

Misshapen Disruption Cooperates with *Ras*^{V12} to Drive Tumorigenesis

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Genotypes of all the figures

Supplemental figure1

Supplemental figure 2

Supplemental figure 3

Supplemental figure 4

Supplemental figure 5

Genotypes of all the figures

Figure 1

(B) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/+; tub-Gal80, FRT79E/FRT79E*; (C) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/+; tub-Gal80, FRT79E/msn³²⁰⁸, FRT79E*; (D and K) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT79E/ FRT79E*; (E) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT80B/msn¹⁷², FRT80B*; (F, L-M) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT79E/msn³²⁰⁸, FRT79E*; (G) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT79E/msn³²⁰⁸, UAS-Msn, FRT79E*; (H, N) *ey-Flp1/UAS-bsk^{DN}; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT79E/msn³²⁰⁸, FRT79E*; (O) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/+; tub-Gal80, FRT79E, puc^{E69}/msn³²⁰⁸, FRT79E*; (P) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT79E, puc^{E69}/msn³²⁰⁸, FRT79E*; (Q) *ey-Flp1/UAS-bsk^{DN}; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT79E, puc^{E69}/msn³²⁰⁸, FRT79E*.

Figure 2

(A, C and G) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT79E/FRT79E*; (B, D and H) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT79E/msn³²⁰⁸, FRT79E*; (E) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/ex-lacZ, UAS-Ras^{V12}; tub-Gal80, FRT79E/FRT79E*; (F) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/ex-lacZ, UAS-Ras^{V12}; tub-Gal80, FRT79E/msn³²⁰⁸, FRT79E*; (I and N) *ey-Flp5, Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT79E/msn³²⁰⁸, FRT79E*; (J and P) *ey-Flp5, Act>y+>Gal4, UAS-GFP/UAS-Wts, UAS-Ras^{V12}; tub-Gal80, FRT79E/msn³²⁰⁸, FRT79E*; (K and O) *UAS-bsk^{DN}/+; ey-Flp5, Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; tub-Gal80, FRT79E/msn³²⁰⁸, FRT79E*; (L and Q) *UAS-bsk^{DN}/+; ey-Flp5, Act>y+>Gal4, UAS-GFP/UAS-Wts, UAS-Ras^{V12}; tub-Gal80, FRT79E/msn³²⁰⁸, FRT79E*; (M) *ey-Flp5, Act>y+>Gal4, UAS-GFP/+; tub-Gal80, FRT79E/FRT79E*.

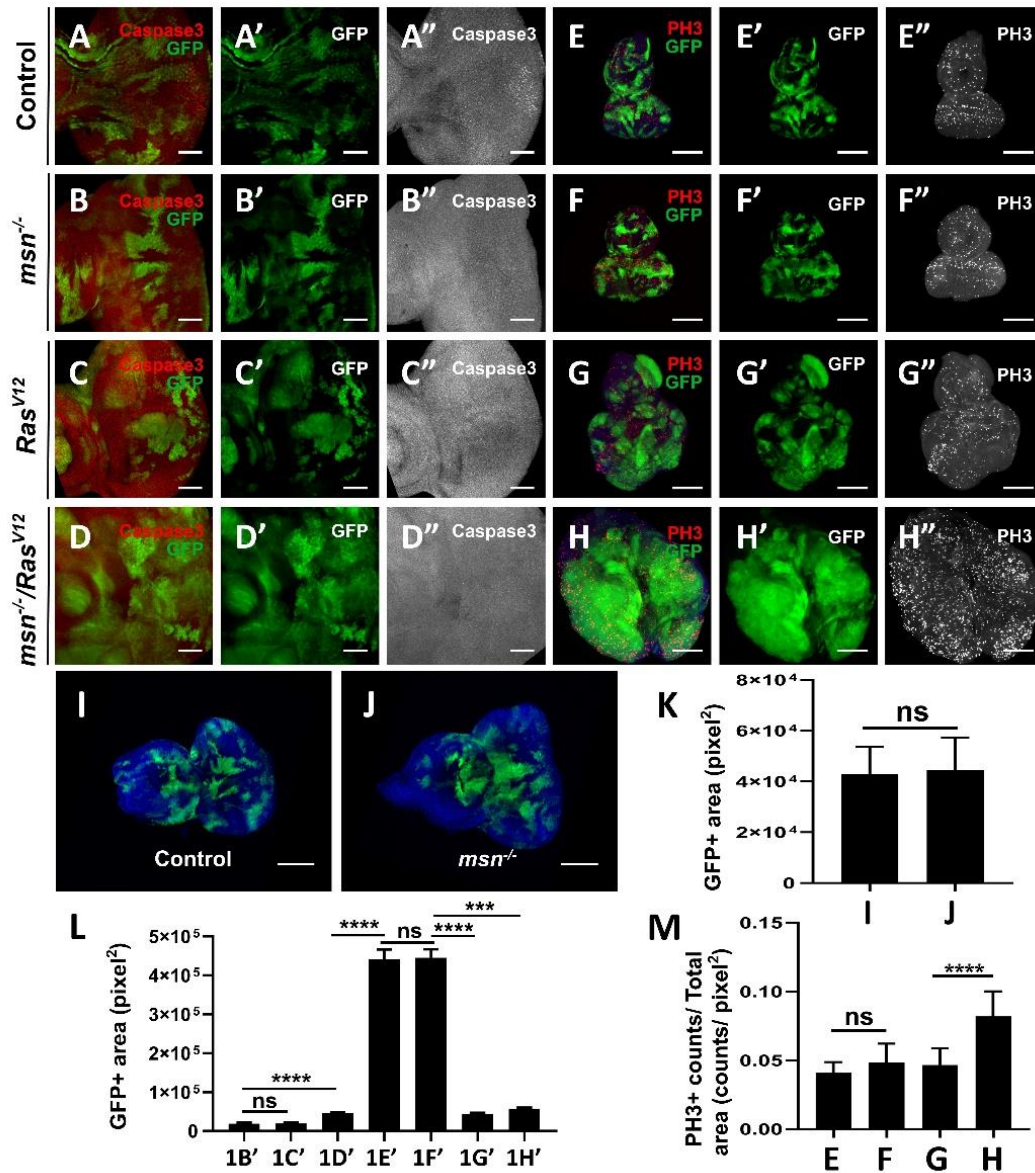
Figure 3

(A, C and G) *ey-Flp5, Act>y+>Gal4, UAS-GFP/+; FRT 82B, tub-Gal80/FRT82B*; (B, D and H) *ey-Flp5, Act>y+>Gal4, UAS-GFP/+; FRT 82B, tub-Gal80/FRT82B, UAS-Msn*; (E) *ey-Flp5, Act>y+>Gal4, UAS-GFP/+; diap1-lacZ, FRT 82B, tub-Gal80/FRT82B*; (F) *ey-Flp5, Act>y+>Gal4, UAS-GFP/+; diap1-lacZ, FRT 82B, tub-Gal80/FRT82B, UAS-Msn*; (I) *ey-Flp5, Act>y+>Gal4, UAS-GFP/+; FRT 82B, tub-Gal80/FRT82B, scrib¹*; (J) *ey-Flp5, Act>y+>Gal4, UAS-GFP/+; FRT 82B, tub-Gal80/FRT82B, UAS-Msn, scrib¹*; (K) *UAS-bsk^{DN}/+; ey-Flp5, Act>y+>Gal4, UAS-GFP/+; FRT 82B, tub-Gal80/FRT82B, scrib¹*; (L) *UAS-bsk^{DN}/+; ey-Flp5, Act>y+>Gal4, UAS-GFP/+; FRT 82B, tub-Gal80/FRT82B, UAS-Msn, scrib¹*; (N) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; FRT 82B, tub-Gal80/FRT82B, scrib¹*; (O) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}; FRT 82B,*

tub-Gal80/ FRT82B, UAS-Msn, scrib¹; (P) *ey-Flp5, Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}*; FRT82B, *tub-Gal80/FRT82B*; (Q) *ey-Flp5, Act>y+>Gal4, UAS-GFP/UAS-Ras^{V12}*; FRT82B, *tub-Gal80/FRT82B, UAS-Msn*.

Figure 4

(A and I) *nub-Gal4/+*; (B) *nub-Gal4/+; UAS-Msn/+*; (C) *nub-Gal4/UAS-ft-IR*; (D) *nub-Gal4/UAS-ft-IR; UAS-Msn/+*; (F) *ex-lacZ/+; hh-Gal4/UAS-Msn*; (G) *ex-lacZ/UAS-ft-IR; hh-Gal4/+*; (H) *ex-lacZ/UAS-ft-IR; hh-Gal4/UAS-Msn*; (J) *nub-Gal4/UAS-Ft^{ΔECD}*; (K) *nub-Gal4/UAS-Ft^{ΔECD}; wts^{X1}/+*; (L) *nub-Gal4/UAS-Ft^{ΔECD}; msn³²⁰⁸/+*; (N) *en-Gal4, UAS-GFP/+; msn-lacZ/+*; (O) *en-Gal4, UAS-GFP/UAS-yki-IR; msn-lacZ/+*; (P) *en-Gal4, UAS-GFP/UAS-sd-IR; msn-lacZ/+*; (Q) *en-Gal4, UAS-GFP/+; msn-lacZ/UAS-Yki*; (R) *en-Gal4, UAS-GFP/+; msn-lacZ/UAS-Yki^{S168A}*; (S) *en-Gal4, UAS-GFP/UAS-sd-IR; msn-lacZ/UAS-Yki^{S168A}*.



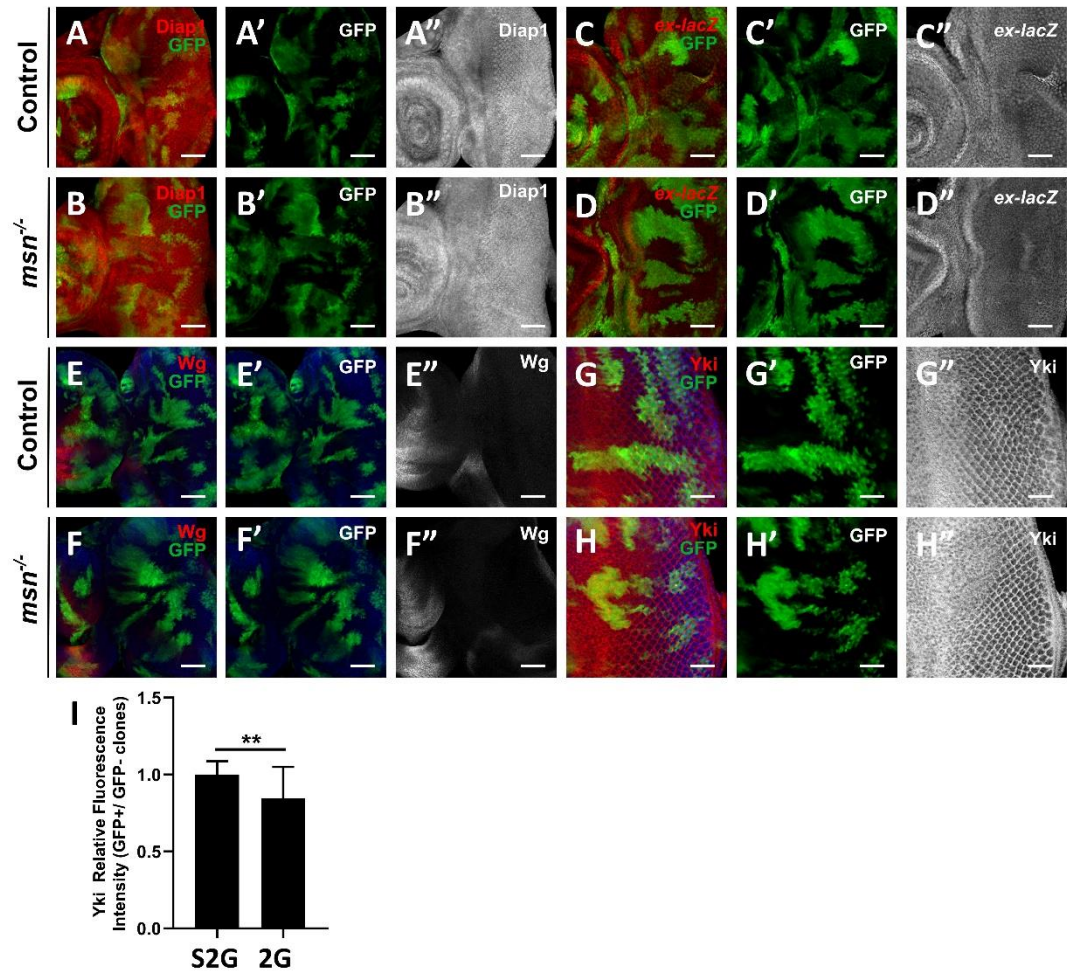


Figure S2. Loss of *msn* alone does not affect Hippo signaling activation

(A-H) Fluorescence micrographs of GFP-labeled clones of eye discs are shown. Compared with control, loss of *msn* alone has no obvious effect on Hippo target genes, including Diap1, Wg, and *ex*. Scale bars represent 50 μ m (A-F), 20 μ m (G-H). (I) Quantification of Yki relative fluorescence intensity in Figures S2G and 2G. ** $P < 0.01$ (mean + S.D.). Genotypes are as follows: (A, E and G) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/+; tub-Gal80, FRT79E/FRT79E*; (B, F and H) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/+; tub-Gal80, FRT79E/msn³²⁰⁸, FRT79E*. (C) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/ex-lacZ; tub-Gal80, FRT79E/ FRT79E*; (D) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/ex-lacZ; tub-Gal80, FRT79E/ msn³²⁰⁸, FRT79E*.

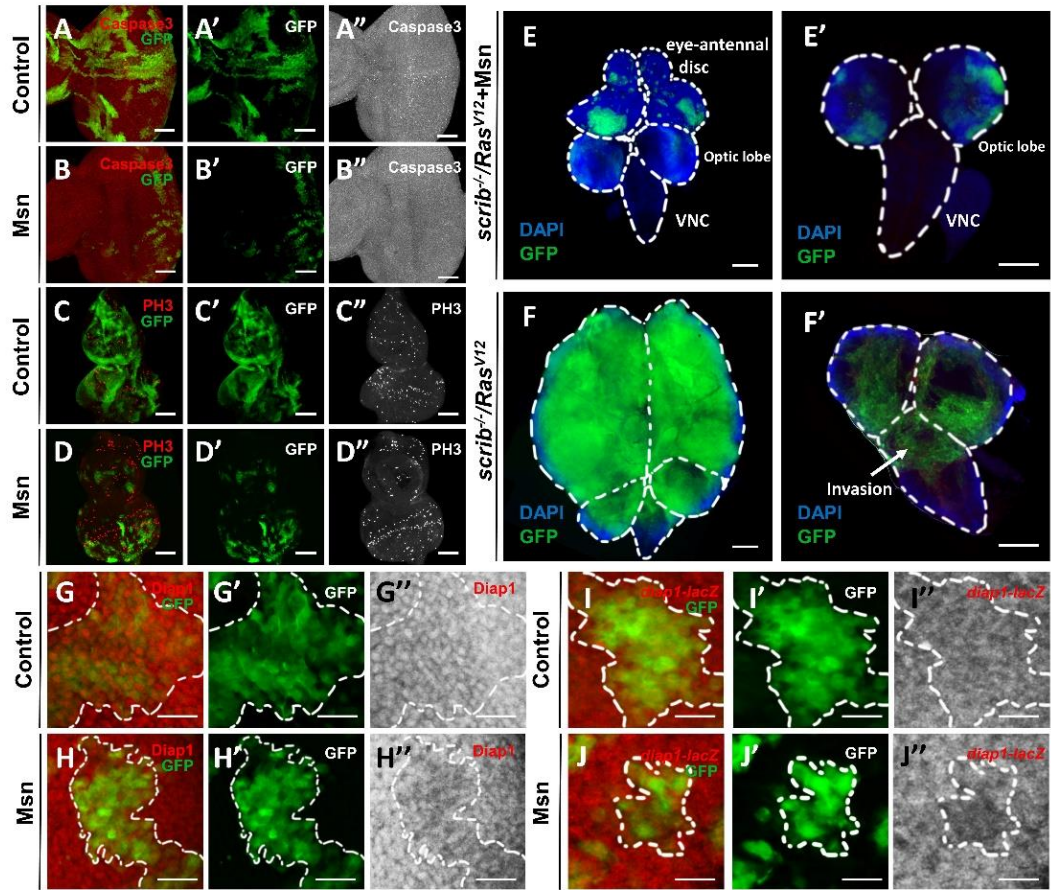


Figure S3. Msn overexpression suppresses tumorigenesis.

(A-D) Fluorescence micrographs of eye discs are shown. Compared with wild-type (A and C), Msn overexpression alone does not induce significant apoptosis (B) or proliferation (D), as indicated by Caspase 3 and PH3 staining, respectively. (E-F) Msn overexpression suppressed *scrib*^{-/-}/*Ras*^{V12} induced tumor overgrowth (E-F) and VNC invasion behavior (E'-F'). (G-J) Zoom in images of Figure 3C-E. Scale bars represent 50 μ m (A-B), 100 μ m (C-F), 20 μ m (G-J). Genotypes are as follows: (A, C, G and I) *ey-Flp5, Act>y+>Gal4, UAS-GFP/+; FRT82B, tub-Gal80/ FRT82B*; (B, D, H and J) *ey-Flp5, Act>y+>Gal4, UAS-GFP/+; FRT82B, tub-Gal80/ FRT82B, UAS-Msn*; (E) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/ UAS-Ras^{V12}; FRT82B, tub-Gal80/ FRT82B, scrib^l, UAS-Msn*; (F) *ey-Flp1/+; Act>y+>Gal4, UAS-GFP/ UAS-Ras^{V12}; FRT82B, tub-Gal80/ FRT82B, scrib^l*.

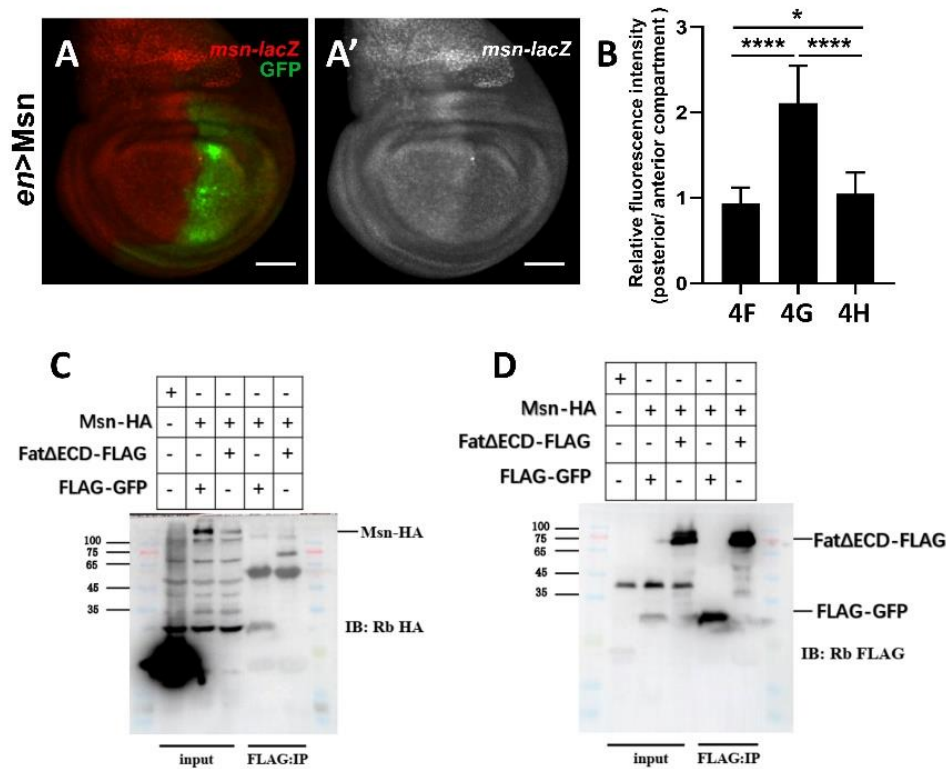


Figure S4. Msn regulates Hippo signaling in a feedback manner

(A-A') Fluorescence micrographs of wing discs are shown. Msn overexpression suppresses *msn* itself transcription. Scale bars represent 100 μ m (A-A'). Genotypes are as follows: (A-A') *en-Gal4*, *UAS-GFP*/+; *msn-lacZ*/*UAS-Msn*. (B) Quantification of Relative fluorescence intensity in Figure 4F-H. * $P < 0.05$ (mean + S.D.), **** $P < 0.0001$ (mean + S.D.). (C-D) Immunoprecipitation experiment indicated that no physical interaction between Ft Δ ECD and Msn in S2 cells. Lysates expressing the indicated constructs were immunoprecipitated (IP) and probed with the indicated antibodies. Msn-HA was not detected in Ft Δ ECD-FLAG immunoprecipitation (C). Conversely, Ft Δ ECD-FLAG was detected in FLAG immunoprecipitation indicated Ft Δ ECD-FLAG plasmid has been transfected into S2 cells (D).

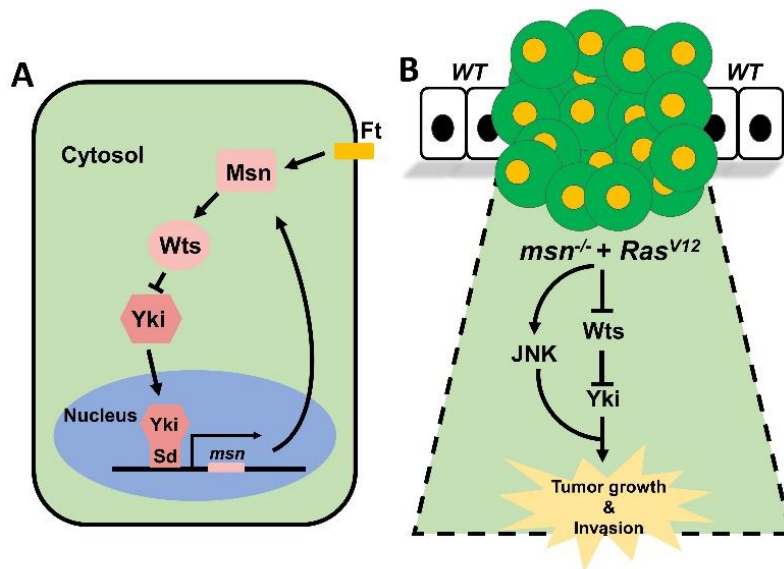


Figure S5. A schematic model depicting the role of Msn in regulating Hippo signaling and tumorigenesis.

(A) Msn acts as downstream of Ft to regulate Hippo pathway in a negative feedback manner. (B) Msn disruption cooperates with *Ras*^{V12} to drive tumorigenesis by inducing JNK pathway activation and Hippo pathway inactivation.