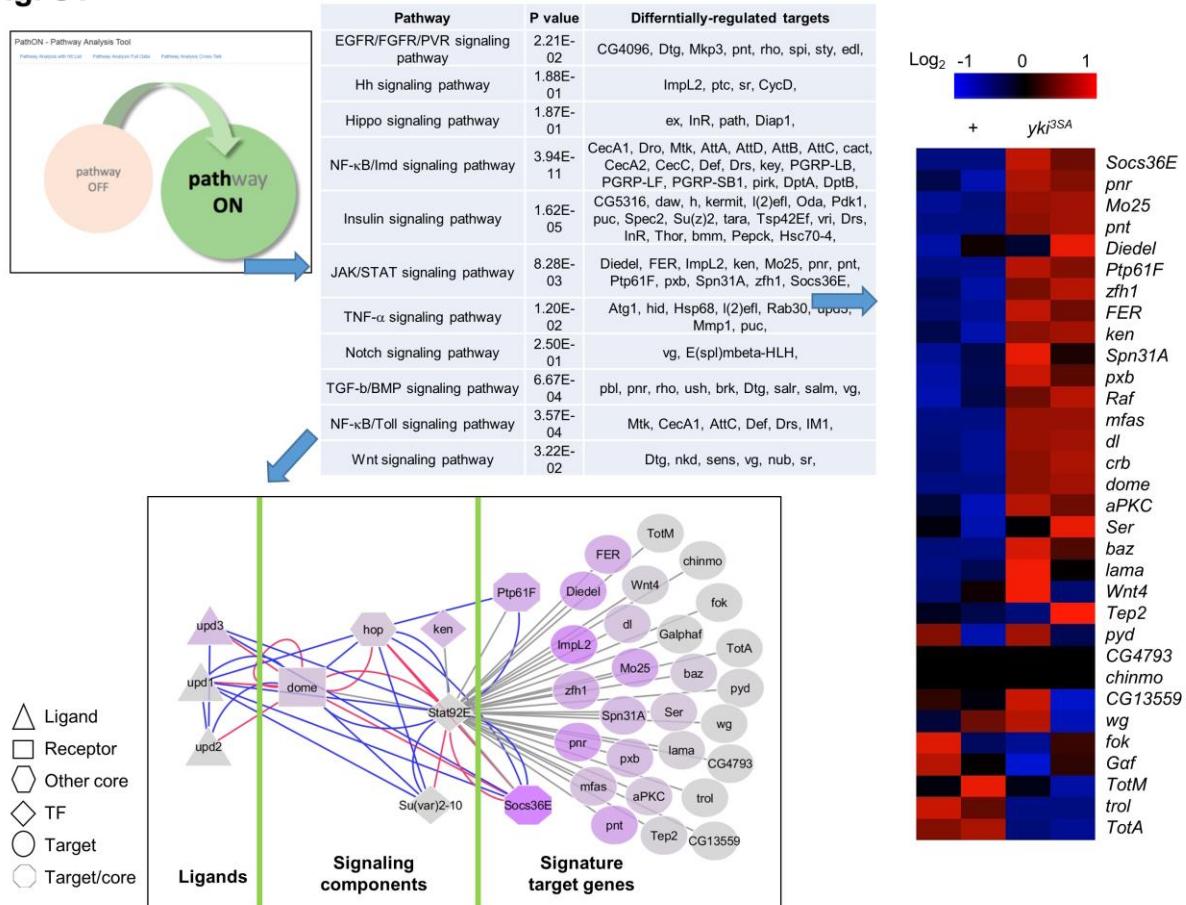


Supplemental information

**Coordination of tumor growth
and host wasting by tumor-derived Upd3**

Guangming Ding, Xiaoxiang Xiang, Yanhui Hu, Gen Xiao, Yuchen Chen, Richard Binari, Aram Comjean, Jiaying Li, Elisabeth Rushworth, Zhenming Fu, Stephanie E. Mohr, Norbert Perrimon, and Wei Song

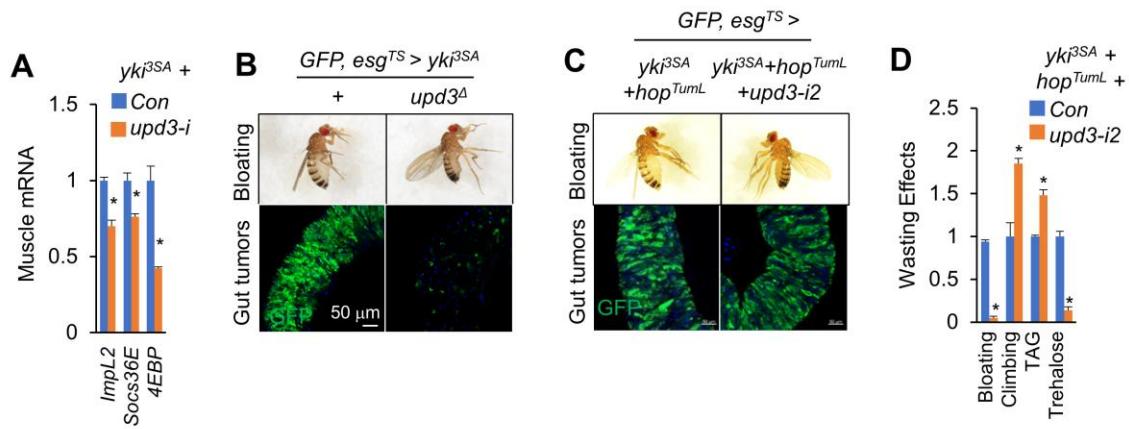
Fig. S1



Supplemental Figure S1. Using PathOn to analyze pathway activation. Related to Figure 1.

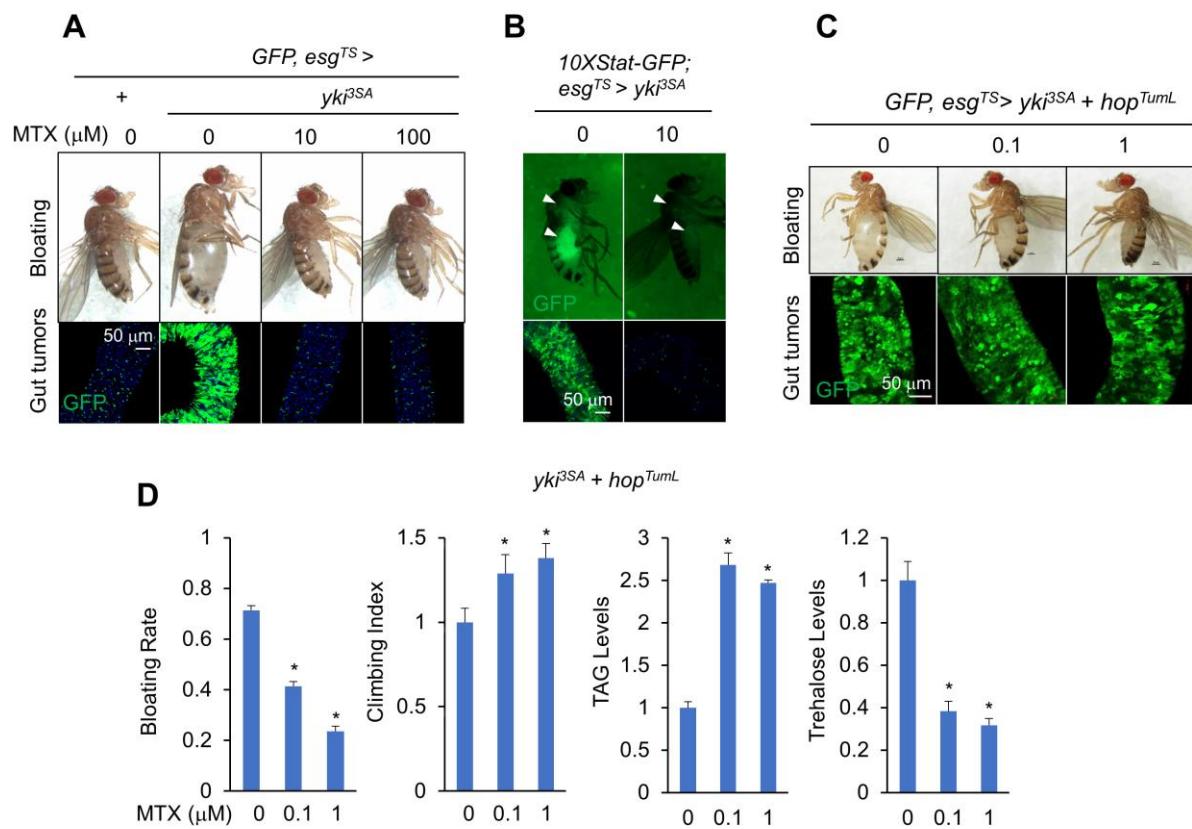
PathOn is a web-based analysis tool useful to predict the ON/OFF status of 14 major signaling pathways by analyzing the enrichment of pathway signaling components or signature target genes based on expression profiles provided by the user. The RNA-seq dataset obtained from the muscle of yki-tumor bearing files, as well as a dataset from control flies, were uploaded and analyzed by PathON. Enrichment *p*-values of signature target genes were calculated using a hypergeometric test. PathOn predicts that the Jak/Stat signaling pathway is activated. A heatmap illustration and a network illustration are also provided by PathOn to visualize the transcriptional changes in mutant vs control samples.

Fig. S2



Supplemental Figure S2. *upd3* knockdown using another RNAi line in *yki^{3SA}+hop^{TumL}* tumors alleviates host wasting. Related to Figures 1 and 2. (A) Gene expression levels in the muscle were analyzed with qPCR at day 8 (n=3, 10 flies/replicate). (B) Bloating phenotypes (up) and gut tumors (bottom, scale bar, 50 μm) of flies bearing *yki^{3SA}* tumors with or without systemic *upd3* mutation at day 8. (C-D) Wasting effects of flies bearing *yki^{3SA}+hop^{TumL}* tumors with or without *upd3* RNAi (HM05061) at day 6: Bloating phenotype (up) and gut tumors (bottom, scale bar, 50 μm) (C), bloating rates (n=4, 20 flies/replicate), climbing rates (n=15), TAG and trehalose storages (n=3, 5 flies/replicate) (D). Data are presented as mean ± SEM. *p < 0.05.

Fig. S3



Supplemental Figure S3. Pharmaceutical Jak/Stat inhibition alleviates host wasting. Related to Figure 2. (A-C)

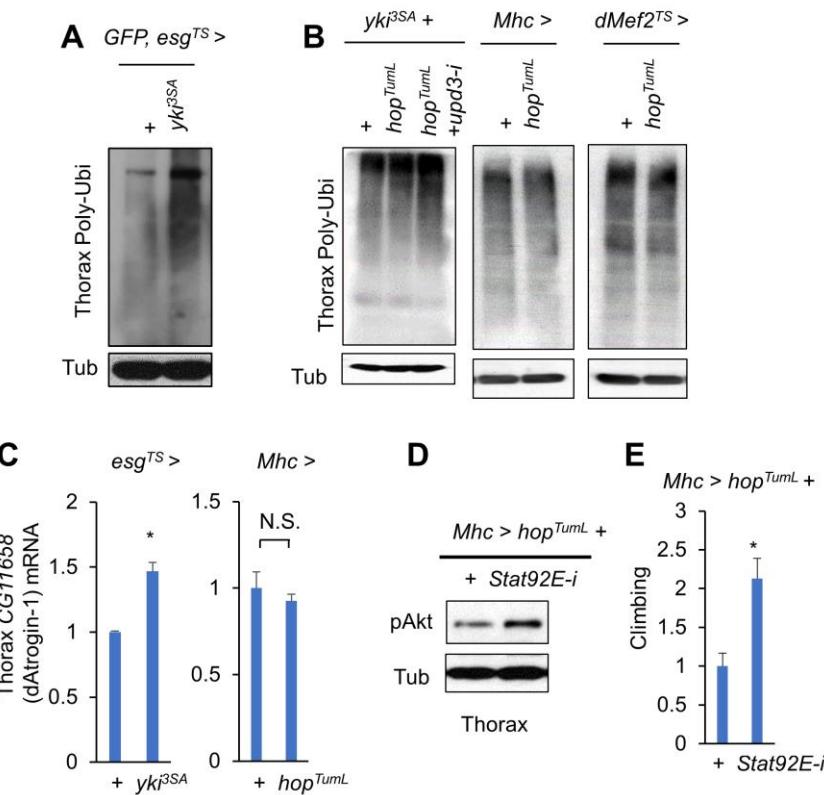
Bloating phenotypes (**A, C**, up) and gut tumors (**A, C**, bottom, scale bar, 50 μm) and 10XStat-GFP signals in the whole

body and midgut (**B**, scale bar, 50 μm) under MTX treatment simultaneously with tumor induction at day 6. (**D**)

Wasting effects such as bloating rates (n=4, 20 flies/replicate), climbing rates (n=15), and TAG and Trehalose storages

(n=3, 5 flies/replicate) at day 6. Data are presented as mean ± SEM. * p < 0.05.

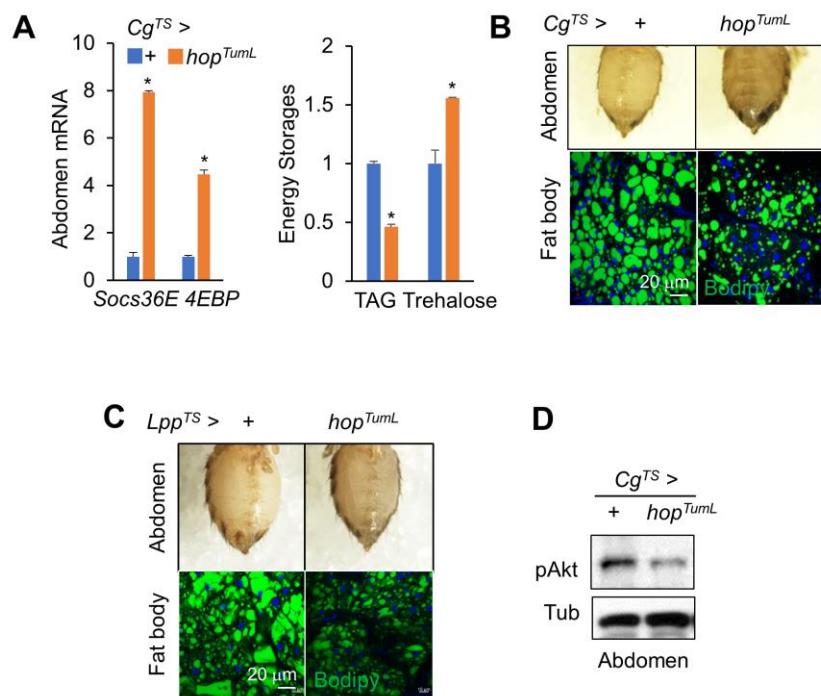
Fig. S4



Supplemental Figure S4. Jak/Stat signaling does not regulate proteasome-associated protein degradation.

Related to Figures 3 and 4. (A-B) Polyubiquitin proteins in adult thoracic muscles (A, day 8) (B, left, day 6; middle, day 4; right, day 4). (C) mRNA levels of *CG11658* in the thoracic muscles of indicated genotypes (left, day 8; right, day 4) (n=3, 5 flies/replicate). (D-E) Thoracic muscle p-Akt levels (D) and fly climbing rates (E, n=15) of adult flies of indicated genotypes at day 4. Data are presented as mean ± SEM. *p < 0.05.

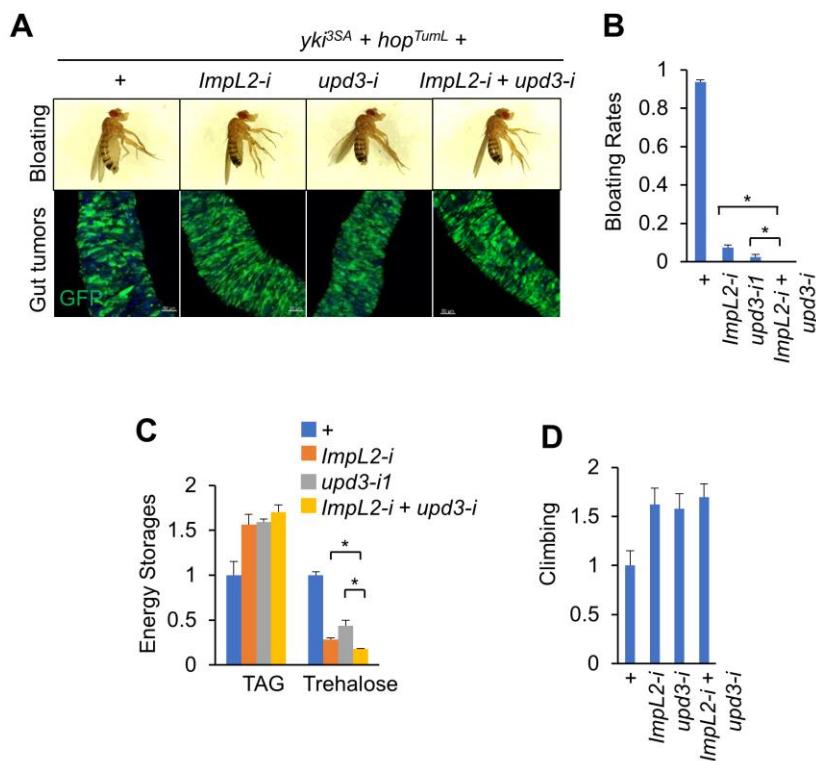
Fig. S5



Supplemental Figure S5. Fat body Jak/Stat signaling regulates lipid loss and hyperglycemia. Related to Figure 6.

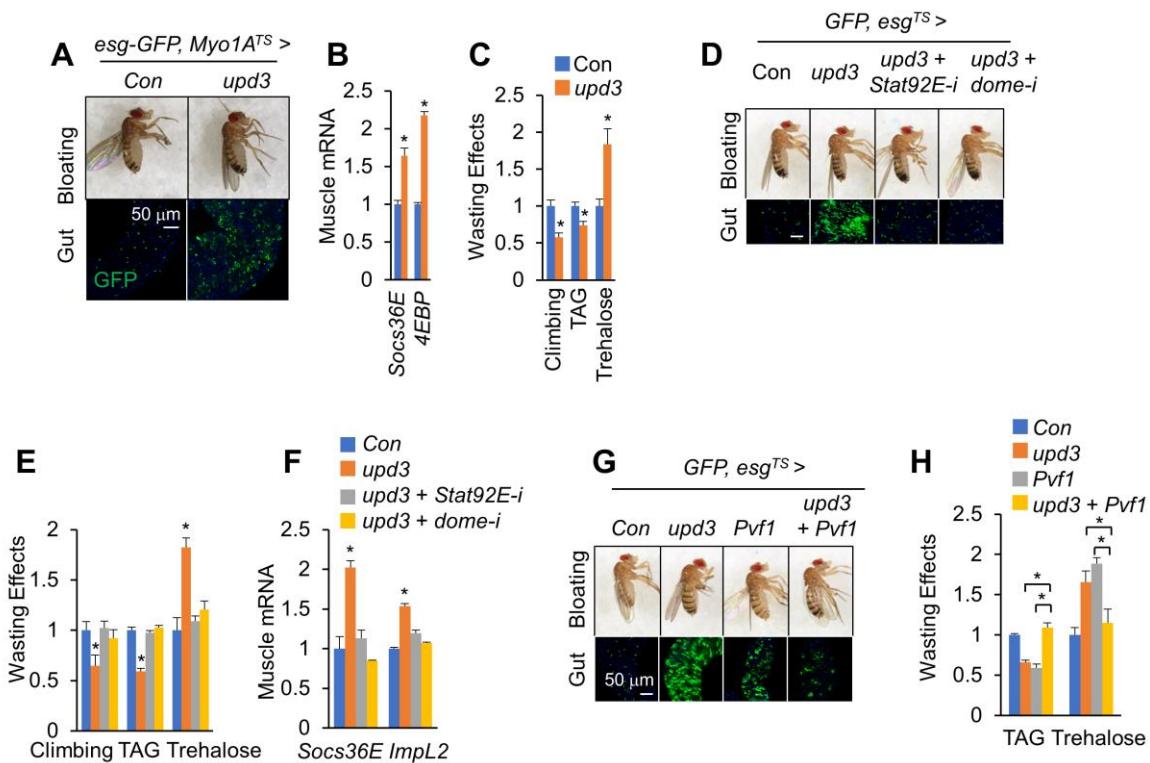
Gene expression in abdomens containing fat bodies (A, left, n=3, 5 flies/replicate), TAG and Trehalose storages (A, right, n=3, 5 flies/replicate), abdomen appearance (B-C, up), Bodipy-labeled lipid droplets in the abdominal fat body (B-C, bottom, scale bar, 20 μ m), as well as abdomen insulin signaling as indicated by p-Akt (D), in flies with activation of Jak/Stat signaling in the fat body at day 4. Data are presented as mean \pm SEM. *p < 0.05.

Fig. S6



Supplemental Figure S6. Double knockdown of *ImpL2* and *Upd3* in *yki3^{SA}+hop^{TumL}* gut tumors further alleviates host wasting. Related to Figure 2. (A-D) Wasting effects of flies bearing *yki3^{SA}+hop^{TumL}* tumors with *ImpL2* RNAi (NIG 15009R-3) and/or *upd3* RNAi (HMS00646) at day 6: (A) Bloating phenotypes (up) and gut tumors (bottom, scale bar, 50 μ m), (B) bloating rates (n=4, 20 flies/replicate), (C) TAG and trehalose levels (n=3, 5 flies/replicate), (D) climbing rates (n=15). Data are presented as mean \pm SEM. *p < 0.05.

Fig. S7



Supplemental Figure S7. Gut-derived Upd3 results in host Jak/Stat activation and organ wasting. Related to Figure 7.

Figure 7. Fly appearances (A, D, G, up), gut tumors (A, D, G, bottom, scale bar, 50 μm), thoracic muscle gene expression (B, F, n=3, 5 flies/replicate), and wasting effects such as climbing (n=15), TAG and Trehalose storages (C, E, H, n=3, 5 flies/replicate) of adult flies with indicated genotypes at day 8. *Stat92E* RNAi (HMS00035, III) and *dome* RNAi (v19717) were used. Data are presented as mean ± SEM. *p < 0.05.

Supplemental Table S3. Genotypes in this study. Related to each Figure.

Figure 1	
A-C,	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP</i>
E-F	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i>
D	<i>10XStat-GFP; esg-GAL4, tub-GAL80^{TS} > +</i> <i>10XStat-GFP; esg-GAL4, tub-GAL80^{TS} > UAS-yki^{3SA}</i>
G	<i>upd3-LacZ; esg-GAL4, tub-GAL80^{TS} > +</i> <i>upd3-LacZ; esg-GAL4, tub-GAL80^{TS} > UAS-yki^{3SA}</i>
H	<i>hs-Flp, FRT19A, upd3-LacZ; tub-Gal4, tub-GAL80 > UAS-GFP</i> <i>hs-Flp, FRT19A, upd3-LacZ; tub-Gal4, tub-GAL80 > UAS-GFP, UAS-yki^{3SA}</i>
I	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-upd3-i1</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-upd3-i2</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-dome-i</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop-i</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-Stat92E-i</i>

Figure 2	
B-E	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}, UAS-upd3-i1</i>

Figure 3	
A-C	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}, UAS-upd3-i1</i>
D-F	<i>Mhc-GAL4 > +</i> <i>Mhc-GAL4 > UAS-hop^{TumL}</i>

Figure 4	
A-B	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}, UAS-upd3-i1</i>
C-G	<i>Mhc-GAL4 > +</i> <i>Mhc-GAL4 > UAS-hop^{TumL}</i> <i>Mhc-GAL4 > UAS-hop^{TumL}, UAS-InR^{AC}</i> <i>dMef2-GAL4, tub-GAL80^{TS} > +</i> <i>dMef2-GAL4, tub-GAL80^{TS} > UAS-hop^{TumL}</i>

Figure 5	
E	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP</i>

	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i>
F	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}, UAS-upd3-i1</i>
G	<i>Mhc-GAL4 > +</i> <i>Mhc-GAL4 > UAS-hop^{TumL}</i> <i>Mhc-GAL4 > UAS-hop^{TumL}, UAS-Stat92E-i</i>

Figure 6

A-C	<i>Mhc-GAL4 > +</i> <i>Mhc-GAL4 > UAS-hop^{TumL}</i> <i>Mhc-GAL4 > UAS-hop^{TumL}, UAS-ImpL2-i</i>
D-F	<i>Cg-GAL4, tub-GAL80^{TS} > +</i> <i>Cg-GAL4, tub-GAL80^{TS} > UAS-hop^{TumL}</i> <i>Cg-GAL4, tub-GAL80^{TS} > UAS-hop^{TumL}, UAS-ImpL2-i</i>

Figure 7

A-D	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-upd3</i>
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Figure S2

A	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-upd3-i1</i>
B	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i> <i>upd3^Δ; esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i>
C-D	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}, UAS-upd3-i2</i>

Figure S3

A	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i>
B	<i>10XStat-GFP; esg-GAL4, tub-GAL80^{TS} > UAS-yki^{3SA}</i>
C-D	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}</i>

Figure S4

A	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i>
B-E	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}, UAS-upd3-i1</i> <i>Mhc-GAL4 > +</i> <i>Mhc-GAL4 > UAS-hop^{TumL}</i> <i>Mhc-GAL4 > UAS-hop^{TumL}, UAS-Stat92E-i</i> <i>dMef2-GAL4, tub-GAL80^{TS} > +</i>

	<i>dMef2-GAL4, tub-GAL80^{TS} > UAS-hop^{TumL}</i>
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Figure S5

A, B,	<i>Cg-GAL4, tub-GAL80^{TS} > +</i>
D	<i>Cg-GAL4, tub-GAL80^{TS} > UAS-hop^{TumL}</i>
C	<i>Lpp-GAL4, tub-GAL80^{TS} > +</i>
	<i>Lpp-GAL4, tub-GAL80^{TS} > UAS-hop^{TumL}</i>

Figure S6

<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}</i>
<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}, UAS-ImpL2-i</i>
<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}, UAS-upd3-i1</i>
<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-yki^{3SA}, UAS-hop^{TumL}, UAS-ImpL2-i, UAS-upd3-i1</i>

Figure S7

A-C	<i>esg-GFP; Myo1A-GAL4, tub-GAL80^{TS} > +</i> <i>esg-GFP; Myo1A-GAL4, tub-GAL80^{TS} > UAS-upd3</i>
D-F	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-upd3</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-upd3, UAS-Stat92E-i</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-upd3, UAS-dome-i</i>
G-H	<i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-upd3</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-Pvf1</i> <i>esg-GAL4, tub-GAL80^{TS} > UAS-GFP, UAS-upd3, UAS-Pvf1</i>

Supplemental Table S4: Oligonucleotide information. Related to STAR Methods.

Oligonucleotides
Upd3 forward: CTGGTCACTGATCTTACTCGCC
Upd3 reverse: GGATTGGTGGGATTGATGGGA
Pvf1 forward: CTGTCGCGTCCGCTGAG
Pvf1 reverse: CTCGCCGGACACATCGTAG
ImpL2 forward: AAGAGCCGTGGACCTGGTA
ImpL2 reverse: TTGGTGAACCTTGAGCCAGTCG
Socs36E forward: GAGATCCTCACAGAGGCCACT
Socs36E reverse: GCGAAACTTCCACCTGACC
4EBP forward: CTCCTGGAGGCACCAAACTTATC
4EBP reverse: TTCCCCTCAGCAAGCAACTG
InR forward: ACAAAATGTAAAACCTTGCACATCC
InR reverse: GCAGGAAGCCCTCGATGA
RpL32 forward: GCTAAGCTGTCGCACAAATG
RpL32 reverse: GTTCGATCCGTAACCGATGT
Stat92E-RB forward: TAGTCCAGTGTGGTGGATTGCCACCATGCCGCTAAATCCCTACAACATGAAC
Stat92E-RB reverse: ATCATGTCTGGATCCCTCGAGTCAAAAGTTCTCAAAGTTGTAATCGTATCGAAGTCC
hop ^{TumL} forward: TAGTCCAGTGTGGTGGATTTCGC CACCATGGCCCTGGCCAACGGG
hop ^{TumL} reverse: ATCATGTCTGGATCCCTCGA GTCACTCGGCATCCGTCGGCT
ImpL2-BS1 forward: TACCGGTGCTAGCCCCGGCTGAGG GAAACTCCAGCCGAAGTTCTAGAAC
ImpL2-BS1 reverse: CTTACTTAGATCGCAGATCTCGAG CGAAATGCAGATCGGATTAACCCA
ImpL2-BS2 forward: TACCGGTGCTAGCCCCGGGCTCGAGGAGCAGGTGTTTCCGCTAACAC
ImpL2-BS2- reverse: CTTACTTAGATCGCAGATCTCGAG GGCGCTATTAATGCGCTGTAATGC
ImpL2-BS3 forward: TACCGGTGCTAGCCCCGGGCTCGA GGGACGTGCTATCCGATTGCGATA
ImpL2-BS3- reverse: CTTACTTAGATCGCAGATCTCGAG CGTCGGCATTGTCATCGAAGTTT
ImpL2-BS4 forward: TACCGGTGCTAGCCCCGGGCTCGA GCACCGTCAACGGGCTTAAAGTTC
ImpL2-BS4- reverse: CTTACTTAGATCGCAGATCTCGA GATCGCTGGCGAATGTTGCATC
ImpL2-BS2-mut-#1 forward: TGCTAGCCCCGGGCTCGAG
ImpL2-BS2-mut-#1- reverse: CTCAAAGTAAAAACCTTGAATTAAAGTCTAG
ImpL2-BS2-mut-#2 forward: CTAGACTTTAATTCAAGGTTTCACTTGAG
ImpL2-BS2-mut-#2 reverse: CTGGAAAACCTCGAATGCCTCAC
ImpL2-BS2-mut-#3 forward: GTGAGGCATTGAGGTTTCCAG
ImpL2-BS2-mut-#3 reverse: AAGCTTACTTAGATCGCAGATCTCGAG
BS2-amplicon forward: TCGCTGAGCAATACTAAACACCTGC
BS2-amplicon reverse: GTGGAACCGAAAACGTGATACCC
BS3-amplicon forward: GGGCGAGAGCCGGAGTCACA
BS3-amplicon reverse: GCGCTGTAATGCGACGCTCTGG
BS4-amplicon forward: GGTGATTGCCGCCGAACGCAG
BS4-amplicon reverse: TGGCGAATGTTGCATCAGACGGT
Negative control- amplicon forward: CACGGCGGCCGGTGCATTCTC
Negative control- amplicon reverse: CAGCGCTTGCAGAGACCGGC