 (E) for 5 min and subjected to sucrose density gradient ultracentrifugation. Data are representative of 3 experiments. The relative positions of the raft and non-raft (soluble) fractions are indicated. One-way ANOVA with Newman-Keuls post-test for multiple comparisons was performed.  $*P < 0.05$ . Data are shown as mean  $\pm$  S.E.M.

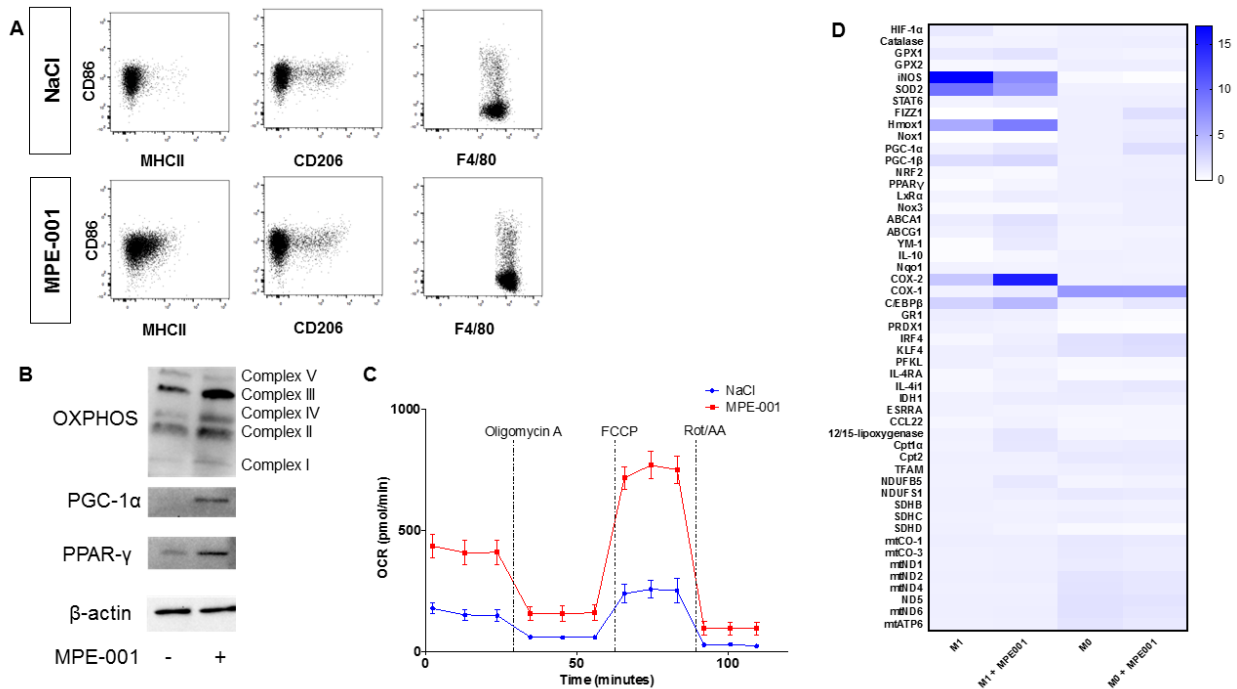


**Figure S4. MPE-001 exerts no effect on macrophage phagocytosis.**

(A-D) Proinflammatory induced-BMDM were incubated with yellow-green microspheres at 4°C, to determine background values (A, C); or at 37°C, to measure the phagocytosed microspheres (B, D). Phagocytosis is expressed as the relative number of cells that have ingested fluorescent beads in vehicle (A, B), or after MPE-001 treatment (C, D). Data are representative of 2 independent experiments.

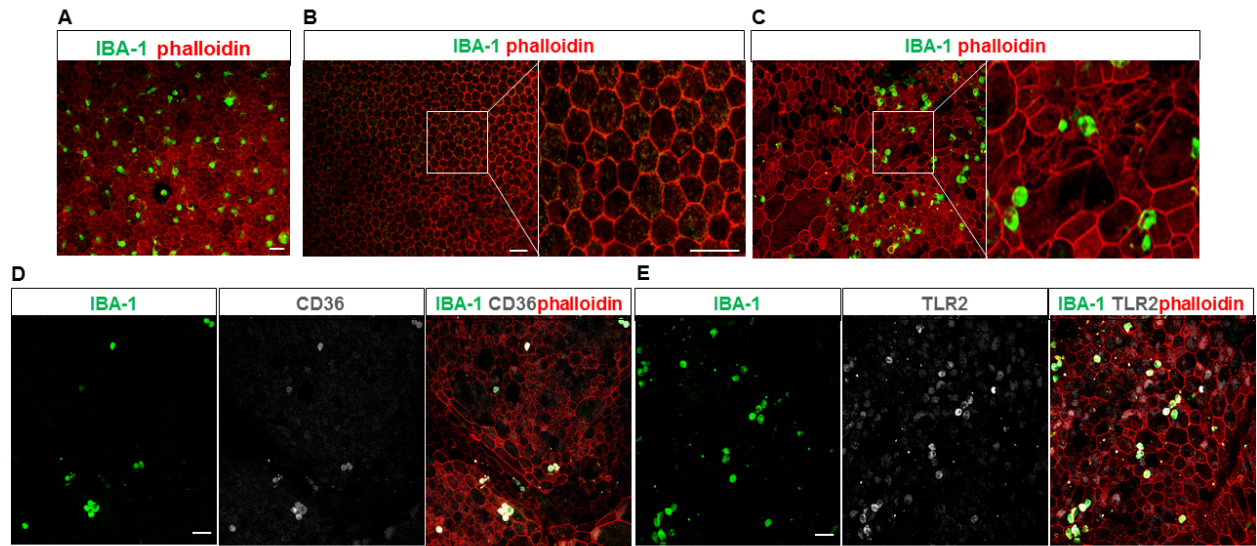


**Figure S5. Azapeptide MPE-001 decreases caspase-1 expression in outer retina.** Confocal microscopy of retina cryosection (A) and neuroretinal flat mounts (photoreceptors side) (B) from illuminated CD36<sup>+/+</sup> mice treated or not with MPE-001 stained with caspase-1 (red), PNA (green) and F4/80 (white); nuclei were counterstained with DAPI (blue). Magnifications of white square show caspase-1 distribution in outer retina. Yellow arrows show caspase-1 expression in subretinal F4/80+ cells. Scale: 10  $\mu$ m.



**Figure S6. Immuno-metabolic effects of MPE-001 on macrophages.**

(A-C) Peritoneal macrophages were isolated from WT mice treated daily *s.c.* injection of MPE-001 (289 nmol/kg) or NaCl for 7 days. (A) Phenotypic analysis of isolated peritoneal macrophages by flow cytometry using MHCII, CD86 and CD206 markers. (B) Protein expression of mitochondrial electron transport chain (mtETC) complex subunits (OXPHOS), PGC-1 $\alpha$  and PPAR- $\gamma$ . (C) OCR analysis of peritoneal macrophages. Data in A-C are representative of 4 independent experiments. (D) The heat map of 51 genes expression from M1 and M0 BMDM treated with vehicle or MPE-001 ( $10^{-7}$  M) for 24 h and analyzed by qPCR (n = 3 per group).



**Figure S7 : Subretinal MPs in  $CD36^{+/+}/TLR2^{+/+}$  in mice and aging human retina. (A-D)** Confocal microscopy of RPE flat mounts obtained from illuminated  $CD36^{+/+}/TLR2^{+/+}$  mouse or aging human donors. **(A)** Double-labeling of subretinal MPs of RPE flat mounts from illuminated mouse showing IBA-1 (green) and phalloidin (for F-actin, red). **(B, C)** Double-labeling of subretinal MPs of RPE flat mounts from human subjects aged 77 **(B)** and 79 **(C)** years old, without and with subretinal inflammation, respectively. Immunofluorescence showing IBA-1 (green) and phalloidin (for F-actin, red). **(C)** White squares magnification (3-fold) show RPE cell morphology (red). **(D)** Triple labeling of subretinal MPs with IBA-1 (green), CD36 (white) and phalloidin (for F-actin, red), of RPE flat mount from a 79-year-old subject with eye inflammation. **(E)** Triple labeling of subretinal MPs with IBA-1 (green), TLR2 (white) and phalloidin (for F-actin, red), of RPE flat mount from a 79-year-old subject with eye inflammation. Scale bar = 100  $\mu\text{m}$ .



## Supplementary tables

**Table S1.** Percentage of pro-inflammatory cytokine/chemokine reduction in response to MPE-001-treatment in TLR2-stimulated peritoneal macrophages.

	TNF $\alpha$	IL-6	CCL2	IL-12
R-FSL1	38.4 ( $P < 0.01$ )	37.6 ( $P < 0.01$ )	41.5 ( $P < 0.01$ )	39.2 ( $P < 0.01$ )
LTA	24.5 ( $P < 0.01$ )	35.0 ( $P < 0.01$ )	31.9 ( $P < 0.01$ )	29.1 ( $P < 0.01$ )
pgLPS	38.4 ( $P < 0.01$ )	42.3 ( $P < 0.01$ )	30.9 ( $P < 0.01$ )	33.4 ( $P < 0.01$ )

**Table S2.** Percentage of IL-1 $\beta$  reduction in response to MPE-001-treatment on R-FSL1-stimulated peritoneal macrophages.

	IL-1 $\beta$
MPE-001 10 <sup>-9</sup>	39.2 ( $P < 0.05$ )
MPE-001 10 <sup>-8</sup>	62.7 ( $P < 0.01$ )
MPE-001 10 <sup>-7</sup>	74.4 ( $P < 0.01$ )

**Table S3.** Primers ID and sequences used for quantitative RT-PCR.

Oligo ID	Gene	UPL Probe	Oligo FWD	Oligo REV	RefSeq	Slope
IR5845	Chil3	88	ggtctgaaagacaagaactgag	gagaccatggcaactgaacg	NM_009892.3	3,2
IR0878	Hif1a	18	catgatggctcccttttca	gtcaactgggtctgcaata	NM_010431.1	3,59
IR1044	Cat	68	gcgaccagatgaagcagtg	gtggtaaggacatcaggctctc	NM_009804.1	3,4
IR1055	Gpx1	2	tttccgtgcaatcagttc	tcggacttactgagggaat	NM_008160.2	3,5
IR1068	Nos2	13	ctttccacggagcagac	tcattgtactctgagggtgac	NM_010927.1	3,2
IR1084	Sod2	3	gaccattgcaaggaaaca	gtagtaagcgtctcccacac	NM_013671.3	3,3
IR1262	Stat6	3	tcctggtcacagttcaataagg	caaaccactgccaaactgtg	NM_009284.1	3,52
IR2537	Retnla	51	ccctccactgtaacgaagactc	cacaccagtagcagtcattcc	NM_020509.3	3,44
IR4021	Ppargc1a	29	gaaagggcacaacagagaga	gtaaatcacagcgctctct	NM_008904.2, NR_027710.1	3,19
IR4024	Nfe2l2	18	catgtggacttgagttgc	cctccaaaggatgcaatcaa	NM_010902.3	3,43
IR4352.3	Pparg	67	caagcctttaccacagttga	caggttctacttctgactcatt	NM_001127330.1, NM_011146.3	3,5
IR5347	Nr1h3	52	gagtgctgacttgcacaatg	cgatctgttctctgacacg	NM_013839.4	3,6
IR6026	Ptgs2	83	caatgtgcaagatccacagc	gtctggatggaggagcact	NM_011198.4	3,48
IR4022	Ppargc1b	88	gacgtggacagcttctact	gagcgtcagagcttctgttt	NM_133249.2	3.40
IR5843	Abca1	26	atggagcagggaagaccac	gtaggcctgcccagaagt	NM_013454.3	3.2
IR5844	Abcg1	64	tgctgtcactcacctatgtaa	ttcccaggataccctgtcc	NM_009593.2	3.3
IR1075	Prdx1	15	gtgagactgtggctcgac	gttccatctggcataacagc	NM_011034.2	3.3
IR5846.2	Il10	48	cagccgggaagacaataact	gtttgccagctggtctttg	NM_010548.2	3.3
IR6027	Cebpd	32	ctgaacgacctatacctcagacc	gcaggtcccaagaaactagc	NM_007679.4	3,38
IR1062	Gr1	64	gttcctcagagaccagat	tccagctgaaagaccatc	NM_010344.3	4
IR1075	Prdx1	15	gtgagactgtggctcgac	gttccatctggcataacagc	NM_011034.2	3,3
IR2319	Irf4	26	accatgacagcacttat	gggtggcatctgattatga	NM_013674.1	3,59
IR2899	Klf4	62	cggaagaggagaagacact	gagttctcagccaacg	NM_010637.3	3,33
IR3771	Cox1	101	cagaccgcaactaaacaca	gggtcccaagaatcagaata	5912286	3,1
IR3773	Cox3	88	cataaatcaagcctactaattacat	ctgaaatggagaatgatgttca	NP_904334.1	3,12
IR3774	Nd1	29	acacttattacaaccaagaacacat	tcatattatggctatgggtcagg	NP_904328.1	3,18
IR3775	Nd2	67	ccataactcaactctcactctatg	gaatcctgttagtgggaagg	NP_904329.1	3,27
IR3776	Nd4	86	gcttaaacgaggattatt	gggttctacatgttttgg	NP_904337.1	3,12
IR3777.2	Nd5	31	agcattcggagactctttg	tttgaggactggaatgctg	NP_904338.1	3,29
IR3778	Nd6	12	cacaactatattgctcctacc	ttggttggagattggttg	NP_904339.1	3,15
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IR4349	Cox5b	26	gtgggttccccactgatga	tgaagctccttggaggta	NM_009942.2	3,6
IR5344	Cpt2	66	ccaaagaagcagcagtg	tagagctcagcagggta	NM_009949.2	3,4
IR5345	Cd36	75	ccaagctattgacacatgatt	ttcaatgtccgagactttca	NM_001159558.1	3,33
IR5353	Cox7b	21	aacgcactaagccgtctcc	catgaaactaggtccctct	NM_025379.2	3,12
IR5354	Ndufab1	78	tgcatgataagaagatgtgtatgaa	ttcactctgtgtcagttgtga	NM_028177.3	3,3
IR5355	Ndufb6	31	agctcgaagcccaggatatt	cttcaatgatgttgcagagaaa	NM_001033305.2	3,37
IR5356	Ndufs1	60	tgactctgacacttatgactga	aagataattggaactgaagctgtacc	NM_145518.2, NM_001160038.1, NM_001160039.1	3,33
IR5358	Sdhb	42	ctggtgaaacggagacaagt	gcgttctctgtgaactctg	NM_023374.3	3,38
IR5359	Sdhc	68	cactgaatgctcagctttgt	ttccagaaacgctcactct	NM_025321.3	3,3
IR5360	Sdhd	3	tctgcttggggctgat	cccatgaacgtatcggtaac	NM_025848.3	3,3
IR5365	Pfkf	42	attgaccggatggaag	aagcccagcctgaacc	NM_008826.4	3,22
IR5641	Cpt1a	56	cttcaactctccgcatcc	gcctctgtgtacacgacaat	NM_013495.2	3,3
IR5972	Il4ra	97	gagtgaggctctagcatcacg	cagaggcaggagatgggtga	NM_001008700.3	3,34
IR6088	Me1	16	cagaggccctgagatgacg	ccgattggcaaaattctcaa	NM_008615.2, NM_001198933.1	3,49
IR6089	Pgd	48	aaagatccgggacagtgct	gagcaaaagacagcttccaa	NM_001081274.2	3,13
IR6090	Esrra	45	gaccttggcagtagctgga	tggcgtacagcttctcaggt	NM_007953.2	3,36
IR6227.2	Ccl22	53	gccagactacatcctgca	cggttcttgcggttatcaaa	NM_009137.2	3,42
IR6228	Il4i1	45	gggccaatcggacttc	caatcctgttatctcctcca	NM_010215.3	3,6
IR6229	Idh1	77	tcgaaagtgacatacatggtaca	ttgaaagaaactgtgtgcaa	NM_010497.3, NM_001111320.1	3,26
IR6230	Ptgs1	76	actgtggatgcttctctc	tctcggactccttgatgac	NM_008969.4	3,18
IR6231.2	Alox15	10	gagatgggttgcagggtg	gatgaagtggcaagcctga	NM_009660.3	3,8
IR3659.3	Actb	56	aagccaacgtgaaaagat	gtggtacgaccagaggcatac	NM_007393.3	3,33
IR3669.2	Gapdh	80	gtctcgtctggatctgac	cctgtccaccactcttctg	NM_008084.2	3,55