

Supplementary Table S1: Oligo sequences of plasmid cloning and qPCR.

Luciferase reporter plasmid construct sequences	
YAP1 Wild Type Sense(52bp)	CAATTCAGAATTCATACCAATCAGTGTTGAAACTCAA CATTGCAAAAAGT
YAP1 Wild Type Antisense(58bp)	CTAGACTTTTGCAATGTTTGAGTTTCAACTGATTGG TATGAATTCTGAATTGAGCT
YAP1 Mutant Type Sense(52bp)	CAATTCAGAATTCATACCAATCCCTCTTGAAACTCAA CATTGCAAAAAGT
YAP1 Mutant Type Antisense(58bp)	CTAGACTTTTGCAATGTTTGAGTTTCAAGAGGGATTG GTATGAATTCTGAATTGAGCT
ChIP assay primers	
CBE1 Forward	GGGACACTGTATTTGGTCTCC
CBE1 Reverse	CCTTTCGGTACTCAGGGT
CBE2 Forward	GAGCTCTGAATCCCACTTGG
CBE2 Reverse	GCCGGATCCACATAGAGTC
CBE3 Forward	GGGGCCCAATACTGAATACC
CBE3 Reverse	GCATAGCACATGACACCAGG
Luciferase reporter plasmid constructed primers	
GROalpha CBE1-3 Forward	ACATGCTAGCACACCTTTCTAC
GROalpha CBE2-3 Forward	TAAAGCTAGCAGGTAAATGAC
GROalpha CBE3 Forward	TTTTCGCTAGCAAGGGGGCCCAA
GROalpha CBEN Forward	ACTGACGCTAGCACTGGATTGAGC
GROalpha CBE Reverse	AGTGGAAGCTTGCGCTGAAGATACCA
CRISPR/Cas9 Knock-out sgRNA	
YAP1 Sense	CACCGGACTCGGAGACCGACCTGG
YAP1Antisense	AAACCCAGGTCGGTCTCCGAGTCCC
TAZ Sense	CACCGGAGATGGCGTCTACCAGAAG
TAZ Antisense	AAACCTTCTGGTAGACGCCATCTCC
CRISPRi sgRNA	
Cxcr1 Sense1	CACCGACCCAGCTGGTGCCTCAGGT
Cxcr1 Antisense1	AAACACCTGAGGCACCAGCTGGGTC
Cxcr1 Sense2	CACCGGTACGCAACTCTGACTTCCT
Cxcr1 Antisense2	AAACAGGAAGTCAGAGTTGCGTACC
Cxcr1 Sense3	CACCGCTGGGTCCGTTCTTGTGTG
Cxcr1 Antisense3	AAACCACACAAGGAACGGACCCAGC
Cxcr2 Sense1	CACCGGTCGTAGAACTACTGCAGGT
Cxcr2 Antisense1	AAACACCTGCAGTAGTTCTACGACC

Cxcr2 Sense2	CACCGATGCGTCATGCCGCTTCCCA
Cxcr2 Antisense2	AAACTGGGAAGCGGCATGACGCATC
Cxcr2 Sense3	CACCGGACGCTGTTTGTGAGCAGAG
Cxcr2 Antisense3	AAACCTCTGCTCACAAACAGCGTCC

TEAD1_TFBS_binding sites						
TEAD1_HUMAN.H10 MO.D	CXCL1(-2k~+2k)	68	81	+	Forward: GGGACTGTATTTGGT CTCC	
\					Reverse: CCTTCGGTTACTCAGG GGT	from -1933 to -1920 AACATTCTAGC ACA
TEAD1_HUMAN.H10 MO.D	CXCL1(-2k~+2k)	861	874	+	Forward: GAGCTCTGAATCCCACT TGG	
					Reverse: GCCGGATCCCACATAG AGTC	from -1140 to -1127 GAAATTCCTGA GAG
TEAD1_HUMAN.H10 MO.D	CXCL1(-2k~+2k)	989	1002	+	Forward: GGGGCCAATACTGAA TACC	
					Reverse: GCATAGCACATGACACC AGG	from -1012 to -999 TGCATGTCTGCT GC