Cancer-associated adipocytes promote the invasion and metastasis in breast cancer through LIF/CXCLs positive feedback loop

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Names	Targets	Concentration (µM)
Stattic	Stat3	10
LY3214996	ERK1/2	2
PD98059	MEK1/2, ERK1/2	10
SB 203580	p38, p38β2	10
LY294002	ΡΙ3Κα, ΡΙ3Κδ, ΡΙ3Κβ, CK2, CK2α2,	5
	DNA-PK	
JSH-23	NF-κB	10
BAY 11-7082	NF-κB, USP7, USP21	5
T-5224	AP-1, c-fos	10
TK216	ETS	1
TAT-DEF-Elk-1	Elk-1	20
SB225002	CXCR2	0.1
Reparixin	CXCR1, CXCR2 (the efficiency of	0.1
	inhibiting CXCR1 activity is 400 times that	
	of CXCR2)	
SB-505124	TGF- β Receptor type I receptor	5
EC330	LIF	1

Supplementary Table 1 Concentration and target of inhibitors.

Su	pplementary	Table 2	The sec	juences of	primers for	or q-PCR.
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Species	Genes	q-PCR Primers (5'-3')		
	GAPDH	F-CATCTTCCAGGAGCGAGACC		
		R-CTCGTGGTTCACACCCATCA		
	S18	F-ATCACCATTATGCAGAATCCACG		
		R-GACCTGGCTGTATTTTCCATCC		
	LIF	F- CCAACGTGACGGACTTCCC		
		R- TACACGACTATGCGGTACAGC		
	LIFR	F- AGATATGCCCTTGGAATGTGC		
		R- TCCACTCTTCGAGACCAGAAAA		
Human	gp130	F- GTGAGTGGGATGGTGGAAGG		
		R- CAAACTTGTGTGTGTGCCCATTC		
	CCL2	F- CAGCCAGATGCAATCAATGCC		
		R-TGGAATCCTGAACCCACTTCT		
	CCL20	F- CAGCACTCCCAAAGAACTGG		
		R- CACTGACATCAAAGCAGCCA		
	IL-1β	F-AGCTACGAATCTCCGACCAC		
		R- CGTTATCCCATGTGTCGAAGAA		
	IL-6	F- CCTGAACCTTCCAAAGATGGC		
		R- TTCACCAGGCAAGTCTCCTCA		
	IL-11	F- GGACCACAACCTGGATTCCCTG		
		R-AGTAGGTCCGCTCGCAGCCTT		

CXCL1F-TCCTGCATCCCCCATAGTTA R-CTTCAGGAACAGCCACCAGTCXCL2F-GAAGCTTGTCTCAAGCCCG R-TCTCCTAAGTGATGCTCAAACACXCL3F-GAGCACCAACTGACAGGAGA R-AGTCCTTTCCAGCTGTCCTACXCL8F-TTTTGCCAAGGAGTGCTAAAGACXCL8F-TTTTGCCAAGGAGTGCTAAAGACXCR2F-AGCACTCATCCCAGAATCACTACXCR2F-AGCACTCATCCAGAACCTCAGGACXCR3F-GCGCTCTCCTGAACCTGAGTA R-AGCCTTGCAACCCAGTACXCR4F-GCGCTCTCCTGAAGACACCCTTCXCR5F-GCGCTGCACACACCCTTCSF3F-GAGCAAGTGAGGACATCCAG R-CAGCTTGAGGTGGCACACASTC-1F-GTGGGGCCCCAAAACTCAG R-GTGGAGCACCTCCCGATGGPTX3F-AGCATGGTGTTTTAGTGCC R-ATGGGTTGGAAGGCTTPAR-γF-ACCAAAGTGCAATCAAGTGGA R-ATGGGTGTTGGACTCGGPSFF-AGGAGCCACCACCACCTT R-AGCTTCTGGTGTGAACCGGAPAR-γF-AGGAGCCAGCATTGAGACAAA R-CGCAGGGCTTCCCCTT R-AGGTGTGTGTACCCGGAATCAAGTGCA R-CTGGTGGAAGTCCATTPAR-γF-ATAGGCTGGGCTTCCCCTT R-AGGAGCCAGCATTGAGCACAA R-CGCAGGATTGAGCACAAA R-CGCAGGATTGACCCAGAACCATPAR-γF-AGGAGCCAGCATTGAGCACAAA R-CGCAGGAATTGACCCAGAACCATPAR-γF-GCTGGGAAGTGACGCCTTCXCL2F-GCTTACTGGGGACACCTT R-CCGTACGGAACCACTACGGAAGACAAA R-CGAGGCACATCAGGTACCACCACCACACGGAAGACAAA R-CGAGGCACATCAGGTACCACCACACACACGATTGAGCACCACCACACGGATGGACACACAC			
R- CTTCAGGAACAGCCACCAGT CXCL2 F. GAAAGCTTGTCCAAACCAACCA R- TCTCTAAGTGATGCTCAAACA CXCL3 F. AGTCCTTTCCAGCAGGAGA R- AGTCCTTTCCAGCTGTCCCTA CXCL8 F. TTTTGCCAAGGAGTGCCAAAGA R- AGTCCTTTCCAGCTGTCCCTA CXCR2 F. AGCCATCCTGACCCAGATTA R- GGCCTTTCCAGCCCAGATCACTA R- GGCCTTTCCACCCAGATCAGCA CSF2 F. GCGCTCCTGAAGCAACCAGCA STC-1 F. GTGGCGCCCAAAACTCAG R- GTGGAGCACCTCCGAATCG R- GTGGAGGCACCCCCGAATCG PTX3 F. AGGCTTGAGTCCTTTAGTGCC R- ATGAGGAGAGTTCGAAAGCAACAGCACAGC R- ATGAGGCAGGCTCCCCTT PAR-7 F. AGGCTTGGGCTGGAACACAGA R- GTGCAGCCAGCCATCGAGAAAACCAGA R- GTGCAGCGGGCTTCCCCTT R- BAPA F. AGCCTTTCGGGGCTCACCCTT R- BAPA F. AGCGTTCGGGGCTTCCCCTT R- GGGGGCTGCGCAGCAGAATTGAGGA R- GCGAGGTGTGGACTCGGGAAGAA R- CGTACTTGGGGAAGTCACACAGCACCTAT R- GTGCCCCCGCAGAATTGACGAAAA R- CCCTTACTTGGGGAAGTCACCAT R- CGGAGGCACCCTAT R- GCGGAGCTACCCCCCAGGAATTGACGAAAAAAAAAAAAA		CXCL1	F- TCCTGCATCCCCCATAGTTA
CXCL2F- GAAAGCTTGTCTCAACCCCG R- TCTCTAAGTGATGCTCAAACACXCL3F- GAGCACCAACTGACAGGAGA R- AGCCTTTCCAGCTGTCCCTACXCL8F- TTTTGCCAAGGAGTGCTAAAGA R- AACCCTCTGCACCCAGTTTTCCXCL8F- TTTTGCCAAGGAGTGCTAAAGA R- AGCCTTGTCACCTGAACCTGAGACXCR2F- AGCACTCATCCCAGAATCACTA R- GGCCTTCCTGAACCTGAGAACXF2F- GGCTCTCCTGAACCTGAGAA R- ATGCTGAAGACTCAGGA R- CAGCTTGTAGGCACACACSF2F- GAGCAAGTGAGGAAGATCCAG R- CAGCTTGTAGGCACACASTC-1F- GTGGCGGCTCAAAACTCAG R- GTGGCAGCACTCCGAATGGPTX3F- AGGCTTGAGTCTTTTAGTGCC R- ATGGATTCCTCTTGTGCCATAGPAR-7F- AGGCTTGAGTCGAATCAAGTGGAR- TGTAATCGGGGGTTGACTCGGHSLF- AGGAGCCAGCCACCTAT R- AGCTTTCGGTGTGACTCAGCTTCPREF1F- CTTCCGGCCAGGAATTGAGCACAT R- TGTCATCCTCGCAGAATTGAGCG R- CCGTAGTTGAGTCACCATPAR-7F- CTGCGGGATCACCAGAACAAA R- CGCAGGTGTTGACTCAGCTTMouseL-6R- GGGGCCACCTAAGGACACT R- AGCTGCCTTAGCCACCATMouseL-6R- GGGGCTTGCCCTAAGGAACAA R- GGAGGCACCATTGGCACACAT R- ATGGTCTTGGTCGAATGCCACCATMouseL-6R- GGCTGTGCCTTAACGCACCACCTAT R- CCCTGTAATGCCCACAAGAAACT R- CCCCTGTAATGCCCACCACAGAGAACT R- ACCAAGGGTTCCCCCAAGAGAACT R- ACCAAGGGGTCCTTACCCCACAAGMOUSELIFF- CCCCCTGTAAATGCCACCACAGAGAACT R- ACCAAGGGGTCCTTACCCCACAGAGAACT R- ACCCAAGGGCTCTGACACCACAGAGAACT R- ACCCAAGGGCTCTTACCCCACAAGPAR-7F- CTCCCAAGAATACCAAAGTGCGAACCACACAGAGAACACCACCACACGACGACCACCAC			R- CTTCAGGAACAGCCACCAGT
R- TCTCCTAAGTGATGCTCAAACA CXCL3 F. GAGCACCAACTGACAGAGAGA R- AGTCCTTTCCAGCTGTCCCTA CXCL8 F. TTTTGCCAAGGAGTGCTAAAGA R- AACCCTCTGCACCCAGTTTTC CXCR2 F. AGCACTCATCCCAGAATCACTA R- GGGCTTTCACCTGAGGACA CXF2 F. GCGTCTCCTGAACCTGAGGACA CSF2 F. GCGCAGTGAGGAGAACCCAG R- CAGCTTGTAGGTGGCACACA R STC-1 F. GTGGACGCCCCAAAACTCAG R- CAGCTTGTAGGTGGCACACA R STC-1 F. GTGGACGCCTCTAAACTCAG R- CAGCATGGAGGAGTCGAAAGTGGA R PTX3 F. AGGAGCAGTGGAGATCACAG R- ATGAGGGAGTTGGAAAGCCATGG R PPAR-7 F. ACCAAAGTGCAATCAAAGTGGA R- ATGAGGGCAGTTGGAAGGCTCT C/EBP-a R- ATGAGGGCAGCATGAGAGACCAT R R- GTGCTGGGGCTTCCCCTT R R- ATGAGGCCAGCATTGAGACAAA R R- CGCAAGGTGTGACCGGGACACCTT R R- CGCAGGGTTGGACCGGAAGAA R R- CGCAGGGCTTCCCCTT R R- CCCAGGGTGTGACCCCT R R- CCCTTACATGGGAACACACAT R R- CCCGTACATCACGCACACAGAAAA R		CXCL2	F- GAAAGCTTGTCTCAACCCCG
CXCL3 F- GAGCACCAACTGACAGGAGA R- AGTCCTTTCCAGCTGTCCCTA CXCL8 F- TTTGCCAAGGAGTGCTAAAGA R- AACCCTCGCACCCAGTTTTC CXCR2 F- GGGCTTTCACCTGAAGCAATCACTA R- GGGCTTTCACCTGAACCGAGAT CSF2 F- GGGCACCAACCCGAGAT CSF3 F- GAGGAGCACCTCCGAACACA STC-1 F- GTGGAGGCACCTCCGAATGG R- GTGGAGCACCTCCGAATGG R PTX3 F- AGGCTTGAGTCTTTAGTGCC R- ATGAGGGAGTTGGAAAGGCTCT C/EBP-a PAR-γ F- TATAGGCTGGGCTTCCCCTT R- ATGAGGGAGTTGGAAGGCTCT C/EBP-a R- ATGAGGGGCTTCCGGGCTTCCCCTT R- ATGAGGGCAGTTGGAAGGCTCT C/EBP-a F- TATAGGCTGGGCTTCCCCTT R- ATGAGGGAGTTGGAAGGCTCT R- ATGAGGCAGGCACTCCGG PAR-γ F- AGGAGCCAGCACCAGGAATCAA R- CCCAAAGTGGAAGACCCTT R- GTGCCAGGAATTGAGCA R- CCCCTTACTCGGCAAGGAACCAT R- CCGTTACTTGGGGAACCCAC PAR-P F- GCTGTCCCTCAACGGAAGAA R- CCCCTTACTTGGGGCATCACCTCAAGGAA R- CCGTGTGCTTAGCCACC Mouse IL-6 F- AGTCCTTCCTACCCAAGGAACAC R- GTGCTTGGTCCTAAGCACCACC R- CCCCCTGTAAAGCACACACAGAACACT R- CCCCCTGTAAAGCCACCA			R- TCTCCTAAGTGATGCTCAAACA
R-AGTCCTTTCCAGCTGTCCCTACXCL8F-TTTTGCCAAGGAGTGCTAAAGA R-AACCCTCGCAGAGTCCCAGAATCACTA R-GGCCTTTCACCCGGAGACACACXCR2F-AGCACTCCAGAACCAGGACACSF2F-GCGCTCCCGAACCTGAGGACACSF3F-GAGCAAGTGAGGAGACACASTC-1R-GTGGAGCACCTCCGAATCGAPTX3F-AGCCTTGAGGGACACAAPTX3F-AGCCTTGAGGGACACAAAPAR-γF-ACCAAAGTGCAGAAGTCCGGAPAR-γF-ACCAAAGTGCAGCACAAARSLF-AGGCTTCGGGCTTCCCCTTRSLF-AGGAGCCACCACAAAGTGCGAPAR-γF-AGGAGCCACCACAAAGTGGARSLF-AGGAGCCACCACAAAGTGGAAGCACAAARCGCAGGTGTGAATCAAGCTTCPREF1F-CTTTCGGCCACAGCACCTATRAGGCTGGCAGTGAAGCACAAARCGCAGGTGTGAATCCAGCTTCPREF1F-CTTTCGGCCACGACCACTATRAGGCAGCACTACCGGAATTGACGRCCGTACTCGGGAATTGACGRCCGTACTCGGGAATTGACGRCCGTACTCGGGAATTGACGRCCGTACTCGGGAATTGACGATCXCL3F-CGTGCCTCCACCGAAGAAR-GGGTGCCTTGGTGTAATCCACGTGCACCACMouseIL-6IL-6F-AGTCCTTCTCCCCCAAGGAACCTTRCCCGTAATGCCCCTTGCCCGTGGCATCXCL2F-CCCTGTGAATGCCACCTGTMOUSEIL-6R-CTCTGTGCCTTACCCCAAGGACCTGR-CTCTGTGCCCTTGGCATRCCCCCTGTGAATGCCACCTGTRCTCTCTGTCCCCACAAGTCCGACRCCCCCTGTGAATGCCACCTGTRCCCCCCCAATGCCCCCAAGGACCTGTRCCCCCCCCCCAAAGTGCCGARCCCCCTGTGCCATGCCCCCCCCCCAAGGGCAACCCGACCCGRCCCCCAGGCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC		CXCL3	F- GAGCACCAACTGACAGGAGA
 CXCL8 F- TTTTGCCAAGGAGTGCTAAAGA R- AACCCTCTGCACCCAGTTTTC CXCR2 F- AGCACTCATCCCAAGATACACTA R- GGGCTTTTCACCTGTAGGACA CSF2 F- GAGCAAGTGAGGAAGATCCAG R- ATGCTGAAGCTCACCCGAGTA R- CAGCTTGTAGGTGGCACACA STC-1 F- GTGGCGGCTCAAAACTCAG R- CTGGAGCACCTCCGAATGG PTX3 F- AGGCTTGAGTCTTTAGTGCC R- ATGAGTCTGTAGTGGAAGATCCAG PPAR-γ F- ACCAAAGTGCAATCAAGTGGA PPAR-γ F- ACCAAAGTGCAACACAA PPAR-γ F- AGGATTGTGATCACAGCAT R- GTGGAGCACCTCCGAATGG PPAR-γ F- AGGAGCAGCATGAGAAAA R- GGCAGGTGTGAATCAAA R- GGCAGGTGTGAATCAAA R- GGCAGGTGTGAATCAAA R- GGCAGGTGTGAATCCAGCAT PREF1 F- ACGAGGCAAGCACCACA R- CGGAGGTAGACAAA R- CGCTGGGATTCACCTCAAG R- CCGTAGCTGGGACACCAT R- CGCTGGCAAGAAGAA R- CGGAGGTGAGCACCTT CXCL2 F- GCTTACCTGGGAAGAAA R- CGTACTAGGGACACCTT CXCL3 F- CCTACCAAGGATCCAT R- AGGTGGCTTGGCACCAT R- GGTGGCCAAGGAATCCAT R- GGTGGCCAAGGAACCT CXCL3 F- CCTACCAAGGGTGAATTTGAGCA R- GGTGGTCTTGGTCTTAGCCAC R- GGTGGTCTTGGTCTTAGCCACTT CXCL3 F- CCTACCAAGGGTTGATTTGAGAC R- GGTGGTGGTAGACTTCC R- GGCTGTTGGTCCTTAGCCAAT R- CCCAAGGATTCCCGAT CXCR2 F- CAAAACAGCGTGTAGAACT R- CCCAGGAATACCAAAGTGCCAA R- GCCTGGTGCAATCCCACAAGTGCCAA R- GCCTGGTGCAATCCCACAAGTGCCAA R- GCCTGGTGCAATCCCACAAGTGCCAA R- GCCTGGTGCAATCCCACAC 			R-AGTCCTTTCCAGCTGTCCCTA
R-AACCCTCTGCACCCAGTTTCCXCR2F-AGCACTCATCCCAGAATCACTA R-GGCTTTCACCTGAGGACACSF2F-GGCTCTCGAACCTGAGTA R-TGCTGAAGCTCACACCCTTCSF3F-GACCAAGTGAGAGAATCCAG R-CAGCTTGAGGTGGCACACASTC-1F-GTGGCGGCTCAAAACTCAG R-CAGCTTGAGGTCCTTAGTGCCPTX3F-AGCCTTGAGCTCCCCAAAGTGPTX3F-AGCCTTGAGCTGGACACAAGPPAR-γF-AGCATGGAGCACTCCCCTT R-AGGGTGGGACTCCCCTTPAR-γF-TATAGGGAGCTGGAAGACTCGGHSLF-AGGAGCAGCATTGAGACAAA R-CGCAGGTGTGAACTCAGGPREF1F-CTTCGGCCACAGCATTGAGACAAA R-CCGTAGGTACCAGCACCTTPAR-γF-ACGGGCCAGGATTGAGACAAA R-CGCAGGTGTGAACTCAGGHSLF-AGGAGCCAGCATTGAGACAAA R-CGCAGGCACCTCCCCTTCXCL1F-TGCCTGGGATTCACCTCAAG R-CCGTACTGGGACACACTMouseIL-6ILFF-AGTGGCCAAGGATTGAGACAA R-CGCTGTCTCCTTGGGGACACCTTCXCL2F-CCACAACGGTGTGATTTGAGAC R-GCTGTCCTCCACCCAATTCC R-GCTGTCCTCCCCCAATTCC R-GCTGTCCTCCCCCAATTCC R-GCTGTCCTCCCCCAATTCC R-GCTGTCCTTGCCGTAGAACT R-CCCAAGGGTTGACTTCCGGAACCTGMouseIL-6PAR-γF-CTCCCAAGAATACCAAAGTGCCA R-GCCTGTGCCTTAGCCACCTG R-GCCTGTGCCTTAGCCACCTGPAR-γF-CTCCAAGAATACCAAAGTGCCAA R-GCCTGATGCCTTATCCCCCCAAGTACCGA R-GCCTGATGCCTTATCCCCCCAACTCCGCACCCPAR-γF-CTCCAAGAATACCAAAGTGCCAACT R-GCCTGATGCCTTATCCCCCCCCCCACAGCACCGPAR-γF-CTCCAAGAATACCAAAGTGCCAACT R-GCCTGATGCCTTATCCCCCCCACACGCACCGCYEBP-aF-CTCCAAGAATACCAAAGTGCCAACCGACCGPAR-γF-CTCCAAGAATACCAAAGTGCCACCGPAR-γF-CTCCAAGAATACCAAAGTGCCAACCGACCGPAR-γF-CTCCAAGAATACCAA		CXCL8	F- TTTTGCCAAGGAGTGCTAAAGA
CXCR2F- AGCACTCATCCCAGAATCACTA R- GGGCTTTCACCTGAGGACACSF2F- GCGTCTCCTGAACCTGAGTA R-ATGCTGAAGCTCACACCCTTCSF3F- GAGCAAGTGAGGAAGATCCAG R- CAGCTTGTAGGTGCCACACASTC-1F- GTGCCGCGCTCAAAACTCAG R- GTGGAGCACCTCCGAATGGPTX3F- AGGCTTGAGTCTTTAGTGCC R- ATGGATTCCTCTTTGTGCCATAGPPAR-γF- ACCAAAGTGCAATCAAGTGGA R- AGGTTGTGGTGGACTCGGHSLF- AGGAGCCAGCACTCGGAATGCGHSLF- AGGAGCCAGCACCTAT R- AGGTTGTGATCAGCTTCPREF1F- CTTTCGGCCACAGCACCTAT R- TGTCATCCTCGGAATCACAGFABP4F- ACTGGGCCAGGAATTGAGCATCXCL2F- GCTGTCCCTCAACGGAAGCACTTMouseIL-6LIFF- CCTTCCGGCACAGGACCTGT R- GGTGTCCTTGGTGCATTCAGCTMouseIL-6PAR-γF- CCTACCAAGGGTTGATTCTGGGAGTGTGTCTCTCTTCGGCAACCAGA R- CCGTGCCACAGCACCTTCXCL2F- GCTGTCCTCAACGGAAGAA R- CGAGGCACATCAGGTAGACT R- GGTGTCCTTAGCCACMouseIL-6AGGTGGCTATGACTCTGGCAACCAGGAACAGA R- CCCGTTAATTGCCACCCAAGGMOUSEILFACCAAAGAGCGCTCCAAGGTAGAACT R- CTCTCTGTCCCCCACAGGTAGAACT R- CCCTGTAAATGCCACCTGT R- CTTCTGTCCCCCCACAGGTAGAACT R- ACCAAGGAGTCCCCACAGGMOUSEILFACCAAGAACAGCGCGCGTAGAACT R- ACCAAGGAGTCCCCCACAGGTACCACCTGT R- ACCAAGACAGCACCAGGTCCCCACAGACCAAGGAGTCCCCCACAGGPAR-γF- CTCCCAGAAACAGCGCGCGTAGAACT R- ACCAAGGAGTCCCCACAGGTACCACGAGACCACGAGTACCAAGGCACCACGAGGACTCCEBP-aF- CAAGAACAGCAACAGCGACCACCACGGACACCACCACCAC			R-AACCCTCTGCACCCAGTTTTC
R- GGGCTTTTCACCTGAGGACACSF2F- GCGTCTCCTGAACCTGAGTA R-ATGCTGAAGTCACACCCTTCSF3F- GAGCAAGTGAGGAGAACACCAG R- CAGCTTGTAGGTGGCACACASTC-1F- GTGGAGCACCTCCGAATGGPTX3F- AGGCTTGAGTCTTTTAGTGCC R- ATGGATTCCTCTTTGTGCCATAGPPAR-γF- ACCAAAGTGCAATCAAGTGGA R- ATGAGGGGCTTCCCCTTC/EBP-αF- TATAGGCTGGCACACCACGGHSLF- AGGACCACCTCGGAATCCAAAGTGGA R- AGCTTCTGGTGGACTCCGGHSLF- AGGAGCCAGCATTGAGACAAA R- CGCAGGTGTTGAATCAACATCAAGPREF1F- CTTTCGGCCACAGCACTAT R- TGTCATCCTCGCAGAATCCATFABP4F- ACTGGGCCACGAATCAAGTGACGATCCCL2F- GCTGTCCCTCAACGGAAGAA R- CGGTGGCATGACGCTTMouseIL-6ILFF- CCTCTCCACCAAGGAACAA R- CGCTGGCAATCAGTTCTGGGAACACTMouseIL-6PAGTCTTCTGGTCCTTAGCCAC R- GGTCTCTCTCTCCCCCAAGGACTMOUSEILFPAR-γF- CCTCCTGAAATGCCACCTCCC22F- CCTCCTGAAATGCCACCTCCC23F- CCTCCTCAACGGAAGAAR- GGCTGGCTTGACTCTCTCTCTGCGGAACCTCCC22F- CCTCCTCAACGGAAGAAR- GGTGTCCTCTACCCCAATTCCR- GGCTGGCTTGACTTAGCCACCTGTR- CCCCTGTAAATGCCACCTGTR- CCCCTGTAAATGCCACCTGTR- CCCCCTGTAATGCCACCTGTR- CCCCCTGTAATGCCACCTGTR- CCCCCTGTAAATGCCACCTGTR- CCCCAGGAGTCCCCACAAGTGCGAR- CCCCCTGTAAATGCCACCTGTR- CCCCCGTAGAATACCAAAGTGCGAR- CCCCTGTAAATGCCAACTGCAAGGACTCCCCACAGPAR-γF- CCCCCAGAATCCAAAGTGCGAAGAAGAACAAGCACAGAGTGCGAACACAGAGTACCAACGCACCACACAGGATCCCACACCACACACA		CXCR2	F- AGCACTCATCCCAGAATCACTA
CSF2F- GCGTCTCCTGAACCTGAGTA R-ATGCTGAAGCTCACACCCTTCSF3F- GAGCAAGTGAGGAAGATCCAG R- CAGCTTGTAGGTGGCACACASTC-1F- GTGGCGGCTCAAAACTCAG R- GTGGAGCACCTCCGAATGGPTX3F- AGGCTTGAGTCTCTTAGTGCC R- ATGGATTCCTCTTTGGCCATAGPPAR-γF- ACCAAAGTGCAATCAAAGTGGAPPAR-γF- ACCAAAGTGCAATCAAGTGGAHSLF- AGGCTTCGGGGCTTCCCCTT R- AGCTTTCTGGTGTGACTCGGHSLF- AGGAGCCAGCCAGCAGTGAAGACAAAPREF1F- CTTCGGCCACAGAATCAAAFABP4F- ACTGGGCCACCAGAATCAAAR- CTCGTGGAAGTGCACCTAT R- TGTCATCCTCGAGAATCCATFABP4F- ACTGGGCCACGAGATCACTCXCL1F- TGGCTGGGATTCACCTCAAG R- CCGTTACTTGGGGACACCTTMouseIL-6IL-6F- AGTCCTCCCCAAGGAGTGACCATCIL-6F- AGTCCTTCCTCCACCAATTCAGCAC R- CGTCTCCTCACCCAATTCCMOUSEIL-6PAR-γF- CCCCTGTAAATGCCACCTFAGAGTCTGTGGTCTTAGCCACFAGAGTGCTTAGCCCCAATTCC R- GGTGCTTAGCTCCCCAAGGATCACTFAGAGTGCTTACCCCAAGGGTCGAACGAC R- GGTGCTGGCCTTGCCCACAGACACCACTFAGAGACACAGCAACGCACCCTGT R- CCCCTGTAAATGCCACCTGT R- CCCCTGTAAATGCCACCTGTFAGAGACAGCAACGGGTCGTAGAACT R- ACCAAGGAGTTCCCCCAAAGPPAR-γF- CCCCTGTAAATGCCACCTGT R- ACCAAGGAGTTCCCCCAAAGFAGCCTGGTGGAACGCACCGGAGACGAGACGAGACGACGACGACGACGA			R-GGGCTTTTCACCTGTAGGACA
R-ATGCTGAAGCTCACACCCTTCSF3F- GAGCAAGTGAGGAAGATCCAG R- CAGCTTGTAGGTGGCACACASTC-1F- GTGGCGGCTCAAAACTCAG R- GTGGAGCACCTCCGAATGGPTX3F- AGGCTTGAGTCTTTTAGTGCC R- ATGGAGTCCTCTTGTGCCATAGPPAR-γF- ACCAAAGTGCAATCAAAGTGGAC/EBP-αF- TATAGGCTGGGCTTCCCCTT R- AGCAGCAGCATGAGACAAAHSLF- AGGAGCCAGCATTGAGACCAAT R- CTCGGCCAGACATTGAGCTCCPREF1F- CTTTCGGCCAGACACCAAT R- CTCGTGGAATCCACTTFABP4F- ACTGGCCAGGATTGACGC R- CTGGGCCAGGAATTGACGKCCL2F- GCTGTCCCTCAACGAAGAA R- CCGAGGCTAGGACACCTTMouseIL-6ILFF- CCCCTGTAAATGCCACCTGT R- GTCTCCTGCCGTAGAACCATTCC R- GGTCTTCGTCCCTAACGGAACCTGMouseIL-6PAR-γF- CCCCTGTAAATGCCACCTGT R- GCTGTCCCTCAACGGAAGACCTGMOUSEILFPAR-γF- CCCCTGTAAATGCCACCTGT R- GCTGTCCTCTGCCGTGGCATMOUSEILFPAR-γF- CCCCTGTAAATGCCACCTGT R- CCCAAGGAGTCCCCAAGGAACCTGT R- CCCCTGTAAATGCCACCTGTPAR-γF- CCCCTGTAAATGCCACCTGT R- CCCAAGAGATTCCCAAGGGAACCTGGGAAACCT R- CCCAAGAACAGCGTGCGTAGAACT R- GCTGATGCCTTAGCCCACAAGGAGAACCT R- GCTGATGCCTTAGCCCACAAGGAGTACCGAGACCCTGT R- CCCAAGAACAGCAGAGTACCGAGAACCGAGAGAACCCAAGGAGTCCCCACAAGGAGTCCCCACAAGGAGTCCCCACAAGGAGTACCGAGACCCCCACAAGGAGTCCCCACAAGGAGTCCCCACAAGGAGTCCCCACAAGGAGTACCGAGACCCGAGACCCGAGACCCGAGACCGAGACCCGACCCGACCCGACCCGACCCGACCCCCC		CSF2	F- GCGTCTCCTGAACCTGAGTA
CSF3F- GAGCAAGTGAGGAAGATCCAG R- CAGCTTGTAGGTGGCACACASTC-1F- GTGGCGGCTCAAAACTCAG R- GTGGAGCACCTCCGAATGGPTX3F- AGGCTTGAGTCTTTAGTGCC R- ATGGATTCCTCTTTGTGCCATAGPPAR-γF- ACCAAAGTGCAATCAAAGTGGA R- ATGAGGGGTTGCACTGGC/EBP-αF- TATAGGCTGGGCTTCCCCTT R- AGCTTCTGGTGTGACTCGGHSLF- AGGAGCCAGCATGAGACAAA R- CGCAGGTGTTGATCAGCTTCPREF1F- CTTTCGGCCACAGCACCTAT R- ACTGGCCAGGAATTGACGCGFABP4F- ACTGGGCCACGAATTGACG R- CCGTGGCATCAGGAAGAA R- CCGAGGTGTAGACGCATMouseIL-6IL-6F- AGTCCTTCGTGGACACCAT R- CTCTCGTGCCATAGCACCATMouseIL-6ILFF- CCTCCTGGAATGCCCCTT R- CCTCTGGTGCATTCCCCCAAGGACCTTMOUSEIL-6F- CCTCCTGGAATGCCCCCGT R- CCTCTCGTGCCATAACGCACCTGT R- CCTCTCTGCCCCTAACGCACCTGT R- CCTCTCTGCCCCTTAGCCAC R- CCTCTCTGCCCGTTGCCATMOUSEIL-6F- CCCCTGTAAATGCCACCTGT R- CCCCTGTAAATGCCACCTGT R- CCCCTGTAAATGCCACCTGT R- CCCCTGTAAATGCCACCTGT R- CCCCTGTAGAATTCCCAAAGTGCGA R- CCCAAACAGCGTGTAGAACT R- ACCAAGGAGTTCCCCCACAG R- GCCTGATGCCTTATCCCCACAF- CACAAACAGCATCCAAGGAT R- GCCTGATGCCTTATCCCACA R- GCCTGATGCCACCCG R- GCCTGATGCCACCCGACCCG			R-ATGCTGAAGCTCACACCCTT
R- CAGCTTGTAGGTGGCACACASTC-1F- GTGGCGGCTCAAAACTCAGR- GTGGAGCACCTCCGAATGGR- GTGGAGCACCTCCGAATGGPTX3F- AGGCTTGAGTCTTTTAGTGCCR- ATGGAGTCCTCTTTGTGCCATAGR- ATGGAGGGAGTTGGAAGCCTCTPAR-γF- ACCAAAGTGGAATCAAAGTGGAR- ATGAGGGAGTTGGAAGCCTCTR- ATGAGGCAGCATTGAAGCCCCTTC/EBP-aF- TATAGGCTGGGCTTGCACTGGGHSLF- AGGAGCCAGCATTGAGACAAAR- CGCAGGTGTGATCAGCTTCR- CGCAGGTGTGATCAGCTTCPREF1F- CTTTCGGCCACAGCATCAGATFABP4F- ACTGGGCCAGGATTGAGCGCCTTCXCL1F- TGCCTGGGATTCACCTCAAGR- CCGTTACTTGGGGACACCTTR- CCGTTACTTGGGGACACCTTCXCL2F- GCTGTCCCTCAACGGAAGAAR- CGAGGCCATGAGTGACTGACTR- CGAGGGCACATCAGGTAGATMouseIL-6F- AGTCCTTCCTACCCCAATTCCILFF- CCCCCTGTAAATGCCACCTGTCCCR2F- CCCCCGTAAATGCCACCTGTR- CTCTCTGTCCCGTTGGCATR- CCCCCGTTAGACACCACILFF- CCCCCGTAAATGCCACCTGTR- CTCCTGTCCCCACAAGGGTGCAAAGACTR- CTCCAGAAATACCAAAGTGCGAR- ACCAAGAGGTTCCCCACAAGPPAR-γF- CTCCAAGAATACCAAAGTGCGAR- GCCTGATGGCTTATCCCACAAGPPAR-γF- CTCCAAGAATACCAAAGGAGACCGR- GCTGATGGCTTATCCCACAAGC/EBP-aF- CAAGAACAGCAACGAATACCAAAGTACCGR- GCETGATGGCTTATCCCACACR- GCTGATGGCTTATCCCCACACC/EBP-aF- CAAGAACAGCAACGAATACCAAAGTACCGR- GCCTGATGGCTCAACCGAGACCGR- GCCTGATGGCACCGGACCG		CSF3	F- GAGCAAGTGAGGAAGATCCAG
STC-1F- GTGGCGGCTCAAAACTCAG R- GTGGAGCACCTCCGAATGGPTX3F- AGGCTTGAGTCTTTAGTGCC R- ATGGATTCCTCTTTGTGCCATAGPPAR-γF- ACCAAAGTGCAATCAAAGTGGA R- ATGAGGGAGTTGGAAGGCTCTC/EBP-αF- TATAGGCTGGGCTTCCCCTT R- AGGATTCAGCACCAGCATTGAGACAAA R- CGCAGGGTTGAATCAGCACCAA R- CGCAGGATTGAGACAAA R- CGCAGGCATTGAGACCACTAT R- TGTCATCCTCGCAGAATCCATPREF1F- AGGAGCCAGCATTGAGCACCAT R- CTCGTGGAAGTGACCCCTTFABP4F- ACTGGGCCAGGAATTGACCAAG R- CCGTACTGGGACACCTTCXCL1F- TGGCTGGGAATCCATCAGCACCTA R- CCGTACTGGGAACACCTTMouseIL-6IL-6F- AGTCCTTCCTACCCCAAGGAAC R- GGTGGTCTTGGTCCTTAGCCACILFF- CCCCTGTAAATGCCACCTGT R- CCCTGTGAATGCCCCCATMouseIL-6PAR-γF- CACAAACAGCGTCGTAGAACT R- CCCCGTGAATCCCAAAGTGCGA R- GCCTGATCCCCAAAGTGCGAACCG R- GCCTGATCACCAAAGTGCGACCG R- GCCTGATACCAAAGTCCCACA C/EBP-αFCACAAGAACGCAACGAGTACCG R- GCCTGGTCAACGAACAACGACGACCG			R- CAGCTTGTAGGTGGCACACA
PTX3R-GTGGAGCACCTCCGAATGGPTX3F-AGGCTTGAGTCTTTAGTGCCR-ATGGATTCCTCTTTGTGCCATAGPPAR-γF-ACCAAAGTGCAATCAAAGTGGAR-ATGAGGGAGTTGGAAGCCTCTC/EBP-αF-TATAGGCTGGGCTTCCCCTTR-AGCTTTCTGGTGTGACTCGGHSLF-AGGAGCCAGCATTGAGACAAAR-CGCAGGTGTTGATTCAGCTTCPREF1F-CTTTCGGCCACAGCACCTATR-BP4F-ACTGGGCAGGAATTGACGR-CCCGTGGAATCCACCTFABP4F-ACTGGGCAGGAATTGACGCCTTCXCL1F-TGCCTCCCCAACGGAACCCTTCXCL2F-CCTACCAAGGATCACCTCAAGR-CGAGGCACATCAGGTACGATCXCL3F-CCTACCAAGGGTTGATTTGAGACMouseIL-6LIFF-CCCCTCTAAATGCCAACGATCXCR2F-CCCCTGTAAATGCCACCTGTR-GGAGGCATGACCTTCGGTGGCATCXCR2F-CCCCTGTAAATGCCACCTGTR-CCCAGAATACCAAAGTGCCACLIFF-CCCCTGTAAATGCCACCTGTR-CCAAACAGCGTCGTAGAACTR-ACCAAGGAGTTCCCCACAAGPPAR-γF-CTCCAAGAATACCAAAGTGCGAR-GCTGGTGCAACGCAACAAGAGAACTR-GCCTGATGCTTATCCCCACAC/EBP-αF-CAAACAGCAACGAGTACCGR-GTCACTGGTCAACCGACACGAGTACCGR-GTCACTGGTCAACCGACCGR-GCCTGGTCAACCGACACCACGACCACCGR-GCCAGGTCTAACCCAACGACCACCGR-GCCAGGTCGTACAACTGCACCGR-GCCAGGTCACACCACGACCACCGR-GCCAGGTCGACACCACGACCACCGR-GCCTGGTCAACCGACCACCACCACACGACCACCACCACCACACCACC		STC-1	F- GTGGCGGCTCAAAACTCAG
PTX3F-AGGCTTGAGTCTTTTAGTGCC R-ATGGATTCCTCTTTGTGCCATAGPPAR-γF-ACCAAAGTGCAATCAAAGTGGA R-ATGAGGGAGTTGGAAGGCTCTC/EBP-αF-TATAGGCTGGGCTTCCCCTT R-AGCTTTCTGGTGTGACTCGGHSLF-AGGAGCCAGCATTGAGACAAA R-CGCAGGTGTTGATTCAGCTTCPREF1F-CTTTCGGCCACAGCACCAT R-TGTCATCCTCGCAGAATCCATFABP4F-ACTGGGCCAGGAATTTGACG R-CCGTTGCTCCCTCAACGGAAGAA R-CGAGGGATCACCTTMouseLL-6MouseLL-6F-AGTCGTCCTCGTAACGCACT R-CCGTTGGTCTTGGCCACCACTTMouseLI-6F-CCCTTCTTCGTCCCCAAGTTCCCCCAAG R-GCTTGGTCCTTAGCCACCTGT R-CCCGTTACTTGGCCACCTTMouseLI-6F-AGTCCTTCGTCCCTAACGCAACT R-CCCAAGGATCCCCCAATTCC R-CTCCTGTCCCTAACCCAAAGTCCATMOUSELI-6F-CCCCTGTAAATGCCACCTGT R-CTCCTTGGCCGTTGCCAACTFCCCCCTGTAAATGCCACCTGT R-CCCAAGAACT R-CCCAAGGAGTTCCCCACAAG R-GCCTGATGCTTTACCCCACAAG R-GCCTGATGCTTTACCCCACAAG R-GCCTGATGCTTACCCCACAAG R-GCCTGATGCTTACCCCACAAG R-GCCTGATGCTTACCCCACAAG R-GCCGATGCTTTACCCCACAAGTGCGA R-GCCTGATGCTTACCCCACAAGTGCGA R-GTCACTCCCACACGAGTACCG R-GTCACTCCCACACCGACCACCACCACGACT			R- GTGGAGCACCTCCGAATGG
R-ATGGATTCCTCTTTGTGCCATAGPPAR-γF-ACCAAAGTGCAATCAAAGTGGAR-ATGAGGGAGTTGGAAGCCTCTR-ATGAGGGAGTTGGAAGCCCTTC/EBP-αF-TATAGGCTGGGCTTCCCCTTR-AGCTTTCTGGTGTGACTCGGR-AGCTTTCTGGTGTGACTCGGHSLF-AGGAGCCAGCAGCAGCACTATR-CGCAGGTGTTGATTCAGCTTCR-TGTCATCCTCGCAGAATCCATPREF1F-CTTTCGGCCACAGCACCTATR-AGCTTCTGGGGAATTTGACGR-CTCGTGGAAGTGACGCCTTCXCL1F-TGGCTGGGATTCACCTCAAGR-CCGTTACTTGGGGACACCTTR-CCGTGCCCTCAACGGAAGAAR-CGAGGCACATCAGGTACGATR-CGAGGCACATCAGGTACGATMouseIL-6F-AGTCCTTCCTACCCCAATTTCCILFF-CCCTGTAAATGCCACCTGTR-CTCTCTGTCCTGGCCATR-CCCTGTGAAATGCCACCTGTMouseIL-6F-AGTCCTTCCTACCCCCAATTCCILFF-CCCCTGTAAATGCCACCTGTR-CCACAAGGAGTTCCCCACAGGR-ACCAAGGAGTTCCCCACAGGPPAR-γF-CTCCCAAGAATACCAAAGTGCGAR-GCCTGATGCTTATCCCCACAGR-GCCTGATGCTTATCCCCACAGPAR-γF-CAAGAACAGCAACGAGTACCGR-GTCACTGGTCAACCCAACGAGTACCGR-GCCTGATGCTTATCCCCACAC/EBP-αF-CAAGAACAGCAACGAAGAACGAGTACCGR-GTCACTGGTCAACTCCAGCACR-GTCACTGCTCAACCCACGAC		PTX3	F- AGGCTTGAGTCTTTTAGTGCC
$ \begin{split} \mbox{PPAR-$$\gamma$} & \mbox{F-ACCAAAGTGCAATCAAAGTGGA} \\ & \mbox{R-ATGAGGGAGTTGGAAGGCTCT} \\ & \mbox{R-ATGAGGCAGGCTTCCCCTT} \\ & \mbox{R-AGCTTTCTGGTGTGACTCGG} \\ & \mbox{HSL} & \mbox{F-AGGAGCCAGCATTGAGACAAA} \\ & \mbox{R-CGCAGGTGTTGATTCAGCTTC} \\ & \mbox{PREF1} & \mbox{F-CTTTCGGCCACAGCACCTAT} \\ & \mbox{R-TGTCATCCTCGCAGAATCCAT} \\ & \mbox{FABP4} & \mbox{F-ACTGGGCAGGATTTGACGCCTT} \\ & \mbox{R-CTCGTGGAAGTGACGCCTT} \\ & \mbox{R-CCGTTACTTGGGGACACCTAC} \\ & \mbox{R-CCGTGGCAATCAGGTACGAA} \\ & \mbox{R-CCGTGCCTCAACGGAAGAA} \\ & \mbox{R-CCGTGCCTCAACGGAAGAA} \\ & \mbox{R-CGAGGCACATCAGGTACGAT} \\ & \mbox{CXCL2} & \mbox{F-GCTGTCCTCACGGAAGAA} \\ & \mbox{R-CGAGGCACATCAGGTACGAT} \\ & \mbox{CXCL3} & \mbox{F-CCTACCAAGGGTTGATTTGAGAC} \\ & \mbox{R-GAGTGGCTATGACTTCTGTCTGG} \\ & \mbox{Mouse} & \mbox{IL-6} & \mbox{F-AGTCCTTCCTACCCCAATTCC} \\ & \mbox{LIF} & \mbox{F-CCCTGTAAATGCCACCTGT} \\ & \mbox{R-CCACGAGGTTCCCCACAGGAACT} \\ & \mbox{R-CCAAGGAGTTCCCCACAAG} \\ & \mbox{PPAR-$$$$\gamma$} & \mbox{F-CTCCAAGGAATACCAAAGTGCGA} \\ & \mbox{R-GCCTGATGCTTTATCCCCACA} \\ & C/EBP-$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$			R-ATGGATTCCTCTTTGTGCCATAG
R-ATGAGGGAGTTGGAAGGCTCTC/EBP-αF. TATAGGCTGGGCTTCCCCTT R-AGCTTCTGGTGTGACTCGGHSLF. AGGAGCCAGCATTGAGACAAA R- CGCAGGTGTGATTCAGCTTCPREF1F. CTTTCGGCCACAGCACCAT R- TGTCATCCTCGCAGAATCCATFABP4F. ACTGGGCCAGGAATTGACG R- CTCGTGGAAGTGACGCCTTCXCL1F. TGGCTGGGATTCACCTCAAG R- CCGTTACTTGGGGACACCTTMouseIL-6IL-6F. AGTCCTTCCTACCCAATTTCC R- GGTCTTGGTCCTTAGCCACLIFF. CCCCTGTAAATGCCACCTGT R- CCCTGTAATGCCACCTGTMouseIL-6IL-6F. AGTCCTTCCTACCCCAATTTCC R- TGGTCTTGGTCCTTAGCCACLIFF. CCCCTGTAAATGCCACCTGT R- CTTCTGTCCCGTTGCCATCXCR2F. CACAAACAGCGTCGTAGAACT R- CCCCTGTAAATGCCACCAGAPAR-γF. CTCCCAAGAATACCAAAGTGCGAA R- GCCTGATGCTTATCCCCACAC/EBP-αF. CAAGAACAGCAACACGAGTACCG R- GTCACTGGTCAACTCCAGCAC		PPAR-γ	F- ACCAAAGTGCAATCAAAGTGGA
C/EBP-αF-TATAGGCTGGGCTTCCCCTT R-AGCTTTCTGGTGTGACTCGGHSLF-AGGAGCCAGCATTGAGACAAAHSLR-CGCAGGTGTTGATTCAGCTTCPREF1F-CTTTCGGCCACAGCACTATFABP4F-ACTGGGCAGGAATTGACGFABP4F-ACTGGGCATGAGACCCTTCXCL1F-TGGCTGGGATTCACCTCAAGCXCL2F-GCTGTCCCTCAACGGAAGAACXCL3F-CCTACCAAGGGTTGATTTGAGACCMouseIL-6ILFF-CCCTGTAAATGCCACTGTCXCR2F-CCCTGTAAATGCCACCTGTCXCR2F-CCCTGTAAATGCCACCTGTCXCR2F-CACAAACAGCGTCGTAGAACTCXCR2F-CACAAACAGCGTCGTAGAACTCXCR2F-CACAAACAGCGTCGTAGAACTCXCR2F-CACAAACAGCGTCGTAGAACTPAR-γF-CTCCAAGAATACCAAAGTGCGAC/EBP-αF-CACAACAGCAACGAGTACCGR-GTCATGGTCTTATCCCCACAF-CACAACAGCAACGAGTACCGAR-GCTGATGCTTATCCCCACAF-CACAACAGCAACGAGTACCGAR-GCTGATGCTTATCCCCACAAGF-CACAACAGCAACGAGTACCGA			R-ATGAGGGAGTTGGAAGGCTCT
R- AGCTTTCTGGTGTGACTCGGHSLF- AGGAGCCAGCATTGAGACAAAR- CGCAGGTGTTGATTCAGCTTCPREF1F- CTTTCGGCCACAGCACCTATR- TGTCATCCTCGCAGAATCCATFABP4F- ACTGGGCCAGGAATTTGACGR- CTCGTGGAAGTGACGCCTTCXCL1F- TGGCTGGGATTCACCTCAAGCXCL2F- GCTGTCCCTCAACGGAAGAACXCL3F- CCTACCAAGGGTTGATTTTGAGACR- GAGTGGCTATGACTTCTGTCTGGMouseIL-6LIFF- CCCTGTAAATGCCACCTGTCXCR2F- CCCTGTAAATGCCACCTGTR- CCAAACAGCGTTGATTTCGCATCXCR2F- CCCCTGTAAATGCCACCTGTR- CCTCCTGTCCCACAAGGATCXCR2F- CCCCCGTAAATGCCACCTGTPAR-γF- CTCCAAGAATACCAAAGTGCGAR- GCTGATGCTTTATCCCACAC/EBP-αF- CAAGAACAGCAACGAGTACCGR- GTCACTGGTCAACTCCACACR- GTCACTGGTCAACTCCAGCAC		C/EBP-α	F- TATAGGCTGGGCTTCCCCTT
HSLF- AGGAGCCAGCATTGAGACAAA R- CGCAGGTGTTGATTCAGCTTCPREF1F- CTTTCGGCCACAGCACCTAT R- TGTCATCCTCGCAGAATCCATFABP4F- ACTGGGCCAGGAATTGACG R- CTCGTGGAAGTGACGCCTTCXCL1F- TGGCTGGGATTCACCTCAAG R- CCGTTACTTGGGGACACCTTCXCL2F- GCTGTCCCTCAACGGAAGAA R- CGAGGCACATCAGGTACGATMouseIL-6IL-6F- AGTCCTTCGTCGCCACACTGA R- CTTCTTGGTCCTTAGCCACLIFF- CCCCTGTAAATGCCACCTGT R- CTTCTCTGTCCGCATCXCR2F- CCACAAACAGCGTCGTAGAACT R- ACCAAGGAGTTCCCCACAAGPPAR-γF- CTCCAAGAATACCAAAGTGCGA R- GCCTGATGCTTATCCCCACAC/EBP-αF- CAAGAACAGCAACGAGTACCG R- GTCACTGGTCAACTCCAGCAC			R-AGCTTTCTGGTGTGACTCGG
R- CGCAGGTGTTGATTCAGCTTCPREF1F- CTTTCGGCCACAGCACCTATR- TGTCATCCTCGCAGAATCCATFABP4F- ACTGGGCCAGGAATTGACGR- CTCGTGGAAGTGACGCCTTCXCL1F- TGGCTGGGATTCACCTCAAGCXCL2F- GCTGTCCCTCAACGGAAGAACXCL3F- CCACCAAGGGTTGATTTGAGACCXCL3F- CCTACCAAGGGTTGATTTGAGACMouseIL-6ILFR- CCCTTGTACCTCGTCGCATCXCR2F- CCCCTGTAAATGCCACCTGTR- CCCTGTAAATGCCACCTGTR- CCCTGTAAATGCCACCTGTR- CTTCTCTGTCCGTTGCCATCXCR2F- CACAAACAGCGTCGTAGAACTR- ACCAAGGAGTTCCCCACAAGPPAR-γF- CTCCAAGAATACCAAAGTGCGAR- GCTGATGCTTATTCCCACAC/EBP-αF- CAAGAACAGCAACGAGTACCGR- GTCACTGGTCAACTCCAGCACR- GTCACTGGTCAACTCCAGCACR- GTCACTGGTCAACCCACGACACCACGAR- GTCACTGGTCAACCCACGAGTACCGR- GTCACTGGTCAACCCACGACGACCACCGAR- GTCACTGGTCAACTCCAGCAC		HSL	F- AGGAGCCAGCATTGAGACAAA
PREF1F- CTTTCGGCCACAGCACCTAT R- TGTCATCCTCGCAGAATCCATFABP4F- ACTGGGCCAGGAATTTGACG R- CTCGTGGAAGTGACGCCTTFABP4F- ACTGGGCAGGAATTTGACG R- CTCGTGGGATTCACCTCAAG R- CCGTTACTTGGGGACACCTTCXCL1F- TGGCTGGGATTCACCTCAAG R- CCGTGTCCCTCAACGGAAGAA R- CGAGGCACATCAGGTACGATMouseL-6MouseL-6LIFF- CCCTGTAAATGCCACCAC R- GGTCTTGGTCCTTAGCCACLIFF- CCCCTGTAAATGCCACCTGT R- CTTCTGTCCCGTTGCCATCXCR2F- CACAAACAGCGTCGTAGAACT R- ACCAAGGAGTTCCCCACAAGPPAR-γF- CTCCAAGAATACCAAAGTGCGA R- GCCTGATGCTTTATCCCCACAC/EBP-αF- CAAGAACAGCAACGAGTACCG R- GTCACTGGTCAACTCCAGCAC			R- CGCAGGTGTTGATTCAGCTTC
R- TGTCATCCTCGCAGAATCCATFABP4F- ACTGGGCCAGGAATTTGACG R- CTCGTGGAAGTGACGCCTTCXCL1F- TGGCTGGGATTCACCTCAAG R- CCGTTACTTGGGGACACCTTCXCL2F- GCTGTCCCTCAACGGAAGAA R- CGAGGCACATCAGGTACGATMouseIL-6IL-6F- AGTCCTTCGTCCCCAATGCCAC R- TGGTCTTGGTCCTTAGCCACLIFF- CCCCTGTAAATGCCACCTGT R- CCCCTGTAAATGCCACCTGT R- CACAAGGAGTTCCCCACAAG R- CCCCTGTAAATGCCACCTGT R- CTCCTGTCCCGCATGCCATPPAR-γF- CTCCAAGAATACCAAAGTGCGA R- GCCTGATGCTTATCCCACA R- GCCTGATGCTTATCCCACAC/EBP-αF- CAAGAACAGCAACGAGTACCG R- GTCACTGGTCAACTCCAGCAC		PREF1	F- CTTTCGGCCACAGCACCTAT
FABP4 F-ACTGGGCCAGGAATTTGACG R- CTCGTGGAAGTGACGCCTT R- CTCGTGGGATTCACCTCAAG R- CCGTTACTTGGGGACACCTT R- CCGTTACTTGGGGACACCTT CXCL2 F- GCTGTCCCTCAACGGAAGAA R- CGAGGCACATCAGGTACGAT R CXCL3 F- CCTACCAAGGGTTGATTTTGAGAC R- GAGTGGCTATGACTTCTGTCTGG R Mouse IL-6 F- AGTCCTTCCTACCCCAATTTCC R- TGGTCTTGGTCCTTAGCCAC ILIF F- CCCCTGTAAATGCCACCTGT R- CXCR2 F- CACAAACAGCGTCGTAGAACT R- ACCAAGGAGTTCCCCACAAG PPAR-γ F- CTCCAAGAATACCAAAGTGCGA R- GCCTGATGCTTATCCCACA C/EBP-α F- CAAGAACAGCAACGAGTACCG R- GTCACTGGTCAACTCCAGCAC			R- TGTCATCCTCGCAGAATCCAT
$ \begin{array}{llllllllllllllllllllllllllllllllllll$		FABP4	F-ACTGGGCCAGGAATTTGACG
$\begin{array}{cccc} CXCL1 & F-TGGCTGGGATTCACCTCAAG \\ R-CCGTTACTTGGGGACACCTT \\ CXCL2 & F-GCTGTCCCTCAACGGAAGAA \\ R-CGAGGCACATCAGGTACGAT \\ CXCL3 & F-CCTACCAAGGGTTGATTTTGAGAC \\ R-GAGTGGCTATGACTTCTGTCTGG \\ Mouse & IL-6 & F-AGTCCTTCCTACCCCAATTTCC \\ R-TGGTCTTGGTCCTTAGCCAC \\ LIF & F-CCCCTGTAAATGCCACCAC \\ LIF & F-CCCCTGTAAATGCCACCAT \\ CXCR2 & F-CACAACAGCGTCGTAGAACT \\ R-ACCAAGGAGTTCCCCACAAG \\ PPAR-\gamma & F-CTCCAAGAATACCAAAGTGCGA \\ R-GCCTGATGCTTTATCCCCAC \\ C/EBP-\alpha & F-CAAGAACAGCAACGAGTACCG \\ R-GTCACTGGTCAACTCCAGCAC \\ \end{array}$			R- CTCGTGGAAGTGACGCCTT
$\begin{array}{llllllllllllllllllllllllllllllllllll$		CXCL1	F- TGGCTGGGATTCACCTCAAG
$\begin{array}{llllllllllllllllllllllllllllllllllll$			R- CCGTTACTTGGGGACACCTT
$\begin{array}{llllllllllllllllllllllllllllllllllll$		CXCL2	F- GCTGTCCCTCAACGGAAGAA
$\begin{array}{llllllllllllllllllllllllllllllllllll$			R- CGAGGCACATCAGGTACGAT
$ \begin{array}{llllllllllllllllllllllllllllllllllll$		CXCL3	F- CCTACCAAGGGTTGATTTTGAGAC
MouseIL-6F- AGTCCTTCCTACCCCAATTTCC R- TGGTCTTGGTCCTTAGCCACLIFF- CCCCTGTAAATGCCACCTGT R- CTTCTCTGTCCCGTTGCCATCXCR2F- CACAAACAGCGTCGTAGAACT R- ACCAAGGAGTTCCCCACAAGPPAR- γ F- CTCCAAGAATACCAAAGTGCGA R- GCCTGATGCTTTATCCCCACAC/EBP- α F- CAAGAACAGCAACGAGTACCG R- GTCACTGGTCAACTCCAGCAC			R- GAGTGGCTATGACTTCTGTCTGG
R- TGGTCTTGGTCCTTAGCCACLIFF- CCCCTGTAAATGCCACCTGTR- CTTCTCTGTCCCGTTGCCATCXCR2F- CACAAACAGCGTCGTAGAACTR- ACCAAGGAGTTCCCCACAAGPPAR-γF- CTCCAAGAATACCAAAGTGCGAR- GCCTGATGCTTTATCCCCACAC/EBP-αF- CAAGAACAGCAACGAGTACCGR- GTCACTGGTCAACTCCAGCAC	Mouse	IL-6	F- AGTCCTTCCTACCCCAATTTCC
LIF F- CCCCTGTAAATGCCACCTGT R- CTTCTCTGTCCCGTTGCCAT CXCR2 F- CACAAACAGCGTCGTAGAACT R- ACCAAGGAGTTCCCCACAAG PPAR-γ F- CTCCAAGAATACCAAAGTGCGA R- GCCTGATGCTTTATCCCCACA C/EBP-α F- CAAGAACAGCAACGAGTACCG R- GTCACTGGTCAACTCCAGCAC			R-TGGTCTTGGTCCTTAGCCAC
R- CTTCTCTGTCCCGTTGCCATCXCR2F- CACAAACAGCGTCGTAGAACTR- ACCAAGGAGTTCCCCACAAGPPAR-γF- CTCCAAGAATACCAAAGTGCGAR- GCCTGATGCTTTATCCCCACAC/EBP-αF- CAAGAACAGCAACGAGTACCGR- GTCACTGGTCAACTCCAGCAC		LIF	F- CCCCTGTAAATGCCACCTGT
CXCR2 F- CACAAACAGCGTCGTAGAACT R- ACCAAGGAGTTCCCCACAAG PPAR-γ F- CTCCAAGAATACCAAAGTGCGA R- GCCTGATGCTTTATCCCCACA C/EBP-α F- CAAGAACAGCAACGAGTACCG R- GTCACTGGTCAACTCCAGCAC			R- CTTCTCTGTCCCGTTGCCAT
R- ACCAAGGAGTTCCCCACAAGPPAR-γF- CTCCAAGAATACCAAAGTGCGAR- GCCTGATGCTTTATCCCCACAR- GCCTGATGCTTTATCCCCACAC/EBP-αF- CAAGAACAGCAACGAGTACCGR- GTCACTGGTCAACTCCAGCAC		CXCR2	F- CACAAACAGCGTCGTAGAACT
PPAR-γ F- CTCCAAGAATACCAAAGTGCGA R- GCCTGATGCTTTATCCCCACA C/EBP-α F- CAAGAACAGCAACGAGTACCG R- GTCACTGGTCAACTCCAGCAC			R-ACCAAGGAGTTCCCCACAAG
R- GCCTGATGCTTTATCCCCACAC/EBP-αF- CAAGAACAGCAACGAGTACCGR- GTCACTGGTCAACTCCAGCAC		PPAR-γ	F- CTCCAAGAATACCAAAGTGCGA
C/EBP-α F- CAAGAACAGCAACGAGTACCG R- GTCACTGGTCAACTCCAGCAC		•	R- GCCTGATGCTTTATCCCCACA
R- GTCACTGGTCAACTCCAGCAC		C/EBP-α	F- CAAGAACAGCAACGAGTACCG
			R- GTCACTGGTCAACTCCAGCAC

Number	Age	Height(m)	Weight(kg)	BMI	Histology	Stage
1	69	148	50	22.83	IDC	3A
2	54	159	53	20.96	IDC	2B
3	57	154	55	23.19	IDC	2B
4	61	155	56	23.31	IDC	3C
5	56	156	60	24.65	IDC	2B
6	33	165	47	17.26	IDC	2A
7	52	158	53	21.23	IDC	2B
8	58	156	52	21.37	IDC	3A
9	51	160	65	25.39	IDC	3A
10	77	145	52	24.73	IDC	3A

Supplementary Table 3 Information on 10 patients with breast cancer.

Note: IDC, invasion ductal carcinoma



Supplementary figure

Supplement figure 1. Formation of CAA. (A) The CD44 and CD90 expressions of the third generation pre-adipocytes were analyzed by flow cytometry. (B) The mRNA expression of differentiation markers in pre-adipocytes and mature adipocytes was detected by q-PCR. (C and D) The mRNA expression of differentiation markers and inflammatory factors in adipocytes and pre-adipocytes was detected by q-PCR. (E) Oil red O staining showed the morphology of lipid droplets in adipocytes cultured alone or co-cultured adipocytes (CAA), scale bar: 200 µm. (F and

G) MDA-MB-231 cells were stimulated by DMEM (negative control group), Adi-CM, CAA-CM and DMEM containing 10% FBS (positive control group), and the cell migration (**F**) and invasion (**G**) were observed by phase contrast microscope. Typical microscopic fields are shown and quantitative data are presented as mean \pm SD from at least three independent experiments. *p < 0.05, **p < 0.01, ***p < 0.001.



Supplement figure 2. CAA-derived LIF promotes breast cancer cell migration and invasion. (A and B) The mRNA expression of LIF in co-cultured pre-adipocytes (A) and mature adipocytes (B) at different time was detected by q-PCR. (C) The protein expression of LIF in co-cultured pre-adipocytes and mature adipocytes was analyzed by western blot. (D) The protein secretion of LIF in co-cultured pre-adipocytes, mature adipocytes and MDA-MB-231 cells was analyzed by ELISA. (E and F) BT549 cells were treated with rhLIF and subjected to migration (E) and Transwell Matrigel invasion assays (F). (G) MDA-MD-231 and BT549 cells were treated with rhLIF (0, 20, 40, 80, 160 ng/ml) for 48 h. DMEM containing 0.2% FBS was used as a control, and the effect of rhLIF on breast cancer cell proliferation was analyzed by MTT assay. (H) MDA-MD-231 cells were treated with rhLIF (0, 20, 40, 80, 15, 15, 30, 60 min). Stat3 phosphorylation was analyzed by western

blot. (**I and J**) BT549 cells cultured in Adi-CM or CAA-CM or DMEM were treated with LIFneutralizing antibody or IgG, and the cell migration (**I**) and invasion (**J**) were observed by phase contrast microscope. Typical microscopic fields and blots are shown, and quantitative data are presented as the mean \pm SD from at least three independent experiments. (**K**) BT549 cells were treated with DMEM or culture media of co-cultured pre-adipocytes or culture media of CAA for 15 min, and Stat3 phosphorylation was analyzed by western blot. (**L**) BT549 cells were treated with different culture media and LIF-neutralizing antibody or IgG for 15 min, and Stat3 phosphorylation was analyzed by western blot. *p < 0.05, **p < 0.01, ***p < 0.001.



Supplement figure 3. Stattic inhibits the migration and invasion and Stat3 phosphorylation of BT549 cells induced by rhLIF and CAA-CM. (A and B) BT549 cells were stimulated with rhLIF or DMEM and Stattic and subjected to migration (A) and Transwell Matrigel invasion (B) assays. (C) BT549 cells were stimulated by rhLIF combined with Stattic for 15 min, the protein

expression level was analyzed by western blot. (**D** and **E**) BT549 cells were stimulated by DMEM, Adi-CM or CAA-CM combined with Stattic and subjected to migration (**D**) and Transwell Matrigel invasion (**E**) assays. (**F**) BT549 cells were treated with DMEM or different adipocyte culture medium with Stattic for 15 min, and the protein expression levels were analyzed by western blot. (**G**) Western blot was performed on MDA-MB-231 cells transfected with non-specific (NS) or Stat3 siRNA and wild-type MDA-MB-231 cells (Con). (**H**) The p-ERK1/2 expression level in pre-adipocytes was analyzed by western blot. Typical microscopic fields and blots are shown and quantitative data are presented as mean ± SD from at least three independent experiments. ${}^{\#}p < 0.05$, ${}^{\#\#}p < 0.01$, ${}^{\#\#}p < 0.001$, rhLIF or CAA-CM VS. Con; ${}^{*}p < 0.05$, ${}^{**}p <$ 0.01, ${}^{***}p < 0.001$, rhLIF + Stattic VS. rhLIF or CAA-CM + Stattic VS. CAA-CM.



Supplement figure 4. The validation of a co-culture system for 3T3-L1 and 4T1 cells. (A) The mRNA expression of differentiation marker genes, LIF, CXCR2 and IL-6 in control and co-cultured 3T3-L1 was detected by q-PCR. (B) The protein expression levels of signaling pathway in control and co-cultured 3T3-L1 were analyzed by western blot. (C) The mRNA expression of CXCL1-3 and IL-6 in control and co-cultured 3T3-L1 was detected by q-PCR. (D) The protein expression level of p-Stat3 in control and co-cultured 4T1 cells was analyzed by western blot. The quantitative data are presented as mean \pm SD from at least three independent experiments. *p < 0.05, **p < 0.01, ***p < 0.001.