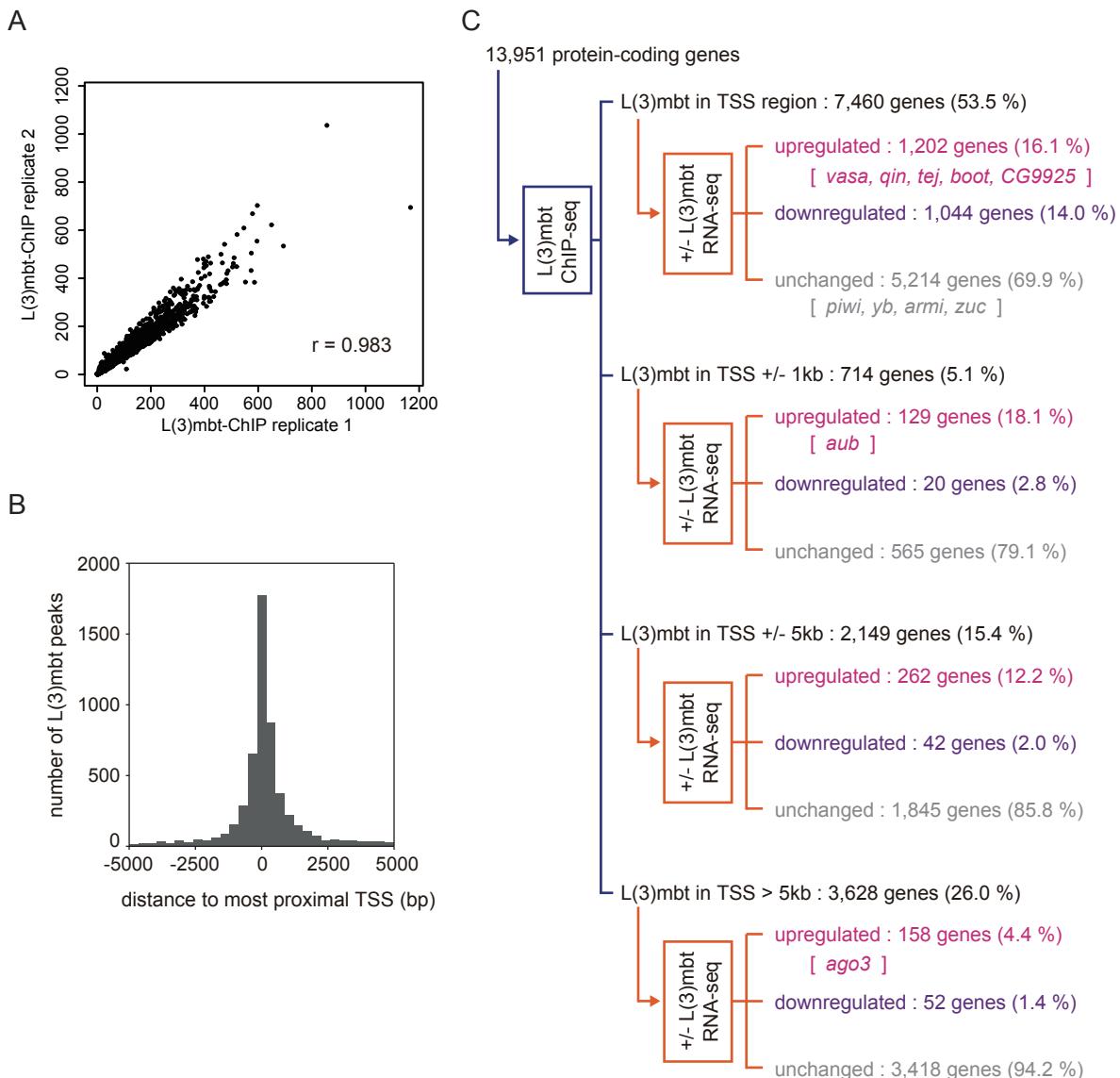


Appendix PDF

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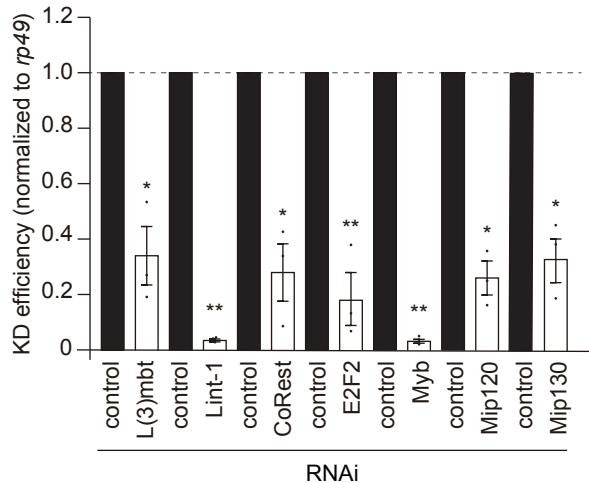
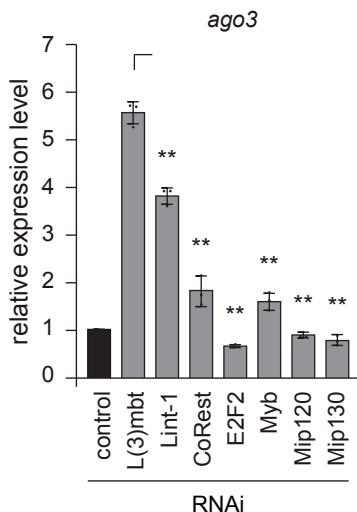
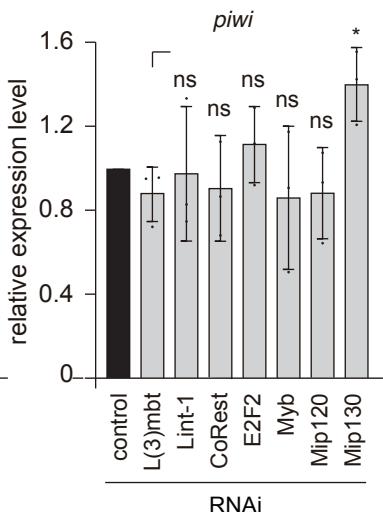
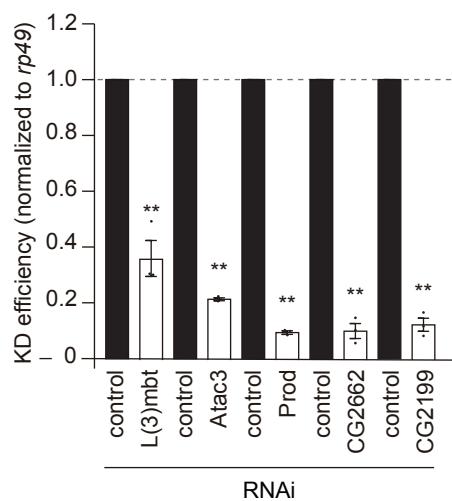


Appendix Figure S1. Relationship between L(3)mbt ChIP peaks and TSS.

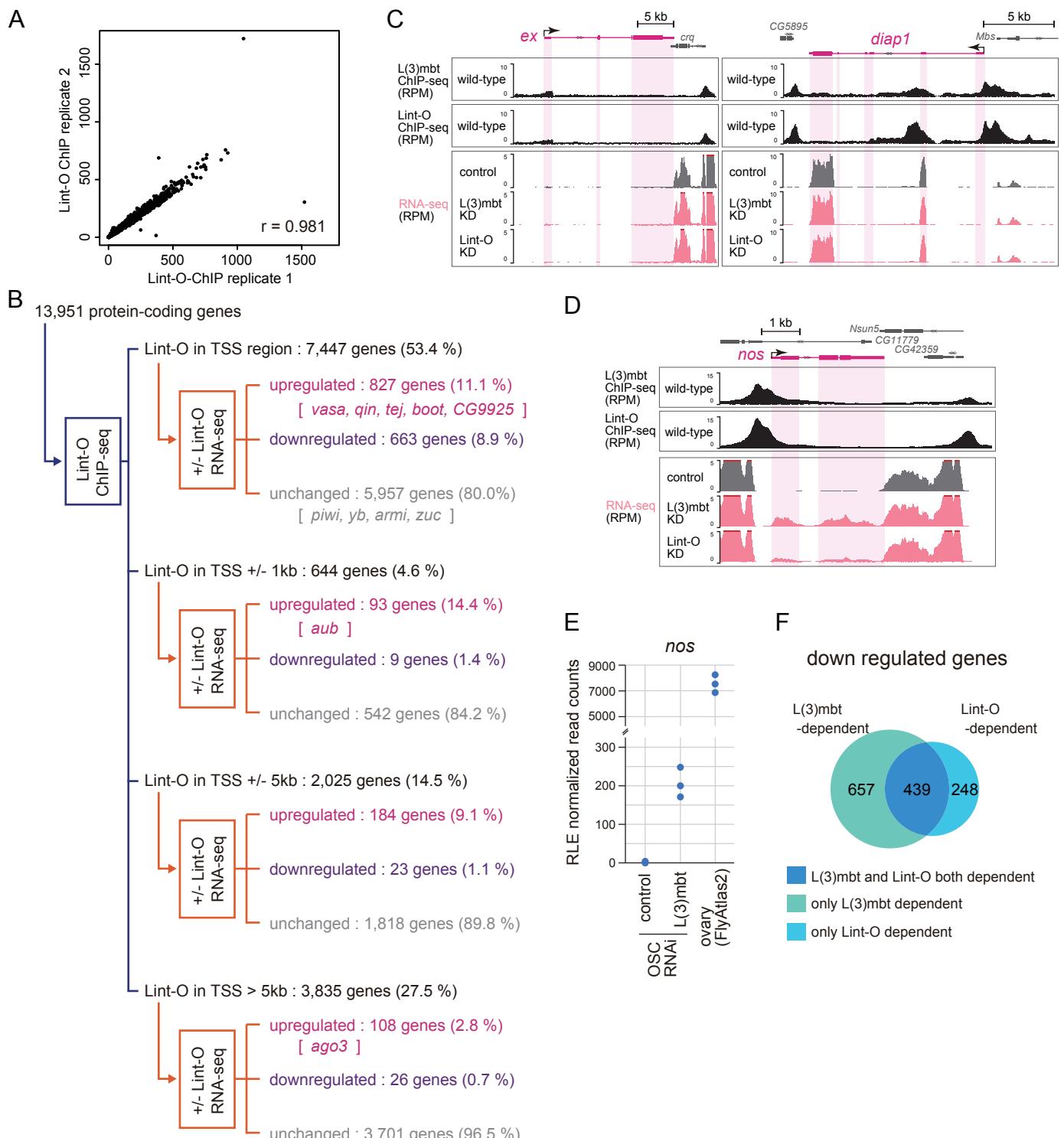
(A) Scatter plot of sequenced reads, mapped onto each L(3)mbt ChIP peak defined by MACS2 for each replicate. Each dot denotes a detected ChIP peak. (B) Distribution of L(3)mbt ChIP peaks is shown according to the distance to most proximal TSS. (C) The 13,951 protein-coding genes of *Drosophila* were classified into “L(3)mbt in TSS region,” “L(3)mbt in TSS +/- 1kb,” “L(3)mbt in TSS +/- 5kb,” and “L(3)mbt in TSS > 5kb,” in accordance with the L(3)mbt ChIP-seq signals, and were subsequently divided into “upregulated,” “downregulated,” and “unchanged” in accordance with the RNA-seq reads from OSCs before and after L(3)mbt depletion [+/- L(3)mbt]. Representatives of piRNA factors are indicated within the groups. Note that the “L(3)mbt in TSS region” corresponds to the “promoter region binding” in Fig 1A.

A

L(3)mbt-containing complex	components
LINT	L(3)mbt, Lint-1, CoRest
dREAM/MMB	L(3)mbt, Myb, E2F2, RBF, p53, Mip120, Mip130, Mip140

B**C****D****E****Appendix Figure S2. The mRNA levels of *ago3* and *piwi* upon the depletion of L(3)mbt interactors in OSCs.**

(A) Known L(3)mbt cofactors within the LINT and dREAM complexes (Lewis et al. 2004; Meier et al. 2012; Blanchard et al, 2014). (B) Knockdown (KD) efficiencies of members of the LINT and dREAM complexes were verified by RT-qPCR. Control: EGFP siRNA was used. Data represent the mean \pm SE ($n = 3$). The p values were calculated with the t-test. * p : < 0.05 , ** p : < 0.01 . (C, D) The mRNA levels of *ago3* (C) and *piwi* (D) were quantified in L(3)mbt-, Lint-1-, CoRest-, E2F2-, Myb-, Mip120-, and Mip130-depleted (RNAi) OSCs, and were compared with those in normal (control) OSCs. (E) KD efficiencies were verified by RT-qPCR for the indicated genes.



Appendix Figure S3. Genes bound and controlled by Lint-O.

(A) Scatter plot of sequenced reads, mapped onto each Lint-O ChIP peak defined by MACS2 for each replicate. Each dot denotes a detected ChIP peak. (B) The 13,951 protein-coding genes of Drosophila were classified into “Lint-O in TSS region,” “Lint-O in TSS +/- 1kb,” “Lint-O in TSS +/- 5kb,” and “Lint-O in TSS > 5kb,” in accordance with the Lint-O ChIP-seq reads, and were subsequently divided into “upregulated,” “downregulated,” and “unchanged” in accordance with the RNA-seq reads from OSCs before and after Lint-O depletion [+/- Lint-O]. Representatives of piRNA factors are indicated within the groups. Note that the “Lint-O in TSS region” corresponds to the “promoter region binding” in Fig 4A. (C, D) The genomic regions harboring the *diap1* (C), *ex* (C), and *nos* (D) genes. The L(3)mbt and Lint-O ChIP-seq reads in normal OSCs (wild-type) and RNA-seq reads from L(3)mbt- and Lint-O-depleted [L(3)mbt KD and Lint-O KD] and control OSCs are shown. The shades in gray correspond to exons. The y-axis shows the number of RPM. (E) The levels of *nos* mRNA in the ovary and cultured OSCs following L(3)mbt depletion are shown. (F) Venn diagram shows that 439 protein-coding genes are commonly present in the L(3)mbt and Lint-O libraries as downregulated genes.

Appendix Table S1. List of proteins in LC/MS/MS analysis.

Accession	Description	Sum PEP Score	PSMs	excluded in first selection, excluded in second selection, excluded in third selection, and final selected proteins			
				Abundances (control)		Abundances (L3)mbt KD)	
E8NH29	LD46071p (Fra)	92.506	778	128.5	50.4	91.6	129.5 0.809136137
Q9VZ22	[L3]mbt_interact	37.045	54	326	70.2	1.9	1.5 104.2631579
Q9VB52	LD05287p OS-	23.163	17	381.4	18.6	0	0 ∞
E4NKN3	GM13963p OS-	17.992	61	247.3	68.8	32.5	51.5 3.763095238
L0MQ04	ATP synthase	16.843	25	39.2	281.2	38.5	41.1 4.025125628
O76866	EG100G10.6 p	14.463	14	383.8	14.2	0	1.9 209.4736842
Q9VLK1	U3 small nucle	14.345	9	274.4	0	27.3	98.3 2.184713376
P56538	Eukaryotic tra	12.677	17	38.8	163.9	188.8	8.6 1.026849037
P29844	Endoplasmic r	12.448	13	168.8	90.3	85.5	55.4 1.838892832
Q9VJD1	Glucosidase 2	11.363	13	131.1	176.4	35.6	56.9 3.324324324
A0A0B4KHY6	Ets at 97D, iso	10.881	7	164.5	94.6	107.7	33.2 1.838892832
Q0ZH21	L23A ribosoma	9.293	5	242.3	54.7	80.4	22.7 2.880698351
Q9VUZO	Transloccon-as	8.169	15	38.8	262.7	67.1	31.4 3.060913706
P17704	40S ribosomal	7.807	14	253	52.9	41.8	52.3 3.250797024
D2NUK9	Hoi-polloi_isof	7.859	18	174.1	161.8	21.2	42.9 5.24024961
A0A0B4LH50	Actin 87E, iso	7.612	18	3.1	193.1	121.8	82 0.962708538
P55830	40S ribosomal	7.37	8	93.7	223.6	28.3	54.4 3.836759371
M9NE89	Lamin, isoform	7.079	7	3.9	214.1	68	114 1.197802198
P50887	60S ribosomal	6.947	51	146.9	95.5	127.6	30 1.538071066
C0HKA1	40S ribosomal	6.738	7	130	136.5	103.4	30.1 1.996254682
Q06559	40S ribosomal	6.598	7	182.4	100.9	56.7	60 2.427592117
Q9VPQ2	DnaJ homolog	5.88	14	90.7	173	94.3	42 1.934702861
X2JF59	Thioredoxin pe	4.989	8	0	261.5	62	76.5 1.888086643
Q7JZ53	CG4866 OS-D	4.883	1	218.1	0	181.9	0 1.199010445
Q9VG9	Probable ribos	4.666	2	205.6	0	194.4	0 1.057613169
P08736	Elongation fac	4.588	12	39.6	193.4	102.3	64.8 1.394374626
A0A023GQA5	Lethal (2) 37C	4.276	2	0	357.9	0	42.1 8.501187648
Q9VTP4	60S ribosomal	3.876	2	302	270.2	85.3	14.3 3.016064257
A0A0B4K6N1	40S ribosomal	3.528	1	400	0	0	0 ∞
P11147	Heat shock 70	3.513	6	61.3	218.8	0	119.9 2.336113428
P46222	60S ribosomal	3.443	2	400	0	0	0 ∞
H0RNL4	FI17108p1 (Fra)	3.391	2	327.7	0	72.3	0 4.532503458
B5RIJ9	FI02875p (Fra)	2.925	1	0	338.7	0	61.3 5.525285481
Q8STI6	RE53401p OS-	2.831	1	400	0	0	0 ∞
P13469	DNA-binding p	2.563	1	0	0	0	0 ∞
Q8T479	AT12489p OS-	2.553	1	0	331.4	0	68.6 4.83090379
A0A0F7J3E9	MIP06027p1 O	2.524	2	225.9	53.4	90.1	30.5 2.315920398
Q9V597	60S ribosomal	2.51	1	195.5	0	204.5	0 0.95599022
P41374	Eukaryotic tra	2.424	1	63.9	259.1	0	77 4.194805195
A0A0B4KH25	Histone H2A C	2.41	6	193.5	85.5	79.8	41.2 2.305785124
Q7KLW6	LD06837p (Fra)	2.318	1	0	400	0	0 ∞
Q4V5X9	IP07888p OS-	2.316	3	287.3	0	112.7	0 2.549245785
A0A23T5E7	GEO01108p1 C	2.224	2	123.8	114.4	86.7	75.1 1.472187886
P02283	Histone H2B C	2.166	10	238	41.1	75	45.9 2.308519438
Q9T183	GH07456p OS-	2.102	3	33.6	305.3	19.7	41.3 5.555737705
A0A0B4LG52	GEO008239p1 C	2.05	2	108.9	139.2	0	152 1.632236842
Q7JNE1	Non-histone c	1.992	2	0	346.7	0	53.3 6.504690432
Q08473	RNA-binding p	1.98	3	68.2	197.7	46.7	87.4 1.98284862
Q9VWD9	LD38919p OS-	1.957	2	213	187	0	0 ∞
C6SV50	40S ribosomal	1.919	2	134.6	216.8	0	48.6 7.230452675
A8DY43	Uncharacterize	1.858	2	0	0	0	0 ∞
Q9VHG5	Insulator bindi	1.854	11	87.5	86.9	212.1	13.6 0.772707133
A9UN96	LD25519p OS-	1.853	1	400	0	0	0 ∞
P54397	39 kDa FK506	1.795	1	400	0	0	0 ∞
P06603	Tubulin alpha-	1.781	1	56.1	0	343.9	0 0.163128817
Q7K5M2	GH05836p OS-	1.748	2	88.3	177.5	80.1	54.1 1.980625931
P09180	60S ribosomal	1.604	1	186.5	0	213.5	0 0.87353563
B7ZWP7	RH14172p (Fra)	1.601	1	91	159.3	81.7	68 1.672010688
M9PFN1	RuvB-like heli	1.576	2	12.3	211.3	94.6	81.9 1.266855524
Q7KRI2	Longitudinali	1.34	1	16.7	31.9	69.4	0 4.763688761
Q9VHX4	LD24679p OS-	1.314	3	27.4	127.3	134.1	111.2 0.630656339
Q8IR16	No on or off tr	1.28	2	305.8	0	94.2	0 3.246284501
COHL66	Histone H3.3A	1.259	2	400	0	0	0 ∞
Q7K4B2	CG7845, isoform	1.259	1	0	256.3	0	143.7 1.783576896
Q9VSW7	LD21862p OS-	1.228	2	0	355.3	0	44.7 7.948545861
Q9ITW5	Something abo	1.154	1	400	0	0	0 ∞
Q9VQ29	Cytochrome b-	1.145	1	0	371.1	0	28.9 12.84083045
Q9VTA4	Integrin alpha-	1.136	4	400	0	0	0 ∞
A8DYH1	Muscle wasted	1.073	14	22.8	332.2	11.6	33.4 7.888888889
Q9V3P3	LD45860p OS-	1.067	2	0	341.5	0	58.5 5.8378068638
Q9VA91	40S ribosomal	1.049	1	400	0	0	0 ∞
A0A0B4LF09	GEO04460p1 C	1.044	1	364.3	0	35.7	0 1.020448179
A0A0B4KFV4	Uncharacterize	1.037	2	12.3	188.2	118.8	80.7 1.005012531
Q9VMT4	GH15768p OS-	0.998	5	0	0	400	0 0
M9PDW8	Uncharacterize	0.99	11	400	0	0	0 ∞
I1V500	Supercoilin f	0.961	1	33.9	294.2	71.9	0 4.563282237
Q9VVH5	AT13736p OS-	0.947	1	266.9	0	133.1	0 2.005259204
A0A0B4KF15	Mangetout, iso	0.942	5	400	0	0	0 ∞
M9NFF9	Ada2a-contain	0.94	2	207.2	114.8	22.2	55.8 4.12805128
Q24156	Transcription f	0.937	2	84.5	56.9	232.6	25.9 0.547001934
Q94901	RNA-binding p	0.919	1	400	0	0	0 ∞
B7Z073	SOSS complex	0.898	1	0	400	0	0 ∞
J7K9E8	Sin3A (Fragme	0.852	1	0	35.2	0	364.8 0.096491228
Q5B131	LD15043p OS-	0.839	2	0	115.5	180	104.5 0.405975395
Q9VLP4	Protein mitoch	0.838	1	400	0	0	0 ∞
F0JAS9	TDD01073p (Fra)	0.828	8	0	0	400	0 0
Q9VNA3	GH21273p OS-	0.824	1	376.6	0	23.4	0 16.09401709
P26686	Serine-arginin	0.815	1	400	0	0	0 ∞
Q95U54	GH06271p OS-	0.813	4	0	0	400	0 0
E1JHR0	Drongo, isoform	0.764	1	0	0	400	0 0
Q967T6	Gag protein OS	0.757	1	155.3	136.4	0	108.3 2.693444137
A1ZBL0	Tubulin beta c	0.757	2	0	196.1	0	203.9 0.961745954
M9Z4Q3	MIP26177p1 O	0.754	1	0	0	0	0 ∞
Q8MT08	Dynein regulat	0.738	1	0	400	0	0 ∞
Q59E37	Uncharacterize	0.735	2	0	0	400	0 0
Q9WIR1	GH10609p OS-	0.725	1	141.1	0	258.9	0 0.544998069
Q9VVN4	ATR-interactin	0.719	1	0	305.1	0	94.9 3.214963119
Q86PC1	RE03224p OS-	0.68	1	0	0	400	0 0
P20007	Phosphoenolp	0.675	9	0	0	400	0 0
C6SUW3	LD13662p OS-	0.644	1	0	240.9	0	159.1 1.514142049

Appendix Table S2. List of siRNAs and primers. RNA is shown in italics.

experiment	name	sense	antisense
RNAi in OSCs	siEGFP	<i>GGCAAGCUGACCCUGAAGU TT</i>	<i>ACUUUCAGGGUCAGCUUGCC TT</i>
	siL(3)mbt	<i>CCUCUGGAGAUGGUCCGUAA TT</i>	<i>UUACGGACAUCUCCAGAGG TT</i>
	siLint-1	<i>GCAGAAUCUCCACCAGUAA TT</i>	<i>UUACUGGUGGAGAUUCUGC TT</i>
	siCoRest	<i>AACGAGGAGUCUGUAUACGAU TT</i>	<i>AUCGUUAUCAGACUCCUCGUU TT</i>
	siE2F2	<i>CCAAAUCUUGGACGUACA TT</i>	<i>UGUACGUCCAAGAUUUUGG TT</i>
	siMyb	<i>GGAGAACGACCAUUCAU TT</i>	<i>AUGGAAUGGUUCGAUUCUCC TT</i>
	siMip120	<i>GUCAUCUUUGUCCAGAAGU TT</i>	<i>UAUGUGGACCACGUUCUUC TT</i>
	siMip130	<i>GGACUCCGUUUCCAUUGCUA TT</i>	<i>UAGCAUGGAAACGGAGGUCC TT</i>
	siAtac3	<i>CGAAUGGAUGGUCCUAGAAA UU</i>	<i>UUUCUAGGACAUCCAUUCG UU</i>
	siProd	<i>CGAUUGGAAGAACACCACU UU</i>	<i>AGUGGUGUUCUUCCAAUCG UU</i>
	siCG2662(siLint-O)	<i>GAUUGAGAACGUAGCCAAA UU</i>	<i>UUUGGCUAGCUUCUCAAUC UU</i>
	siCG2199	<i>CAGACGUACUGGAGACCAU UU</i>	<i>AUGGUCUCCAGUACGUCUG UU</i>
experiment	name	forward primer (5' > 3')	reverse primer (5' > 3')
RT-qPCR	L(3)mbt	AGGAGACGTCTTCGCTTGG	AGGAGACGTCTTCGCTTGG
	Lint-1	CGCCTAACGATAGCCTACA	TATCACGAGCCATACCGAC
	CoRest	CCAGTTCCACGGGAAGAGC	CTGGCTATGGATTGTCTGGC
	E2F2	CCAAGGACCAAGAGAACTAC	ATACGACCTATTGAGGGAT
	Myb	TTGAAGATATGGATCTGCGG	GACTCTGTGTGGAGTTCT
	Mip120	GAAACCAATACCCAGCAGTA	TCTGGACAAAGATGACTTGG
	Mip130	CTGCCAGATTAAGATCTCCC	AAGTTCTCCGACACAATCTC
	Atac3	CGATGGCACAAAGCTAGTC	CTTGTCCCCTAAGCGAACTG
	Prod	ATGTCGAGGAGGATCCCCAA	ACGGCGCGCATTTAACTT
	CG2662 (siLint-O)	AAGTGTGGCAAAAGCGAAC	ACGTTGCATAACCGGCAGTA
	CG2199	TGCAAAAAGAGGTGCGATGC	ATGGGATGCTCTGTGGGTG
	Vasa	AGCCGTCCAACGATGCA	AACATCGCTGCCGGTCA
	Aub	ACGGCATTGTGACGAGTAG	ACGTCCACGAGCAATTACAG
	AGO3	CTGCATTGTGGCCTCCATA	GGGGAGTTGCCATTCTTT
	Piwi	CAAGGCCGGATAATTGGACA	CCATCGCTGGAGTGGTAAG
RT-qPCR using flies	control(heterochromatin)	CAGTTGATGGATGAATTGG	TGCCTGTGGTTCTATCCAAC
	vasa promoter	TGAATGAATCACTTAGGTTGCTTG	TGGTGBAATTCCATTGTGC
	rp49	CCGCTTCAAGGGACAGTATCTG	ATCTCGCCGCAGTAAACGC
	rp49	CCGCTTCAAGGGACAGTATCTG	ATCTCGCCGCAGTAAACGC
	CG2662 (Lint-O)	CCAAGGACGAAGACAAGGAG	CGAGGGATCGTACACCTGTT

Sequence Validation of Lint-O fly mutant	CG2662gRNA1_seq	AACTGATCTCCTTATCCAACCGATGGAAGG	GCTGCTCCGAAACAGGACTGGTTGTGGAG
experiment	name	sequence	vector
Expression vector for OSC	L3mbt-ATGR	CCCGATCCGGGGTCTCTGG	pAcF-L(3)mbt-S-3xFLAG
	L3mbt-ATG4	ATGAGCCTAATTAAATCCAAGGCCG	pAcF-L(3)mbt-S-3xFLAG
	pAcM-Lint-O_F	AACAAGCTTGGTACCAACAAATCCAAGGACGAAGA	pAcM-Lint-O
	pAcM-Lint-O_R	GACTCGAGCGGCCCTAATCTAGCCAGGCGAGGC	pAcM-Lint-O
	pAcM_F	GCGGCCGCTCGAGTCTAGAG	pAcM
	pAcM_R	GGTACCAAGCTTGTTCAGGT	pAcM
	pAc-C3F_F	CTCGAGGACTACAAAGACCA	pAc-C3F_vector
	pAc-C3F_R	CATGGTGGCTAGCCCCGATCC	pAc-C3F_vector
	pAc-L3mbt-3xFLAG_F	GGGCTAGGCCACCATGCTGCCATTGCGATGGCCAG	pAc-L3mbt-3xFLAG
	pAc-L3mbt-3xFLAG_R	TTTGAGTCCTCGAGAGAGGACGTGCGCAAGGGCG	pAc-L3mbt-3xFLAG
	pAc-Lint-O-3xFLAG_F	GGGCTAGCCACCATGAAACAAATCCAAGGACGAAGA	pAc-Lint-O-3xFLAG
	pAc-Lint-O-3xFLAG_R	TTTGAGTCCTCGAGATCTAGCCAGGGAGGCCAA	pAc-Lint-O-3xFLAG
	siRNA resistant_F	AAAATCGAAAAACTGGCTAACGCTAGACAGCCCTT	pAc-Lint-O_siRNA_resistant
	siRNA resistant_R	CTTAGCCAGTTTCGATTTCGCTGGCTTTCTT	pAc-Lint-O_siRNA_resistant
	8CA1_vector_F	GCCCCGGTTAGCCAACGTGAGGGGGCCA	pAc-Lint-O_8CA-3xFLAG
	8CA1_vector_R	GGCCTTCTGGCCGTGCGAAGTTGCG	pAc-Lint-O_8CA-3xFLAG
	8CA1_insert_F	CGGCCCAGAAGGCCGGCAAAGCGAACCCA	pAc-Lint-O_8CA-3xFLAG
	8CA1_insert_R	GTTGGCTAACGGGGCGTAGCGACACTTGTC	pAc-Lint-O_8CA-3xFLAG
	8CA2_vector_F	GCCAGTTAGCCGTGGATGCCTATCAC	pAc-Lint-O_8CA-3xFLAG
	8CA2_vector_R	GGCGCCCTGGCAGTGAGGAAGTTGCT	pAc-Lint-O_8CA-3xFLAG
	8CA2_insert_F	ACTGCCAAGGGCGCCATGCAGAAATGGCAC	pAc-Lint-O_8CA-3xFLAG
	8CA2_insert_R	CACGGCTAAACTGGCAATGGAAAGACCTGG	pAc-Lint-O_8CA-3xFLAG
	Lint-O-ΔSAM-3xFLAG_F	AAGCCATCCATCCAATGACGTTGGCCTCGC	pAc-Lint-O-ΔSAM-3xFLAG
	Lint-O-ΔSAM-3xFLAG_R	CGTCATTGGATGGATGGCTTCTCTCTA	pAc-Lint-O-ΔSAM-3xFLAG
	Lint-O-ΔPHD-3xFLAG_F	CTTCGCAACTTCGACGCGTGTAAACATCGGC	pAc-Lint-O-ΔPHD-3xFLAG
	Lint-O-ΔPHD-3xFLAG_R	GCCGATGTTACACGCGTCGAAGTTGCGAAG	pAc-Lint-O-ΔPHD-3xFLAG
	L(3)mbt-L/S_ΔSAM_F	GGACCACAGTCTGGTCGTCAGTTGATTC	pAcF-L(3)mbt-L/S_ΔSAM
	L(3)mbt-L/S_ΔSAM_R	AAACTTGAGACGACCAGGAACGTGTTGATGT	pAcF-L(3)mbt-L/S_ΔSAM

experiment	name	sequence	vector
Lint-O fly mutagenesis	gRNA_CG2662_1F	CTTCGACGACCCACAATTCCAGCGC	pBFv-U6.2-Lint-O-gRNA1
	gRNA_CG2662_1R	AAACGCGCTGGAATTGTGGGTCGTC	pBFv-U6.2-Lint-O-gRNA1
Lint-O-Venus transgenic fly construction	Lint-O gRNA5_F	CTTCGACATTAGGTGCTAAGGAGGT	pBFv-U6.2-Lint-O-gRNA5
	Lint-O gRNA5_R	AAACACCTCCTTAGCACCTAACGTC	pBFv-U6.2-Lint-O-gRNA5
	Lint-O 5'arm_F	GCTTGATATCGAATTCCCTGCCGATCACGTTCCACAAAC	pBS-Lint-Oarm-Venus-3xP3-dsRed-Express2
	Lint-O 5'arm_R	AGTTGGGGCGTAGGATCTAGCCAGGCAGGGCAACGTC	pBS-Lint-Oarm-Venus-3xP3-dsRed-Express2
	Lint-O 3'arm_F	TAGTATAGGAACTCCGCCCCACCCACCTCCTTAGCAC	pBS-Lint-Oarm-Venus-3xP3-dsRed-Express2
	Lint-O 3'arm_R	CGGGCTGCAGGAATTCTAAAAGGCGTATTTACTCCATC	pBS-Lint-Oarm-Venus-3xP3-dsRed-Express2

Appendix Table S3. Genes downregulated in L(3)mbt KD

FBgn0000036	FBgn0010265	FBgn0023081	FBgn0027836	FBgn0031169	FBgn0033086	FBgn0034543	FBgn0036035	FBgn0037648	FBgn0039274
FBgn0000064	FBgn0010340	FBgn0023091	FBgn0027873	FBgn0031233	FBgn0033088	FBgn0034617	FBgn0036038	FBgn0037652	FBgn0039303
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FBgn0000339	FBgn0010551	FBgn0023521	FBgn0028479	FBgn0031321	FBgn0033244	FBgn0034741	FBgn0036173	FBgn0037794	FBgn0039419
FBgn0000377	FBgn0010611	FBgn0023527	FBgn0028484	FBgn0031360	FBgn0033304	FBgn0034792	FBgn0036187	FBgn0037810	FBgn0039464
FBgn0000395	FBgn0010651	FBgn0024245	FBgn0028500	FBgn0031456	FBgn0033352	FBgn0034849	FBgn0036207	FBgn0037852	FBgn0039555
FBgn0000426	FBgn0010770	FBgn0024277	FBgn0028504	FBgn0031497	FBgn0033354	FBgn0034858	FBgn0036237	FBgn0037882	FBgn0039558
FBgn0000455	FBgn0010772	FBgn0024285	FBgn0028506	FBgn0031500	FBgn0033357	FBgn0034909	FBgn0036254	FBgn0037891	FBgn0039600
FBgn0000559	FBgn0010786	FBgn0024329	FBgn0028509	FBgn0031516	FBgn0033376	FBgn0034915	FBgn0036257	FBgn0037899	FBgn0039627
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FBgn0001168	FBgn0011769	FBgn0025582	FBgn0028697	FBgn0031660	FBgn0033507	FBgn0035039	FBgn0036460	FBgn0038057	FBgn0039765
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FBgn0001258	FBgn0013435	FBgn0025781	FBgn0028962	FBgn0031851	FBgn0033548	FBgn0035087	FBgn0036514	FBgn0038235	FBgn0039920
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FBgn0001316	FBgn0013749	FBgn0025809	FBgn0028990	FBgn0031885	FBgn0033556	FBgn0035121	FBgn0036570	FBgn0038271	FBgn0039993
FBgn0001324	FBgn0013750	FBgn0025830	FBgn0029006	FBgn0031893	FBgn0033557	FBgn0035131	FBgn0036571	FBgn0038272	FBgn0040011
FBgn0001341	FBgn0013756	FBgn0025832	FBgn0029114	FBgn0031904	FBgn0033571	FBgn0035137	FBgn0036579	FBgn0038275	FBgn0040022
FBgn0001565	FBgn0013762	FBgn0026015	FBgn0029117	FBgn0031914	FBgn0033592	FBgn0035142	FBgn0036614	FBgn0038294	FBgn0040064
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FBgn0002566	FBgn0014366	FBgn0026149	FBgn0029706	FBgn0032016	FBgn0033688	FBgn0035205	FBgn0036686	FBgn0038426	FBgn0040389
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FBgn0002780	FBgn0015222	FBgn0026250	FBgn0029804	FBgn0032077	FBgn0033741	FBgn0035295	FBgn0036811	FBgn0038473	FBgn0041588
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FBgn0003278	FBgn0015521	FBgn0026630	FBgn0030061	FBgn0032242	FBgn0033907	FBgn0035462	FBgn0036916	FBgn0038742	FBgn0044047
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FBgn0003366	FBgn0015754	FBgn0026666	FBgn0030092	FBgn0032248	FBgn0033924	FBgn0035524	FBgn0036922	FBgn0038746	FBgn0044452
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FBgn0004404	FBgn0016726	FBgn0027087	FBgn0030588	FBgn0032620	FBgn0034113	FBgn0035722	FBgn0037232	FBgn0038964	FBgn0050423
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FBgn0004456	FBgn0016917	FBgn0027090	FBgn0030634	FBgn0032656	FBgn0034158	FBgn0035727	FBgn0037328	FBgn0038989	FBgn0050491
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FBgn0004893	FBgn0020236	FBgn0027094	FBgn0030658	FBgn0032699	FBgn0034214	FBgn0035763	FBgn0037391	FBgn0039055	FBgn0051133
FBgn0004901	FBgn0020278	FBgn0027101	FBgn0030671	FBgn0032724	FBgn0034223	FBgn0035802	FBgn0037487	FBgn0039056	FBgn0051223
FBgn0004907	FBgn0020445	FBgn0027296	FBgn0030676	FBgn0032728	FBgn0034243	FBgn0035805	FBgn0037489	FBgn0039109	FBgn0051224
FBgn0004908	FBgn0020497	FBgn0027330	FBgn0030687	FBgn0032731	FBgn0034246	FBgn0035833	FBgn0037529	FBgn0039120	FBgn0051278
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FBgn0005278	FBgn0020764	FBgn0030720	FBgn0032799	FBgn0034259	FBgn0035868	FBgn0037537	FBgn00		

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FBgn0052594	FBgn0262125	FBgn0029896	FBgn0040259
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FBgn0052649	FBgn0262160	FBgn0030870	FBgn0042173
FBgn0052672	FBgn0262166	FBgn0031011	FBgn0045843
FBgn0053052	FBgn0262527	FBgn0031530	FBgn0051547
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FBgn0053158	FBgn0262647	FBgn0032723	FBgn0052641
FBgn0053180	FBgn0262733	FBgn0034389	FBgn0052655
FBgn0061469	FBgn0262954	FBgn0034501	FBgn0063491
FBgn0061476	FBgn0262955	FBgn0034734	FBgn0085249
FBgn0063492	FBgn0263110	FBgn0034903	FBgn0085407
FBgn0063493	FBgn0263120	FBgn0035023	FBgn0263116
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FBgn0064766	FBgn0263316	FBgn0037613	FBgn0266261
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FBgn0085294	FBgn0263510	FBgn0038720	
FBgn0085339	FBgn0263594	FBgn0039209	
FBgn0085376	FBgn0263599	FBgn0039257	
FBgn0085437	FBgn0263605	FBgn0039266	
FBgn0085451	FBgn0263606	FBgn0039829	
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FBgn0086251	FBgn0263929	FBgn0040299	
FBgn0086347	FBgn0263995	FBgn0040496	
FBgn0086371	FBgn0264006	FBgn0043841	
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FBgn0086451	FBgn0264560	FBgn0052311	
FBgn0086472	FBgn0264751	FBgn0062978	
FBgn0086679	FBgn0264978	FBgn0082585	
FBgn0086687	FBgn0265003	FBgn0086906	
FBgn0086706	FBgn0265187	FBgn0243512	
FBgn0086710	FBgn0265413	FBgn0250753	
FBgn0086711	FBgn0265574	FBgn0259245	
FBgn0086895	FBgn0265998	FBgn0259935	
FBgn0086912	FBgn0266019	FBgn0259938	
FBgn0243511	FBgn0266053	FBgn0260660	
FBgn0250785	FBgn0266268	FBgn0263219	
FBgn0250814	FBgn0266284	FBgn0263256	
FBgn0250868	FBgn0266409	FBgn0264907	
FBgn0259152	FBgn0266411	FBgn0266099	
FBgn0259170	FBgn0266581	FBgn0267796	
FBgn0259481	FBgn0266668	FBgn0285963	
FBgn0259711	FBgn0266673	FBgn0000037	
FBgn0259742	FBgn0266674	FBgn002576	
FBgn0259821	FBgn0266696	FBgn002645	
FBgn0259832	FBgn0266717	FBgn002783	
FBgn0259937	FBgn0266722	FBgn0002930	
FBgn0259984	FBgn0266723	FBgn0003285	
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FBgn0260456	FBgn0267976	FBgn0019990	
FBgn0260632	FBgn0270926	FBgn0027505	
FBgn0260648	FBgn0283427	FBgn0028888	
FBgn0260745	FBgn0283468	FBgn0029697	
FBgn0260755	FBgn0283472	FBgn0030438	
FBgn0260855	FBgn0283473	FBgn0030693	
FBgn0260857	FBgn0283510	FBgn0030882	
FBgn0260965	FBgn0283659	FBgn0030960	
FBgn0260990	FBgn0283680	FBgn0030983	
FBgn0261020	FBgn0283681	FBgn0031227	
FBgn0261049	FBgn0283724	FBgn0031728	
FBgn0261068	FBgn0283741	FBgn0031857	
FBgn0261258	FBgn0284245	FBgn0031888	
FBgn0261274	FBgn0284246	FBgn0032139	
FBgn0261276	FBgn0284248	FBgn0032258	
FBgn0261285	FBgn0284252	FBgn0032259	
FBgn0261286	FBgn0285911	FBgn0032645	
FBgn0261380	FBgn0285954	FBgn0032969	
FBgn0261444	FBgn0285955	FBgn0033190	
FBgn0261530	FBgn0285962	FBgn0033696	
FBgn0261535	FBgn0286075	FBgn0033697	
FBgn0261550	FBgn0286786	FBgn0033725	
FBgn0261554	FBgn0286813	FBgn0033888	
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FBgn0261800	FBgn000326	FBgn0036970	
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FBgn0261933	FBgn003507	FBgn0038680	
FBgn0261934	FBgn0017581	FBgn0039494	

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FBgn0032055	FBgn0259216	FBgn0031920	FBgn0038261
FBgn0032218	FBgn0260874	FBgn0031974	FBgn0038315
FBgn0032345	FBgn0261016	FBgn0032003	FBgn0038407
FBgn0032726	FBgn0261859	FBgn0032082	FBgn0038435
FBgn0032749	FBgn0261996	FBgn0032178	FBgn0038492
FBgn0033096	FBgn0262057	FBgn0032225	FBgn0038530
FBgn0033110	FBgn0262477	FBgn0032252	FBgn0038761
FBgn0033203	FBgn0262509	FBgn0032253	FBgn0038957
FBgn0033222	FBgn0262531	FBgn0032262	FBgn0039006
FBgn0033246	FBgn0262867	FBgn0032374	FBgn0039124
FBgn0033657	FBgn0262871	FBgn0032375	FBgn0039282
FBgn0033904	FBgn0263111	FBgn0032422	FBgn0039294
FBgn0033963	FBgn0263219	FBgn0032601	FBgn0039321
FBgn0034052	FBgn0264815	FBgn0032683	FBgn0039504
FBgn0034094	FBgn0264895	FBgn0032774	FBgn0039629
FBgn0034247	FBgn0265045	FBgn0032783	FBgn0039655
FBgn0034489	FBgn0265052	FBgn0032803	FBgn0039678
FBgn0034497	FBgn0265274	FBgn0032900	FBgn0039818
FBgn0034706	FBgn0266129	FBgn0032972	FBgn0039911
FBgn0034943	FBgn0266709	FBgn0032973	FBgn0039915
FBgn0034966	FBgn0284435	FBgn0033051	FBgn0039927
FBgn0034974	FBgn0000146	FBgn0033079	FBgn0040370
FBgn0034985	FBgn0000153	FBgn0033113	FBgn0041233
FBgn0035034	FBgn0001259	FBgn0033115	FBgn0041710
FBgn0035094	FBgn0001296	FBgn0033221	FBgn0042696
FBgn0035103	FBgn0001991	FBgn0033234	FBgn0043043
FBgn0035234	FBgn0003137	FBgn0033268	FBgn0043575
FBgn0035264	FBgn0003308	FBgn0033387	FBgn0043806
FBgn0035505	FBgn0003475	FBgn0033658	FBgn0045980
FBgn0035612	FBgn0003486	FBgn0033659	FBgn0050345
FBgn0035903	FBgn0003886	FBgn0033753	FBgn0050485
FBgn0036195	FBgn0003890	FBgn0033763	FBgn0051017
FBgn0036282	FBgn0004118	FBgn0033875	FBgn0051370
FBgn0036479	FBgn0004512	FBgn0033917	FBgn0051436
FBgn0036568	FBgn0005427	FBgn0034047	FBgn0051463
FBgn0036727	FBgn0005666	FBgn0034279	FBgn0051464
FBgn0036806	FBgn0010241	FBgn0034413	FBgn0051997
FBgn0036875	FBgn0010258	FBgn0034530	FBgn0052006
FBgn0036978	FBgn0010473	FBgn0034538	FBgn0052037
FBgn0036993	FBgn0011296	FBgn0034602	FBgn0052073
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FBgn0037618	FBgn0011828	FBgn0034789	FBgn0052783
FBgn0037714	FBgn0013348	FBgn0034928	FBgn0053481
FBgn0037717	FBgn0013813	FBgn0034956	FBgn0053775
FBgn0037750	FBgn0014000	FBgn0034957	FBgn0058470
FBgn0037751	FBgn0014395	FBgn0034972	FBgn0065110
FBgn0037916	FBgn0014396	FBgn0035056	FBgn0069969
FBgn0038076	FBgn0014417	FBgn0035104	FBgn0085245
FBgn0038079	FBgn0015032	FBgn0035286	FBgn0085298
FBgn0038290	FBgn0015519	FBgn0035317	FBgn0085483
FBgn0038330	FBgn0020379	FBgn0035412	FBgn0087040
FBgn0038602	FBgn0022709	FBgn0035434	FBgn0250819
FBgn0038740	FBgn0023416	FBgn0035571	FBgn0250904
FBgn0039201	FBgn0023495	FBgn0035880	FBgn0250907
FBgn0039415	FBgn0024732	FBgn0035904	FBgn0259977
FBgn0039500	FBgn0026144	FBgn0035985	FBgn0260430
FBgn0039594	FBgn0026562	FBgn0036040	FBgn0260658
FBgn0039651	FBgn0026593	FBgn0036131	FBgn0260866
FBgn0039789	FBgn0027070	FBgn0036168	FBgn0261341
FBgn0039886	FBgn0027073	FBgn0036208	FBgn0261393
FBgn0039897	FBgn0027521	FBgn0036454	FBgn0261561
FBgn0039932	FBgn0027550	FBgn0036547	FBgn0262035
FBgn0040091	FBgn0028988	FBgn0036575	FBgn0262794
FBgn0040096	FBgn0029091	FBgn0036687	FBgn0264443
FBgn0040372	FBgn0029710	FBgn0036738	FBgn0264894
FBgn0041234	FBgn0029856	FBgn0036752	FBgn0265048
FBgn0041723	FBgn0029986	FBgn0036765	FBgn0265137
FBgn0042133	FBgn0029987	FBgn0036890	FBgn0265417
FBgn0050502	FBgn0030028	FBgn0036891	FBgn0265767
FBgn0051161	FBgn0030244	FBgn0036910	FBgn0266534
FBgn0051202	FBgn0030310	FBgn0037028	FBgn0266566
FBgn0051231	FBgn0030590	FBgn0037060	FBgn0267408
FBgn0051279	FBgn0030653	FBgn0037222	FBgn0285896
FBgn0051706	FBgn0030691	FBgn0037225	FBgn0287209
FBgn0052036	FBgn0030748	FBgn0037304	
FBgn0053099	FBgn0030859	FBgn0037347	
FBgn0053257	FBgn0031004	FBgn0037375	
FBgn0053468	FBgn0031018	FBgn0037563	
FBgn0053519	FBgn0031157	FBgn0037659	
FBgn0061356	FBgn0031184	FBgn0037807	
FBgn0083960	FBgn0031255	FBgn0037835	
FBgn0085384	FBgn0031343	FBgn0037915	
FBgn0085421	FBgn0031515	FBgn0037934	
FBgn0085431	FBgn0031548	FBgn0037973	
FBgn0085434	FBgn0031715	FBgn0037974	
FBgn0086906	FBgn0031716	FBgn0037975	
FBgn0250816	FBgn0031737	FBgn0038017	
FBgn0250869	FBgn0031869	FBgn0038067	

Appendix Table S3. Genes downregulated in Lint-O KD

FBgn0000116	FBgn0011726	FBgn0026666	FBgn0031651	FBgn0034447	FBgn0036686	FBgn0039419	FBgn0261380	FBgn0030983
FBgn0000181	FBgn0011737	FBgn0027055	FBgn0031660	FBgn0034451	FBgn0036702	FBgn0039488	FBgn0261535	FBgn0031227
FBgn0000250	FBgn0011762	FBgn0027057	FBgn0031673	FBgn0034494	FBgn0036734	FBgn0039602	FBgn0261560	FBgn0031805
FBgn0000253	FBgn0013269	FBgn0027079	FBgn0031710	FBgn0034528	FBgn0036754	FBgn0039626	FBgn0261565	FBgn0031857
FBgn0000261	FBgn0013432	FBgn0027081	FBgn0031736	FBgn0034534	FBgn0036805	FBgn0039636	FBgn0261596	FBgn0031888
FBgn0000326	FBgn0014020	FBgn0027082	FBgn0031832	FBgn0034579	FBgn0036828	FBgn0039654	FBgn0261599	FBgn0033592
FBgn0000370	FBgn0014028	FBgn0027086	FBgn0031851	FBgn0034617	FBgn0036838	FBgn0039735	FBgn0261822	FBgn0033725
FBgn0000395	FBgn0014163	FBgn0027088	FBgn0031881	FBgn0034631	FBgn0036886	FBgn0039740	FBgn0261931	FBgn0035158
FBgn0000412	FBgn0014366	FBgn0027090	FBgn0031902	FBgn0034703	FBgn0036892	FBgn0039765	FBgn0261933	FBgn0035189
FBgn0000426	FBgn0014411	FBgn0027091	FBgn0031992	FBgn0034724	FBgn0036916	FBgn0039828	FBgn0261984	FBgn0035281
FBgn0000560	FBgn0014877	FBgn0027094	FBgn0032016	FBgn0034741	FBgn0036973	FBgn0039868	FBgn0262081	FBgn0035727
FBgn0000579	FBgn0015000	FBgn0027330	FBgn0032034	FBgn0034792	FBgn0037011	FBgn0039909	FBgn0262169	FBgn0036597
FBgn0001092	FBgn0015024	FBgn0027554	FBgn0032050	FBgn0034802	FBgn0037012	FBgn0039929	FBgn0262527	FBgn0037672
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FBgn0001168	FBgn0015222	FBgn0027599	FBgn0032138	FBgn0034914	FBgn0037073	FBgn0040066	FBgn0262734	FBgn0040259
FBgn0001186	FBgn0015229	FBgn0027619	FBgn0032198	FBgn0034915	FBgn0037081	FBgn0040075	FBgn0262736	FBgn0040609
FBgn0001202	FBgn0015245	FBgn0027785	FBgn0032217	FBgn0034918	FBgn0037102	FBgn0040271	FBgn0262954	FBgn0041087
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FBgn0001218	FBgn0015288	FBgn0027932	FBgn0032229	FBgn0034982	FBgn0037248	FBgn0040319	FBgn0263200	FBgn0043841
FBgn0001220	FBgn0015527	FBgn0028331	FBgn0032236	FBgn0035020	FBgn0037293	FBgn0040389	FBgn0263396	FBgn0051547
FBgn0001247	FBgn0015754	FBgn0028341	FBgn0032248	FBgn0035027	FBgn0037382	FBgn0040394	FBgn0263510	FBgn0063491
FBgn0001258	FBgn0015765	FBgn0028429	FBgn0032258	FBgn0035039	FBgn0037391	FBgn0040529	FBgn0263594	FBgn0259243
FBgn0001301	FBgn0015776	FBgn0028473	FBgn0032261	FBgn0035063	FBgn0037487	FBgn0040985	FBgn0263605	FBgn0263118
FBgn0001316	FBgn0015778	FBgn0028504	FBgn0032296	FBgn0035085	FBgn0037489	FBgn0041147	FBgn0263740	FBgn0264907
FBgn0001341	FBgn0015790	FBgn0028527	FBgn0032321	FBgn0035121	FBgn0037543	FBgn0041184	FBgn0263773	
FBgn0001961	FBgn0015795	FBgn0028690	FBgn0032346	FBgn0035131	FBgn0037555	FBgn0041210	FBgn0263929	
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FBgn0002413	FBgn0015829	FBgn0028836	FBgn0032408	FBgn0035165	FBgn0037608	FBgn0041629	FBgn0264294	
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FBgn0003231	FBgn0016797	FBgn0029133	FBgn0032514	FBgn0035232	FBgn0037743	FBgn0046214	FBgn0265935	
FBgn0003257	FBgn0017578	FBgn0029664	FBgn0032586	FBgn0035295	FBgn0037773	FBgn0046685	FBgn0266284	
FBgn0003274	FBgn0019624	FBgn0029672	FBgn0032587	FBgn0035336	FBgn0037794	FBgn0050000	FBgn0266581	
FBgn0003275	FBgn0020238	FBgn0029689	FBgn0032620	FBgn0035337	FBgn0037852	FBgn0050035	FBgn0266668	
FBgn0003278	FBgn0020278	FBgn0029755	FBgn0032703	FBgn0035346	FBgn0037891	FBgn005104	FBgn0266669	
FBgn0003310	FBgn0020388	FBgn0029766	FBgn0032728	FBgn0035357	FBgn0037899	FBgn0050159	FBgn0266673	
FBgn0003360	FBgn0020414	FBgn0029785	FBgn0032731	FBgn0035383	FBgn0037958	FBgn0050349	FBgn0266720	
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FBgn0004050	FBgn0022069	FBgn0030208	FBgn0032981	FBgn0035541	FBgn0038294	FBgn0052176	FBgn0284251	
FBgn0004177	FBgn0022238	FBgn0030291	FBgn0032988	FBgn0035589	FBgn0038306	FBgn0052267	FBgn0284253	
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FBgn0004638	FBgn0023000	FBgn0030569	FBgn0033260	FBgn0035656	FBgn0038455	FBgn0052649	FBgn0000308	
FBgn0004657	FBgn0023091	FBgn0030588	FBgn0033304	FBgn0035714	FBgn0038472	FBgn0052672	FBgn0002645	
FBgn0004828	FBgn0023177	FBgn0030598	FBgn0033379	FBgn0035760	FBgn0038473	FBgn0053100	FBgn0005775	
FBgn0004901	FBgn0023212	FBgn0030603	FBgn0033395	FBgn0035805	FBgn0038488	FBgn0053158	FBgn0014388	
FBgn0004907	FBgn0023214	FBgn0030605	FBgn0033428	FBgn0035853	FBgn0038519	FBgn0053180	FBgn024183	
FBgn0004926	FBgn0023514	FBgn0030608	FBgn0033454	FBgn0035866	FBgn0038585	FBgn0053299	FBgn0026160	
FBgn0005278	FBgn0023519	FBgn0030672	FBgn0033483	FBgn0035877	FBgn0038593	FBgn0053542	FBgn0026319	
FBgn0005616	FBgn0023527	FBgn0030687	FBgn0033485	FBgn0035878	FBgn0038666	FBgn0062449	FBgn0028541	
FBgn0005648	FBgn0023542	FBgn0030692	FBgn0033507	FBgn0035901	FBgn0038678	FBgn0085339	FBgn0029896	
FBgn0005672	FBgn0024245	FBgn0030720	FBgn0033540	FBgn0036000	FBgn0038742	FBgn0086347	FBgn0030318	
FBgn0001038	FBgn0024321	FBgn0030752	FBgn0033544	FBgn0036018	FBgn0038746	FBgn0086443	FBgn0030870	
FBgn0010173	FBgn0024362	FBgn0030850	FBgn0033547	FBgn0036035	FBgn0038768	FBgn0086656	FBgn0032640	
FBgn0010220	FBgn0024364	FBgn0030853	FBgn0033548	FBgn0036038	FBgn0038787	FBgn0086711	FBgn0035023	
FBgn0010235	FBgn0024556	FBgn0030854	FBgn0033571	FBgn0036124	FBgn0038854	FBgn0086855	FBgn0035049	
FBgn0010238	FBgn0024993	FBgn0030863	FBgn0033649	FBgn0036126	FBgn0038864	FBgn0086904	FBgn0035101	
FBgn0010278	FBgn0025366	FBgn0030894	FBgn0033737	FBgn0036173	FBgn0038890	FBgn0086912	FBgn0035470	
FBgn0010333	FBgn0025394	FBgn0030904	FBgn0033741	FBgn0036187	FBgn0038893	FBgn00250785	FBgn0036316	
FBgn0010340	FBgn0025582	FBgn0030969	FBgn0033816	FBgn0036207	FBgn0038964	FBgn00259152	FBgn0039257	
FBgn0010348	FBgn0025608	FBgn0031040	FBgn0033907	FBgn0036237	FBgn0038989	FBgn00259481	FBgn0039829	
FBgn0010389	FBgn0025674	FBgn0031043	FBgn0033918	FBgn0036258	FBgn0039004	FBgn00259711	FBgn0039858	
FBgn0010409	FBgn0025781	FBgn0031047	FBgn0033989	FBgn0036333	FBgn0039109	FBgn00259742	FBgn0250753	
FBgn0010504	FBgn0025809	FBgn0031057	FBgn0034029	FBgn0036334	FBgn0039175	FBgn00259832	FBgn0263256	
FBgn0010551	FBgn0025830	FBgn0031114	FBgn0034049	FBgn0036350	FBgn0039205	FBgn00259878	FBgn0285963	
FBgn0010611	FBgn0025832	FBgn0031238	FBgn0034065	FBgn0036512	FBgn0039207	FBgn00259937	FBgn0260501	
FBgn0010651	FBgn0025879	FBgn0031281	FBgn0034086	FBgn0036514	FBgn0039223	FBgn0260407	FBgn0000037	
FBgn0010770	FBgn0026079	FBgn0031321	FBgn0034113	FBgn0036516	FBgn0039252	FBgn0260456	FBgn0002783	
FBgn0010774	FBgn0026084	FBgn0031392	FBgn0034118	FBgn0036522	FBgn0039259	FBgn0260632	FBgn0003285	
FBgn0010786	FBgn0026196	FBgn0031456	FBgn0034232	FBgn0036570	FBgn0039274	FBgn0260648	FBgn0004034	
FBgn0010803	FBgn0026250	FBgn0031493	FBgn0034243	FBgn0036580	FBgn0039303	FBgn0260755	FBgn0004436	
FBgn0010808	FBgn0026252	FBgn0031500	FBgn0034259	FBgn0036614	FBgn0039306	FBgn0260855	FBgn0015037	
FBgn0010926	FBgn0026317	FBgn0031540	FBgn0034345	FBgn0036627	FBgn0039350	FBgn0260936	FBgn0022349	
FBgn0011204	FBgn0026372	FBgn0031597	FBgn0034401	FBgn0036661	FBgn0039358	FBgn0261258	FBgn0029804	
FBgn0011638	FBgn0026378	FBgn0031643	FBgn0034432	FBgn0036666	FBgn0039407	FBgn0261286	FBgn0030373	

Appendix table S4

Related to Figs 5-6 and Figs EV5. Experimental genotypes used in this study.

Fig 5B-5E

yw	$y^1 w^{1118}$
$Lint-O^{KO}$	$y^1 w^{1118} Lint-O^{KO} / y^1 w^{1118} Lint-O^{KO}$

Fig 6A, 6B, 6D-6G

yw	$y^1 w^{1118},$ $y^1 w^{1118} / Y$
$Lint-O^{KO}$	$y^1 w^{1118} Lint-O^{KO} / y^1 w^{1118} Lint-O^{KO},$ $y^1 w^{1118} Lint-O^{KO} / Y$
$L(3)mbt^{ts1}$	$w^* ; + ; L(3)mbt^{ts1} / L(3)mbt^{ts1},$ $w^* / Y ; + ; L(3)mbt^{ts1} / L(3)mbt^{ts1}$

Fig 6C

$Lint-O^{KO} -/+$	$y^1 w^{1118} Lint-O^{KO} / FM7 Kruppel>GFP,$ $FM7 Kruppel>GFP / FM7 Kruppel>GFP,$ $FM7 Kruppel>GFP / Y$
$Lint-O^{KO} -/-$	$y^1 w^{1118} Lint-O^{KO} / y^1 w^{1118} Lint-O^{KO},$ $y^1 w^{1118} Lint-O^{KO} / Y$
$L(3)mbt^{ts1} -/-$	$w^* ; + ; L(3)mbt^{ts1} / L(3)mbt^{ts1},$ $w^* / Y ; + ; L(3)mbt^{ts1} / L(3)mbt^{ts1}$

Fig EV5

yw	$y^1 w^{1118},$ $y^1 w^{1118} / Y$
$Lint-O^{KO}$	$y^1 w^{1118} Lint-O^{KO} / y^1 w^{1118} Lint-O^{KO},$ $y^1 w^{1118} Lint-O^{KO} / Y$
$L(3)mbt^{ts1}$	$w^* ; + ; L(3)mbt^{ts1} / L(3)mbt^{ts1},$ $w^* / Y ; + ; L(3)mbt^{ts1} / L(3)mbt^{ts1}$
$yw Lint-O-Venus$	$y^1 w^{1118} Lint-O-Venus / y^1 w^{1118} Lint-O-Venus$