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## **Supplemental Information**

## **Tumor-Derived Ligands Trigger**

### **Tumor Growth and Host Wasting**

### via Differential MEK Activation

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#### **Supplemental Figure and Table Legends**

Supplemental Figure 1. Host wasting in yki<sup>3SA</sup>-tumor-bearing flies, related to Figure 1 and 2. (A-D) TAG (A), glucose (B), and trehalose levels (C) (n=3, 30 flies/group), and climbing speeds (D, n=60) in yki<sup>3SA</sup>-tumor-bearing flies. Thorax muscle degenerative phenotypes (E) protein amounts (F, n=3, 30 flies/group), muscle protein synthesis (G, left, n=3, 30 flies/group) and degradation rates (G, right, n=3, 30 flies/group), and of yki<sup>3SA</sup>-tumor-bearing flies at day 8. (H) TAG levels of indicated flies after 8 days at 29°C (n=3, 30 flies/group). (I) Lipolysis rates of fat bodies, isolated from indicated flies at 29°C at day 6, with or without Tram treatment (n=3, 30 abdomens/group). (J-K) TAG levels (J, n=3, 30 flies/group) and climbing speeds (K, n=60) of flies bearing yki<sup>3SA</sup>-gut tumors with dERK-RNAi (*esg-GAL4*, *UAS-GFP*, *tub-GAL80<sup>TS</sup>*/+; *UAS-yki<sup>3SA</sup>/UAS-dERK-RNAi*). (L-N) Appearance (L, up), midgut ISCs (L, down), TAG levels (M, n=3, 30 flies/group) and climbing speeds (N, n=60) of flies with dERK<sup>SEM</sup> overexpression in ISCs (*esg-GAL4*, *UAS-GFP*, *tub-GAL80<sup>TS</sup>*/*UAS-dERK<sup>SEM</sup>*). Data are presented as means ± SEM. \* p < 0.05.

Supplemental Figure 2. Pharmaceutical MEK inhibition in host tissues improves yki<sup>3SA</sup> tumor-induced systemic wasting, related to Figure 2. (A-B) Protocol used for feeding yki<sup>3SA</sup>-tumor-bearing flies with Tram after tumor induction (A) and schematic MEK suppression by Tram (B). (C-M) Phenotypes of systemic wasting in yki<sup>3SA</sup>-tumor-bearing flies (*esg-GAL4, UAS-GFP, tub-GAL80<sup>TS</sup>* /+; *UAS-yki<sup>3SA</sup>*/+) at day 8: bloating phenotypes (C, up), gut tumor phenotypes (C, middle), muscle degenerative phenotypes (C, down), MEK activation in the fat body and muscle (D), bloating rates (E,

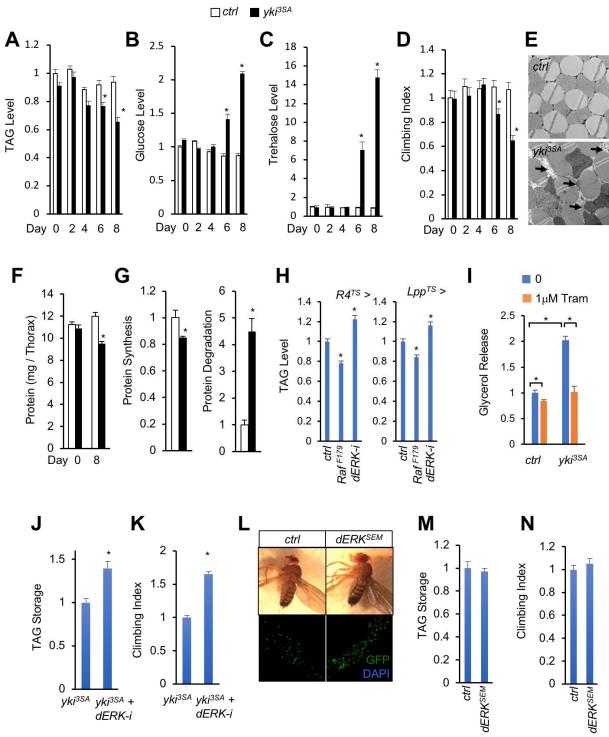
n=3, 60 flies/group), climbing speeds (**F**, n=60), lipid storages (**G**, n=3, 30 flies/group), trehalose and glucose levels in the hemolymph (**H**, **J**, n=3, 60 flies/group) and whole fly (**I**, **K**, n=3, 30 flies/group), and lifespan (**L** and **M**, n=120). (**N**) Midgut phenotypes of yki<sup>3SA</sup>-tumor-bearing flies, which were treated with Tram from day 4, at day 30. Data are presented as means  $\pm$  SEM. \* p < 0.05.

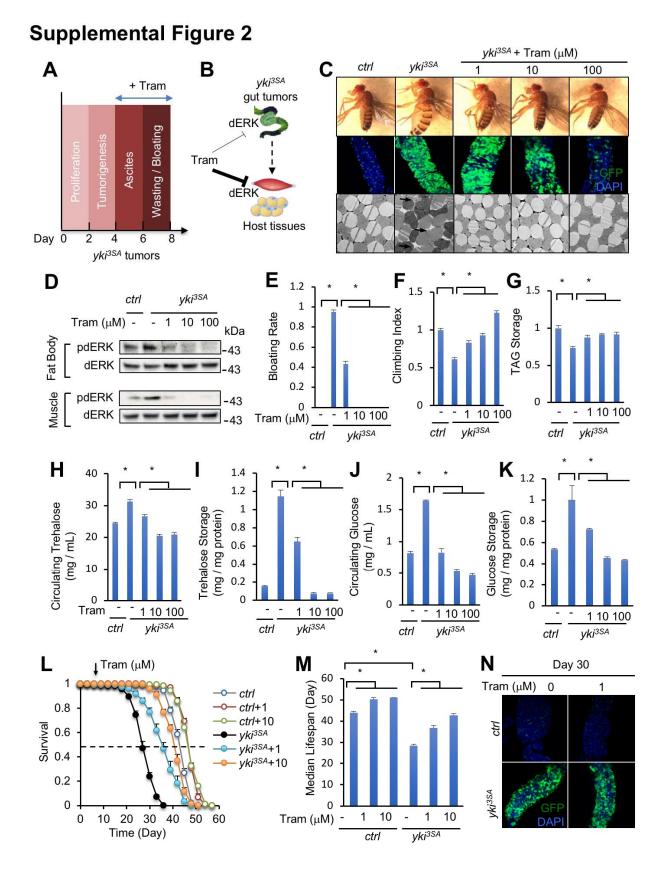
Supplemental Figure 3. Pharmaceutical MEK inhibition in host tissues only improves yki<sup>3SA</sup>+dERK<sup>SEM</sup> tumor-induced systemic wasting, related to Figure 2 and 3. (A-B) Appearances (A, up), midgut morphology (A, down), lipid and trehalose storage (**B**, n=3, 30 flies/group), and climbing speed (**B**, n=60) of control flies treated with Tram from day 0. (C-E) Bloating (C, up) and gut tumor (C, down), ERK activation in the fat body and muscles (**D**), TAG storage and trehalose levels (**E**, n=3, 30 flies/group), and climbing speed (E, n=60) of yki<sup>3SA</sup>+dERK<sup>SEM</sup>-tumor-bearing flies at day 8 that were treated with PD from simultaneously with tumor induction. (F) Heat map indicating differentially expressed genes that encode trypsins (up) and gut hormones (down) in *yki*<sup>3SA</sup> midguts. (G-I) Midgut phenotypes (Pros<sup>+</sup> cells) and appearances (G), wasting phenotypes including TAG and trehalose (H, n=3, 30 flies/group) levels and climbing speed (**H**, n=60), and survival rates (**I**, n=120) of indicated flies. Genotype for  $N^{intra}$  is esq-GAL4, UAS-GFP, tub-GAL80<sup>TS</sup>/UAS-N<sup>intra</sup>. (J) Midgut Pvf1 mRNA levels of flies bearing yki<sup>3SA</sup> or yki<sup>3SA</sup>+dERK<sup>SEM</sup> tumors at day 8. (K-M) Midgut (K, left, n=3, 30 guts/group) and head (K, right, n=3, 30 heads/group) mRNA levels, Ilp2 levels in IPCs (L), and food intake (M, n=4, 80 flies/group) of yki<sup>3SA</sup>+dERK<sup>SEM</sup> tumor-bearing flies (*esg*-GAL4, UAS-GFP, tub-GAL80<sup>TS</sup>/UAS-dERK<sup>SEM</sup>; UAS-vki<sup>3SA</sup>/+) at day 8 that were fed 10

 $\mu$ M Tram simultaneously with tumor induction. (**N**) Midgut *ImpL2* mRNA levels in flies bearing yki<sup>3SA</sup> tumors with RNAi at day 10 (n=3, 30 guts/group). Data are presented as means ± SEM. \* p < 0.05.

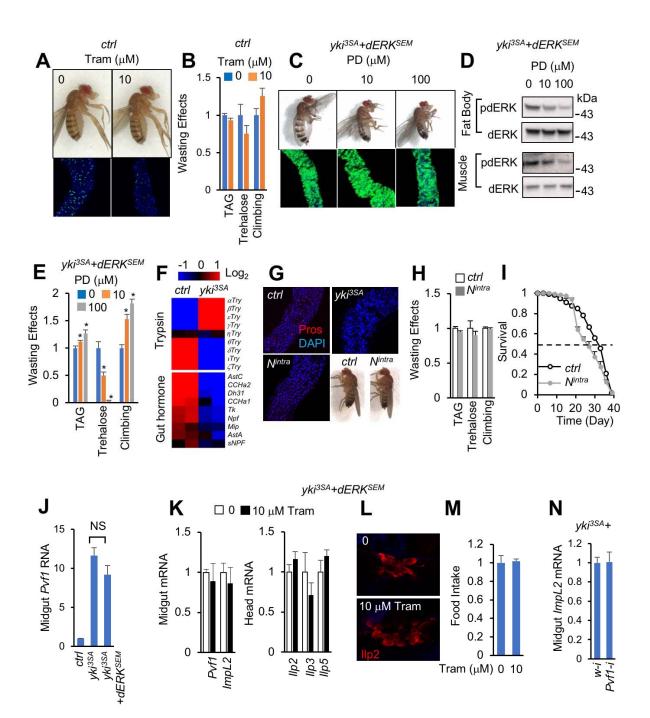
Supplemental Figure 4. MEK inhibition alleviates mammalian myotube atrophy and lipolysis, related to Figure 2 and 3. (A) MEK/ERK activation in differentiated C2C12 myotubes (day 5) treated with LLC conditioned differentiation medium (3:1 diluted with fresh differentiation medium) at different time points (up) or with conditioned differentiation medium plus Tram for 1h (down). (**B-E**) Gene expression (**B**, n=3), protein degradation rates (**C**, n=3), myotube diameters (**D**, n=30, quantification of myotube images), and myotube morphologies (E) in C2C12 myotubes (day 5) treated with LLC conditioned differentiation medium with or without 0.1 µM trametinb for 24h. (F) MEK/ERK activation in differentiated 3T3-L1 adipocytes (day 10) treated with LLC conditioned growth medium (3:1 diluted with fresh growth medium) at different time points (up) or with conditioned growth medium plus Tram for 1h (down). (G) Lipolysis rates in 3T3-L1 adipocytes (day 10) treated with control or LLC-conditioned growth medium for 24h and then treated with 0.1  $\mu$ M Tram for 1h (n=3). (H-I) Lipid storages (H, n=3) and lipid staining (I) in 3T3-L1 adipocytes (day 10) treated with LLC-conditioned growth medium with or without 0.1  $\mu$ M Tram for 24h. Data are presented as means ± SEM. \* p < 0.05.

# **Supplemental Figure 1**





# **Supplemental Figure 3**



# **Supplemental Figure 4**

