

## SUPPLEMENTARY TABLES

### Table S1 – Materials and Methods

#### Plasmids

pLenti6-miRNA-Ctrl  
pLenti6-miRNA-KMT9 $\alpha$

#### miRNA sequences (5'-3')

##### Human miKMT9 $\alpha$

###### *Top strand*

TGCTGATGTGAACTTTGTTACAGCGTGT TTTGGCCACTGACTGACACGCTGTAA  
AAGTTCACAT

###### *Bottom strand*

CCTGATGTGAACTTTTACAGCGTGTCAGTCAGTGGCCAAAACACGCTGTAACAA  
AGTTCACATC

##### Human miCtrl

###### *Top strand*

TGCTGAAATGTACTGCGCGTGGAGACGTTTTGGCCACTGACTGACGTCTCCAC  
GCAGTACATTT

###### *Bottom strand*

CCTGAAATGTACTGCGTGGAGACGTCAGTCAGTGGCCAAAACGTCTCCACGCG  
CAGTACATTTTC

Control miRNA (miCtrl) or miRNA directed against *KMT9 $\alpha$*  (miKMT9 $\alpha$ ) were designed using BLOCK-IT™ RNAi Designer. Each oligonucleotides pair encodes for a pre-miRNA and contains a linker sequence (purple), the mature miRNA sequence (green), a loop sequence (black) and a sequence containing the nucleotides 1-8 and 11-21 of the 21-mer target sequence (red).

#### Primer sequences qRT-PCR (5'-3')

<i>Aurkb</i>	CAGACTTTGGCTGGTCCGGT and TATGCATGCGCCCCTCAATC
<i>E2f1</i>	CAGCTGCAACTGCTTTCCGG and AGGAGGGGCCTTGATCACTA
<i>Esco2</i>	CACCACCGATTTCTGGAAGGA and TGGCAGGACCAACACAATCT
<i>Hprt</i>	GTTAAGCAGTACAGCCCCAAA and AGGGCATATCCAACAACAAACTT
<i>HPRT</i>	CCTGGCGTCGTGATTAGTGAT and AGACGTTTCAGTCCTGTCCATAA
<i>Kmt9<math>\alpha</math></i>	CAGCCGCATGTACCTTGGAA and TCCACGACTTCTACCTCTTCA
<i>KMT9<math>\alpha</math></i>	ACGTTTCTGCTTTTGGACGC and TCAGTGCACATGTACAAAGCC
<i>Mcm6</i>	ACCTTCTCTTTGCCCGACAG and CTGGCGGAGACGTTTGTACT
<i>Pclaf</i>	CTGGCGGAGACGTTTGTACT and TGGGTTCCCTCCTGCATACTT
<i>Pmaip1</i>	GAAGTCGCAAAGAGCAGGA and GTTGAGCACACTCGTCCTTCA
<i>Prr11</i>	TTGTGCAAGCTCCGAGAAAAC and ATTCGTCGTTGGAGCACAGG
<i>Rad51c</i>	GGTTTCAGACGGCGGAGGA and CGGCACATCTTGGTTTATTTGT
<i>Rpa2</i>	ACCAGGATGTGGAATAGCGG and ACAATATGCTGGGCTCGGAC
<i>Tgfb2</i>	CAGCGCTACATCGATAGCAA and CCTCGAGCTCTTCGCTTTTA
<i>Top2a</i>	CCTCGGGGCAAAGAGTCAT and CTATTCGTTGCCGGAG

**Table S2 – List of the 12 significantly differentially expressed genes in KMT9 $\alpha$ -deficient compared to KMT9 $\alpha$ -proficient colon ( $p < 1e-6$ )**

Gene name	Fold change
<i>Ceacam2</i>	12.04
<i>Dmtn</i>	20.86
<i>Gm15915</i>	7.05
<i>H2-Aa</i>	0.22
<i>H2-Ab1</i>	0.26
<i>Krt12</i>	7.65
<i>Lpo</i>	36.74
<i>N6amt1</i>	0.51
<i>Phyhip</i>	19.84
<i>Scpep1</i>	1.62
<i>Serinc3</i>	1.49
<i>Slpi</i>	0.13

**Table S3 – Gene list of pro-apoptotic genes used to calculate gene scores with Seurat depicted in figure 3I.**

<b>Gene symbol</b>	<b>Full gene name</b>
<b><i>Aifm3</i></b>	apoptosis inducing factor mitochondria associated 3
<b><i>Anxa1</i></b>	annexin a1
<b><i>App</i></b>	amyloid beta precursor protein
<b><i>Atf3</i></b>	activating transcription factor 3
<b><i>Bax</i></b>	bcl2 associated x, apoptosis regulator
<b><i>Bcap31</i></b>	b cell receptor associated protein 31
<b><i>Bcl10</i></b>	bcl10 immune signaling adaptor
<b><i>Bcl2l11</i></b>	bcl2 like 11
<b><i>Bid</i></b>	bh3 interacting domain death agonist
<b><i>Bik</i></b>	bcl2 interacting killer
<b><i>Bmf</i></b>	bcl2 modifying factor
<b><i>Bmp2</i></b>	bone morphogenetic protein 2
<b><i>Bnip3l</i></b>	bcl2 interacting protein 3 like
<b><i>Brca1</i></b>	brca1 dna repair associated
<b><i>Btg2</i></b>	btg anti-proliferation factor 2
<b><i>Btg3</i></b>	btg anti-proliferation factor 3
<b><i>Casp1</i></b>	caspase 1
<b><i>Casp2</i></b>	caspase 2
<b><i>Casp3</i></b>	caspase 3
<b><i>Casp4</i></b>	caspase 4
<b><i>Casp6</i></b>	caspase 6
<b><i>Casp7</i></b>	caspase 7
<b><i>Casp8</i></b>	caspase 8
<b><i>Casp9</i></b>	caspase 9
<b><i>Cdkn1a</i></b>	cyclin dependent kinase inhibitor 1a
<b><i>Cdkn1b</i></b>	cyclin dependent kinase inhibitor 1b
<b><i>Cyld</i></b>	cyld lysine 63 deubiquitinase
<b><i>Dap</i></b>	death associated protein
<b><i>Dap3</i></b>	death associated protein 3
<b><i>Ddit3</i></b>	dna damage inducible transcript 3
<b><i>Dffa</i></b>	dna fragmentation factor subunit alpha
<b><i>Diablo</i></b>	diablo iap-binding mitochondrial protein
<b><i>Dnaja1</i></b>	dnaj heat shock protein family (hsp40) member a1
<b><i>Dnajc3</i></b>	dnaj heat shock protein family (hsp40) member c3
<b><i>Dnm1l</i></b>	dynamamin 1 like
<b><i>Egr3</i></b>	early growth response 3
<b><i>Fas</i></b>	fas cell surface death receptor
<b><i>Fdxr</i></b>	ferredoxin reductase
<b><i>Gadd45a</i></b>	growth arrest and dna damage inducible alpha
<b><i>Gadd45b</i></b>	growth arrest and dna damage inducible beta
<b><i>Hmgb2</i></b>	high mobility group box 2
<b><i>Ifngr1</i></b>	interferon gamma receptor 1
<b><i>Il18</i></b>	interleukin 18
<b><i>Irf1</i></b>	interferon regulatory factor 1
<b><i>Madd</i></b>	map kinase activating death domain
<b><i>Nedd9</i></b>	neural precursor cell expressed, developmentally down-regulated 9
<b><i>Nefh</i></b>	neurofilament heavy
<b><i>Pdcd4</i></b>	programmed cell death 4
<b><i>Pmaip1</i></b>	phorbol-12-myristate-13-acetate-induced protein 1

<b><i>Ppp2r5b</i></b>	protein phosphatase 2 regulatory subunit b'beta
<b><i>Psen1</i></b>	presenilin 1
<b><i>Psen2</i></b>	presenilin 2
<b><i>Rara</i></b>	retinoic acid receptor alpha
<b><i>Rela</i></b>	rela proto-oncogene, nf-kb subunit
<b><i>Rhob</i></b>	ras homolog family member b
<b><i>Rock1</i></b>	rho associated coiled-coil containing protein kinase 1
<b><i>Sat1</i></b>	spermidine/spermine n1-acetyltransferase 1
<b><i>Slc20a1</i></b>	solute carrier family 20 member 1
<b><i>Smad7</i></b>	smad family member 7
<b><i>Tap1</i></b>	transporter 1, atp binding cassette subfamily b member
<b><i>Tgfb2</i></b>	transforming growth factor beta 2
<b><i>Tgfb3</i></b>	transforming growth factor beta receptor 3
<b><i>Tnf</i></b>	tumour necrosis factor
<b><i>Tnfrsf12a</i></b>	tnf receptor superfamily member 12a
<b><i>Tnfsf10</i></b>	tnf superfamily member 10
<b><i>Tspo</i></b>	translocator protein
<b><i>Txnip</i></b>	thioredoxin interacting protein
<b><i>Vdac2</i></b>	voltage dependent anion channel 2