

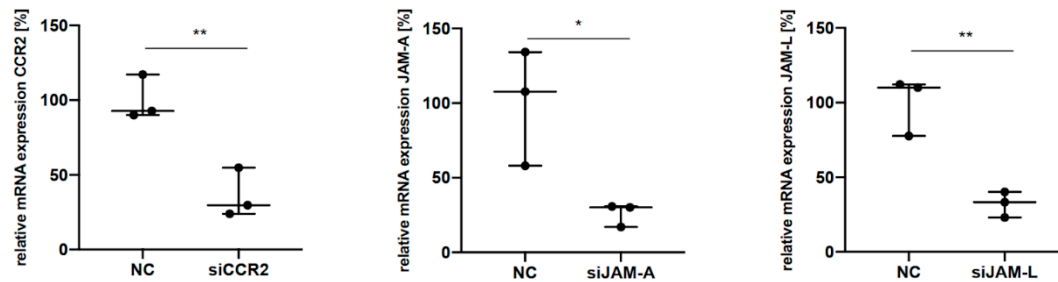
Supplementary Information

Table S1. The primer sequences.

Method	Name	Sequence
Cloning	JAM-A forward	5' GTGTGTTGGAGGGCTTCTTAG 3'
	JAM-A reverse	5' ATGCAGACCACCCATGATCT 3'
Cloning	JAM-L forward	5' AGTCCCTTCATCTCAGCAGC 3'
	JAM-L reverse	5' CAACCCATTACCCCAAACC 3'
qRT-PCR	CCR2 forward	5'TGAGACAAGCCACAAGCTGA 3'
	CCR2 reverse	5' TTCTGATAAACCGAGAACGAGAT 3'
qRT-PCR	JAM-A forward	5'AAGTTGTCCTGTGCCTACTCG 3'
	JAM-A reverse	5' CGGTCCTCATAGGAAGCTGT 3'
qRT-PCR	JAM-L forward	5' CATGTTTTGCCCACTGAAAC 3'
	JAM-L reverse	5' CATTGAGCCCAAGGAATAA 3'
qRT-PCR	B2M forward	5'TTCTGGCCTGGAGGCTATC 3'
	B2M reverse	5' TCAGGAAATTTGACTTTCCATTC 3'
qRT-PCR	TBP forward	5' GAACATCATGGATCAGAAACA 3'
	TBP reverse	5' ATAGGGATTCCGGGAGTCAT 3'

Supplementary Figure S1

A



Supplementary Figure S1 Knockdown efficiency of primary human monocytes transfected with siRNAs targeting CCR2, JAM-A and JAM-L.

(A) Knockdown efficiency of siRNAs indicated as relative mRNA expression of respective target mRNA after transfection of individual siRNAs compared to negative control (n=3, * $p < 0.05$, ** $p < 0.01$).

Supplementary Figure S2. Gene names of regulated genes on the heatmap ordered by clusters.

Cluster 1

11726524 s at - ZFH3	11720057 a at - POLR1D	
11727707 a at - BBC3	11733389 a at - DEF6	
11742871 a at - PIK3IP1	11724081 a at - TOR2A	
11748806 a at - ARID3B	11724082 x at - TOR2A	
11720739 s at - ITPKB	11756694 a at - RNF44	
11751393 a at - PIK3IP1	11743343 at - RMND5A	
11741051 x at - IMPDH1	11724499 at - MFNG	
11729130 at - ANAPC16	11755502 x at - TECR	
11725297 a at - MAPRE2	11718676 x at - TMEM245	
11737440 a at - BMF	11763731 a at - VPS36	
11720768 at - METTL9	11722039 x at - FES	
11730935 a at - AKNA	11756983 a at - F11R	
11740932 a at - CSF3R	11758645 s at - EIF4EBP2	
11723521 x at - ZNF385A	11755355 s at - RCSD1	
11749329 a at - ARHGAP9	11730478 at - RNF125	
11721421 s at - RAP1GAP2	11753321 a at - NA	
11734778 a at - PCTP	11752590 s at - CCR2	
11750840 s at - TGFBR2	11723849 a at - MS4A6A	
11717720 a at - ST6GAL1	11718062 a at - NME3	
11735157 at - CYSLTR1	11718675 s at - TMEM245	
11737504 x at - SIRPB2	11744475 x at - TRABD	
11756441 x at - CSF3R	11729748 a at - IL10RA	
11727552 at - FZD2	11745326 x at - DUS1L	
11749377 a at - CECR1	11716402 at - EIF4EBP2	
11741447 s at - RNASE4	11738091 a at - ABTB1	
11755975 x at - PKIB	11722888 s at - NA	
11743171 a at - RCSD1	11756846 a at - PRAM1	
11763426 a at - TRAF3IP3	11722038 a at - FES	
11727643 s at - TRERF1	11740874 a at - FES	
11742816 a at - FAM13A	11755715 x at - CD46	
11738980 a at - CD4	11750001 x at - CLEC7A	
11743170 a at - RCSD1	11751549 a at - CD46	
11721583 x at - LRMP	11722952 a at - SLC27A1	
11760710 a at - MS4A6A	11720461 a at - F11R	
11731066 x at - GIMAP1	11720763 a at - SELPLG	
11737820 a at - MAP3K3	11729758 at - RGS18	
11724061 s at - OSBPL1A	11728559 a at - ASGR1	
11763695 a at - ABTB1	11717341 at - PTGFRN	
11725830 a at - PLCL2	11719667 s at - STMN1	
11739079 a at - ANP32A	11718988 s at - RNASE4	
11734777 a at - PCTP	11738982 at - CD4	
11729013 at - ACPP	11738981 x at - CD4	
11718818 s at - SLC26A6	11724979 a at - DPEP2	
11721107 a at - TRABD	11757026 x at - SERPINF1	
11734920 s at - PCYOX1L	11757025 s at - SERPINF1	
11758673 s at - SORD	11724008 a at - ENTPD1	
11726393 at - FGD4	11722970 a at - CREB5	
11750157 at - CD4	11722971 a at - CREB5	
11752633 a at - OLFM1	11733801 s at - CPPED1	
11723885 at - NATD1	11753705 x at - STMN1	
11754362 a at - TTC7A	11732351 at - HGF	
11727385 a at - PCCA	11722681 at - RBP7	
11730331 at - BRI3BP	11752825 a at - ARHGEF6	
11745554 a at - ASGR1	11735309 a at - MYCL	
11724292 s at - TRIT1	11728522 a at - MYCL	
11718283 a at - EIF4EBP1	11751425 x at - TRAF3IP3	
11732135 a at - ADCK3	11736681 at - SLC46A1	
11756746 s at - NA	11755091 a at - ABCA7	
11739744 x at - MX11	11724291 a at - TRIT1	
11757868 a at - PIK3IP1	11721582 a at - LRMP	
11757767 s at - NAV1	11721569 a at - ANG	
11754492 x at - PLCB2	11746729 a at - LRMP	
11720595 a at - CD46	11745165 a at - TRAF3IP3	
11725092 s at - RFX5	11759029 at - BRI3BP	
11742219 a at - CD46	11751424 a at - TRAF3IP3	
11751532 x at - CD46	11749685 a at - FAM13A	
11731200 a at - TLR5	11750052 a at - ASGR2	
11758893 at - MLXIP	11722448 at - KCNM84	
11727007 at - SORT1	11759152 at - CYSLTR1	
11753491 s at - ERP29	11749188 a at - ENTPD1	
11718284 x at - EIF4EBP1	11724992 x at - GTPBP3	
		11743378 a at - NIN
		11723433 at - TBC1D22A
		11751525 a at - CCDC69
		11730323 at - RPS29
		11740088 s at - PIK3IP1
		11717339 s at - SORD
		11752942 s at - DBP
		11716232 at - CTNNB1P1
		11736962 a at - MBNL2
		11758238 s at - ABCC4
		11749411 a at - SPIDR
		11720856 a at - INSR
		11746542 x at - SPIDR
		11747171 a at - MXD4
		11718304 s at - NA
		11748699 a at - RGS14
		11735728 x at - ZNF385A
		11725091 at - RFX5
		11720464 x at - F11R
		11751394 s at - PIK3IP1
		11742872 s at - PIK3IP1
		11759028 at - BRI3BP
		11744281 a at - FAM53B
		11749056 a at - RCSD1
		11730477 at - RNF125
		11718081 a at - ATP2B4
		11743879 at - MAML3
		11719332 at - ARHGAP15
		11750682 a at - TPCN1
		11725093 a at - RFX5
		11741179 a at - ABTB1
		11749057 x at - RCSD1
		11749214 a at - SELPLG
		11741250 a at - SH2D3C
		11747498 s at - SORD
		11729268 a at - STAC3
		11740018 a at - INSR
		11754685 a at - RGS14
		11754990 a at - EPB41
		11755873 a at - ACS1
		11727553 a at - PKIB
		11737038 at - C19orf38
		11743677 at - P2RY8
		11717508 at - IRF4
		11731147 at - TLR7
		11756371 a at - VSI64
		11731148 at - TLR7
		11762274 x at - HLA-DMB
		11725827 s at - CD1D
		11729871 a at - ASGR2
		11760018 at - CLEC10A
		11727554 s at - PKIB
		11726485 at - CPED1
		11741874 x at - SEPP1

Cluster 2

11753719 a at - GMFB
11753358 a at - ACVR1
11758185 s at - TWF1
11746929 a at - MOB3C
11735676 a at - OSGIN1
11717524 s at - RDX
11748968 x at - RDX
11724523 a at - CDK5RAP2
11728498 a at - SVIL
11745431 a at - SVIL
11728499 x at - SVIL
11722593 s at - BCAP29
11757628 s at - TSPYL2
11717843 at - ETNK1
11752386 a at - NAB1
11730052 a at - SNX16
11716945 s at - TWF1
11746201 a at - NAB1
11743972 a at - DDIT4
11734657 s at - NA
11745021 a at - MYC
11721308 at - PHLDA1
11758781 at - EMP1
11744648 s at - AIFM2
11733240 at - IRAK2
11757558 s at - LONRF1
11729751 a at - LPAR1
11724524 x at - CDK5RAP2
11716654 a at - EML4
11716653 x at - EML4
11716500 a at - PIR
11763806 x at - KCNN4
11721307 at - PHLDA1
11729750 a at - LPAR1
11715711 a at - AKR1C3
11721775 s at - ASPH
11745104 a at - PJA1
11724715 a at - PJA1
11748596 a at - KLHL2
11757836 s at - SEC24A
11763581 a at - SLC30A6
11724676 a at - KLHL2
11731452 s at - SEC62
11720444 s at - ERLIN1
11748951 a at - CTNNB1
11748952 x at - CTNNB1
11747388 x at - CTNNB1
11747361 x at - HMGA1
11750680 x at - CTNNB1
11723135 s at - SRC
11720755 at - STAT3
11746889 a at - CKAP4
11733110 s at - TLK1
11736217 at - MAP3K8
11747602 a at - MAP3K8
11749022 x at - CTNNB1
11720769 a at - PPIF
11717716 at - FNDC3B
11724424 at - GPR68
11717567 a at - NQO1
11718875 a at - PNMA1
11733109 a at - TLK1
11737748 a at - GMFB
11718218 at - RAB22A
11736513 s at - NMT2
11736006 at - MAPK11P1L
11732718 at - EREG
11717715 at - FNDC3B
11722796 a at - MAPK6
11763785 x at - KCNN4
11719227 x at - KCNN4
11729847 a at - CCL7

Cluster 3

11744921 a at - SLC43A1
11756191 a at - MAN1C1
11750879 a at - CREB5
11738889 at - IQCD
11720857 a at - INSR
11728286 at - RBMS2
11746198 a at - METTL9
11751817 x at - MBNL2
11725410 at - NAV1
11737027 a at - FLT3
11748399 s at - VCL
11721626 a at - SEPT6
11746773 a at - SLC27A1
11719036 at - UVRAG
11725053 x at - TOP1MT
11755265 s at - NT5DC1
11741251 a at - SIRPB2
11731879 a at - NLRP1
11747618 a at - ADCK3
11748695 a at - SYNGR1
11735306 at - C9orf139
11733650 s at - FAM228B
11755804 a at - SEMA4D
11730068 a at - HGF
11740366 a at - SHMT1
11753045 a at - MBNL2
11732417 at - FAM105A
11720204 at - NT5DC1
11728285 at - RBMS2
11732699 a at - PKIB
11732700 x at - PKIB
11758583 s at - TMEM154
11724763 s at - DEPTOR
11737134 x at - ZNF385A
11731602 at - TMEM154
11752189 a at - ARHGEF6
11728961 a at - NA
11754521 a at - MBNL2
11734791 x at - SEPT6
11742063 a at - BTN3A1
11753046 x at - MBNL2
11724293 a at - TRIT1
11739770 s at - TMOD2
11759015 at - SPECC1
11747732 a at - ADCK3
11753704 a at - STMN1
11751748 a at - CHN2
11727642 a at - TRERF1
11719666 a at - STMN1
11729227 a at - GRK5
11716403 at - EIF4EBP2
11730843 a at - MX1
11731578 s at - ACSS1
11749445 a at - ARHGAP15
11726459 at - FAM117B
11762132 a at - ΔMICΔ1

Cluster 4

11722391 at - MTF1
11725045 a at - UAP1
11749608 a at - RNF11
11753001 a at - PPARG
11758864 at - SRP19
11736274 at - TLR2
11718139 at - PDXK
11759869 x at - PJA1
11746814 x at - NAB1
11739508 a at - NAA30
11721330 s at - RNGTT
11739746 s at - SVIL
11717383 at - FOPNL
11761935 at - HNRNPPL
11730914 a at - ERLIN1
11763822 at - EREG
11747839 a at - LPAR1
11717527 a at - RDX
11729235 s at - E2F6
11725276 a at - CDK5RAP2
11731077 a at - LONRF1
11731076 a at - LONRF1
11763594 a at - HS3ST3B1
11739367 at - ASPH
11757312 x at - SERPINE1
11726809 x at - SLC12A8
11724275 s at - TMEM158
11735268 s at - SLC41A2
11727186 a at - WNT5A
11734811 a at - EGR2
11735116 a at - MET
11721774 a at - ASPH
11731577 a at - SLC22A1
11715751 a at - DPYSL3
11737251 at - CD226
11745300 a at - DNAJC6
11758577 s at - ELOVL7
11727189 x at - WNT5A
11751610 a at - DPYSL3
11717376 a at - SLC25A25
11724915 at - TMTC3
11746856 a at - SERPINE1
11735724 a at - TP53BP1
11718808 a at - TP53BP1
11752291 x at - SLC30A6

Cluster 5

11759611 at - SMIM11A
11724007 s at - ENTPD1
11725568 a at - ATP8A1
11751656 a at - TMEM154
11721955 at - RFX7
11756758 x at - SPECC1
11758230 s at - NFIA
11749401 a at - TRIT1
11732747 at - HGF
11716390 a at - BACE1
11724625 a at - SPATA7
11756155 a at - NHS
11724762 a at - DEPTOR

Cluster 6

11728984 a at - MTHFD1L
11756610 a at - CD226
11727188 at - WNT5A
11719215 a at - NA
11727840 at - HPD
11752813 a at - MSC-AS1
11745301 a at - DNAJC6

Cluster 7

11715635 x at - TMEM14C
11744429 a at - STX10
11719780 at - TNFAIP8L2
11739119 s at - CNPY3
11739146 a at - MKNK2
11739720 a at - GIMAP8
11757612 a at - ASRGL1
11732589 a at - ZNF467
11742220 x at - CD46
11716633 a at - TMEM50A
11719025 a at - ZNF652
11716404 s at - EIF4EBP2
11751550 x at - CD46
11718610 at - TP53INP1
11722767 s at - MBNL2
11723692 a at - CD302
11718090 a at - TPCN1
11732418 at - FAM105A
11729759 at - RGS18
11719845 a at - SMAP2
11747948 a at - SMAP2
11720219 a at - MAP3K3
11739118 a at - CNPY3
11719157 x at - ASRGL1
11717903 s at - IER3IP1
11723712 x at - MLXIP
11742744 a at - STX10
11733402 a at - CLEC7A
11754976 x at - CNPY3
11732415 s at - TIAM1
11745553 s at - SMAP2
11720218 a at - MAP3K3
11730936 s at - AKNA
11730498 s at - MXI1
11737870 s at - FAM78A
11743771 at - GIMAP7
11757602 x at - ERP29
11759405 x at - STX10
11743903 a at - CD46
11735060 a at - CD46
11744364 s at - INPP5D
11717618 s at - PXN
11723368 a at - ARHGEF6
11726947 a at - TLR8
11717340 at - PTGFRN
11718002 at - C10orf54
11715816 at - ZFP36L2
11740238 a at - SULF2
11720460 x at - F11R
11716527 a at - GSTK1
11721159 x at - DRAM2
11715627 x at - ALDH2
11717762 a at - CPVL
11749293 x at - MS4A6A
11730996 a at - AMICA1
11728523 a at - MYCL
11752737 a at - CLEC7A
11733403 x at - CLEC7A
11727788 a at - PIGS
11741510 a at - HERPUD1
11730383 a at - SNX18
11716359 a at - FAM96A
11735681 x at - CD46
11726820 a at - ARHGAP9
11718971 x at - CD46
11720380 a at - IMPDH1
11742681 s at - ERP29
11743931 x at - ERP29
11752331 s at - NA
11736017 s at - PXN
11721702 a at - CMTM7
11719844 at - SMAP2

11750531 a at - PYCARD
11721169 a at - PYCARD
11730090 a at - NA
11716554 a at - HLA-DMA
11745989 a at - HLA-DMA
11757815 s at - METTL7A
11723691 at - CD302
11752948 a at - PRCP
11721158 a at - DRAM2
11729747 a at - IL10RA
11726948 at - TLR8
11715848 a at - ATP5G2
11718160 at - KLF13
11754471 a at - HLA-DMB
11750769 s at - CCR2
11716846 a at - MS4A6A
11746962 x at - HLA-DMB
11735352 x at - SLC37A2
11731676 s at - CCR2
11740531 a at - CLEC10A
11745500 a at - CLEC10A
11746961 a at - HLA-DMB
11750856 s at - CCR2
11749803 s at - ARL8B
11754863 s at - VPS37C

Cluster 8

11749803 s at - ARL8B
11754863 s at - VPS37C
11745421 a at - BID
11755603 a at - ARL8B
11749021 s at - CTNNB1
11731110 at - ADAM17
11716718 at - GLA
11720458 at - CAPRIN1
11756634 a at - GSR
11721502 at - SWAP70
11719354 s at - SH3BP5
11736273 a at - TLR2
11747640 a at - TDP2
11743066 s at - PTDSS1
11722031 s at - RNF11
11747482 a at - HNRNPR
11726583 s at - CKAP4
11721801 at - ARL8B
11742819 at - PLAUR
11718648 s at - ACVR1
11722705 at - B3GN15
11720771 x at - PPIF
11722728 a at - EGR2
11744680 a at - SLC7A11
11739475 s at - NAB1
11752899 s at - SEC62
11764019 s at - NAB1
11753204 a at - PTDSS1
11719368 s at - MAPKAPK2
11717326 at - KLF9
11721448 a at - ETF1
11747387 a at - CTNNB1
11717526 a at - RDX
11720370 x at - RTCA
11729163 a at - BTG3
11717525 s at - RDX
11721306 at - PHLDA1
11739366 at - ASPH
11717582 a at - RGCC
11717566 at - EMP1
11719355 s at - SH3BP5
11721305 a at - PHLDA1
11720770 a at - PPIF

Cluster 9

11741350 a at - ANXA11
11757303 s at - BNIP3L
11741103 a at - ANXA11
11754666 a at - CORO1A
11742944 a at - CORO1A
11718001 a at - C10orf54
11715817 at - ZFP36L2
11715818 at - ZFP36L2
11754681 s at - HLA-DMB
11717012 s at - VAMP8
11716377 s at - GRN
11757420 s at - ZFP36L2
11748543 a at - TXNIP
11759666 x at - HLA-DPB1
11724682 s at - TGFBR2
11756073 x at - HLA-DPB1
11756393 x at - CORO1A

Cluster 10

11750538 x at - TXN
11753775 x at - TXN
11731454 a at - SEC62
11721876 s at - SLC7A11
11720772 s at - PPIF
11753804 x at - TXN
11715470 x at - TXN

Cluster 11

11756431 s at - TXNIP
11759642 x at - HLA-DRB1

Cluster 12

11731450 s at - SEC62
11719819 s at - EMC10
11759080 at - GSR
11755523 a at - RBM47
11722167 s at - CTNNB1
11754833 a at - TLR2
11740390 a at - RBM47
11721802 s at - ARL8B
11715649 s at - TXNRD1
11721875 at - SLC7A11
11717565 s at - EMP1
11754026 a at - CXCL8
11763226 x at - CXCL8