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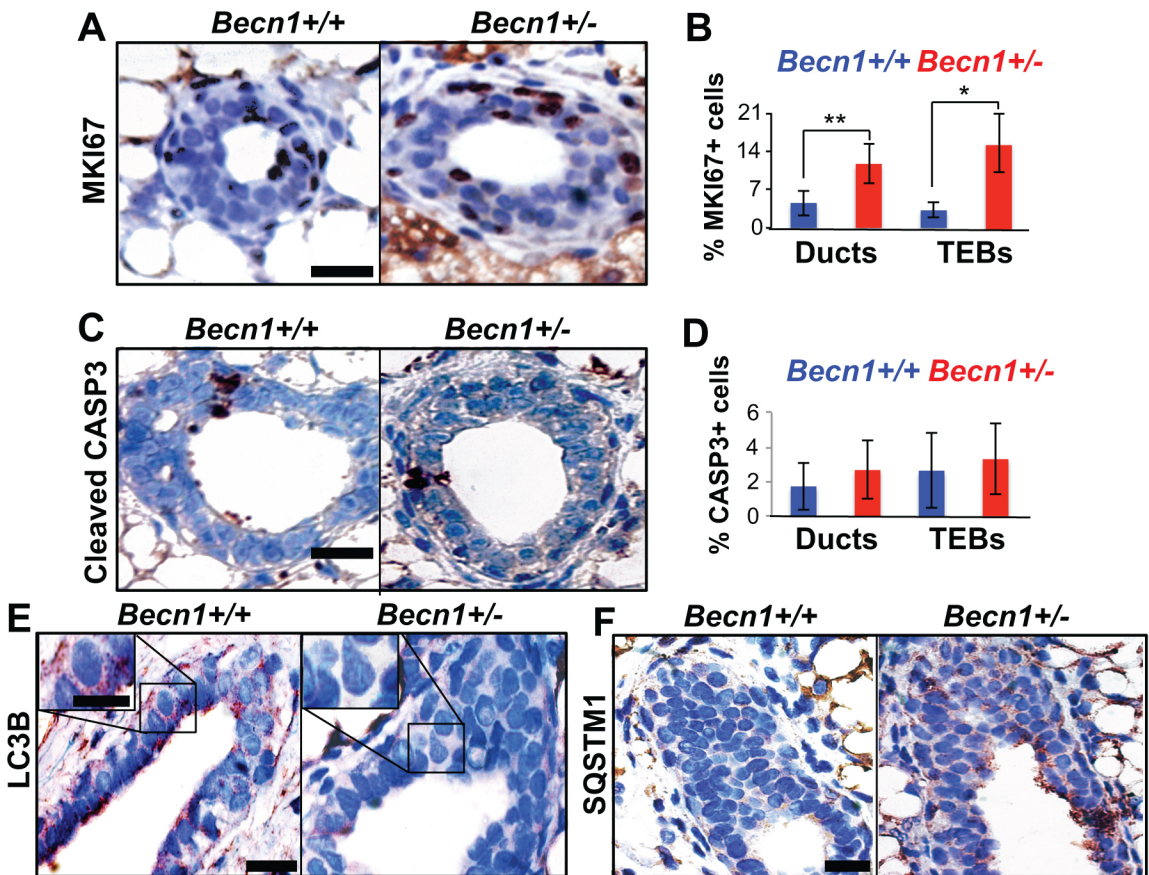
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Sandy Price, Ritu Nahar, Fred Lozy, Hua Zhong,
Alexei Vazquez, Yibin Kang, and Vassiliki Karantza**

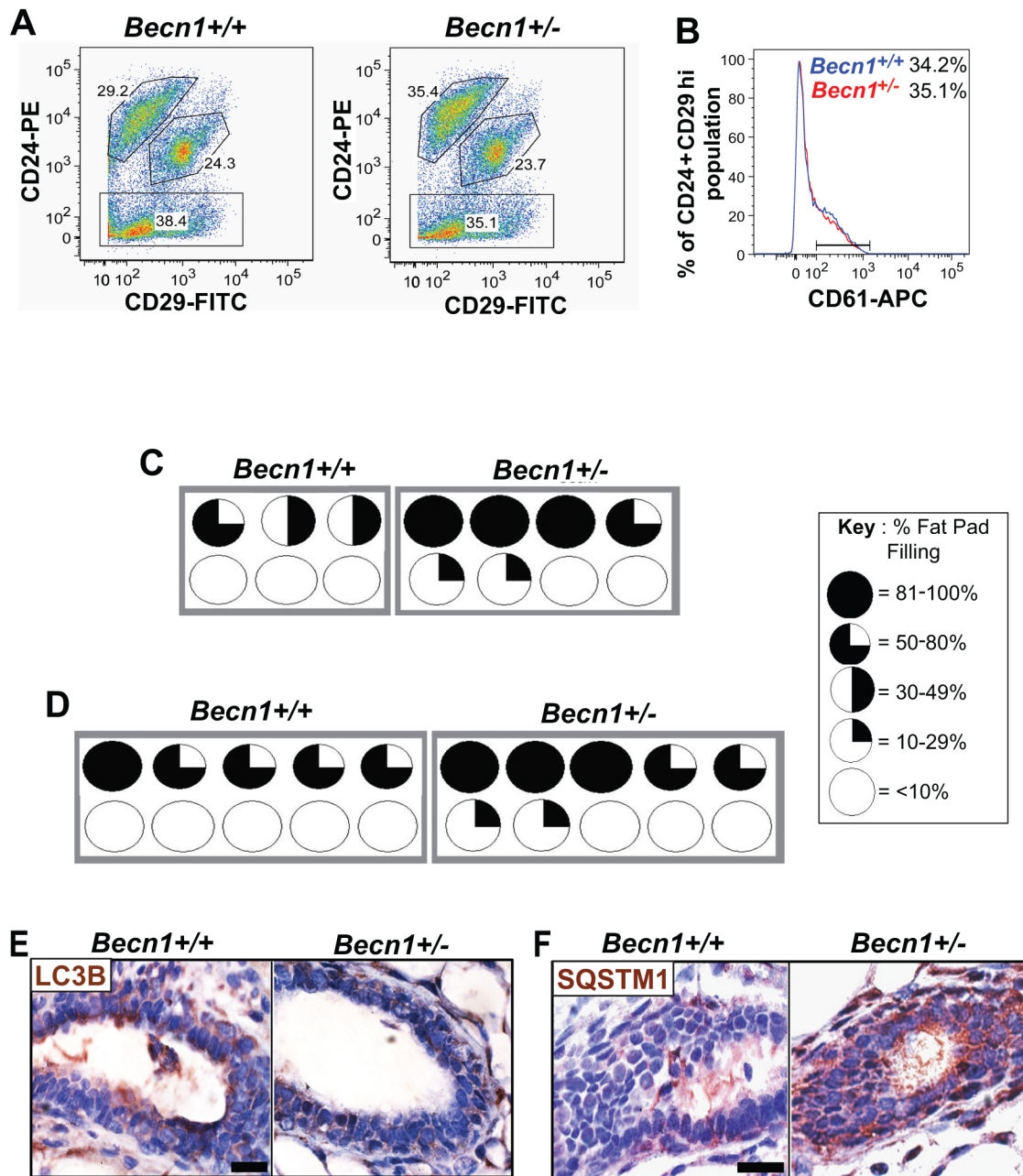
**Autophagy regulator BECN1 suppresses mammary
tumorigenesis driven by WNT1 activation
and following parity**

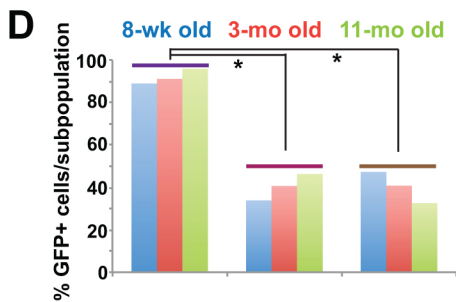
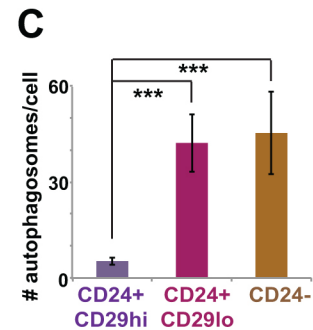
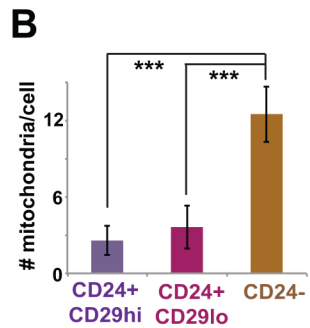
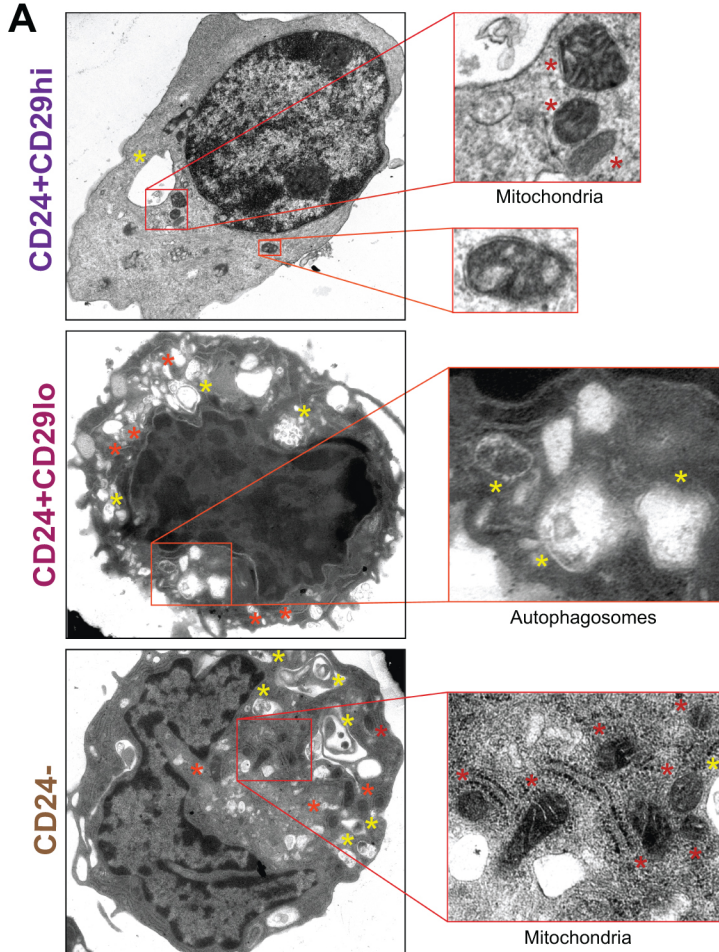
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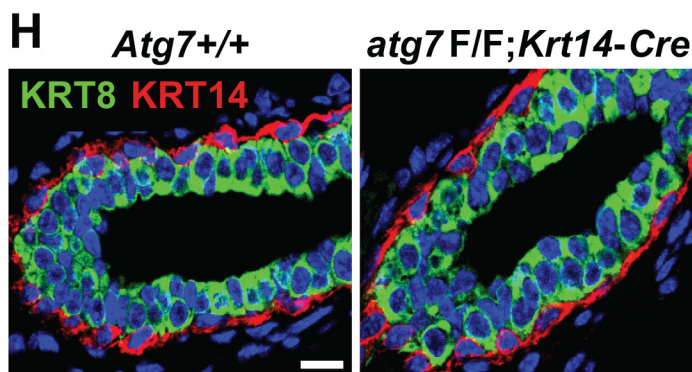
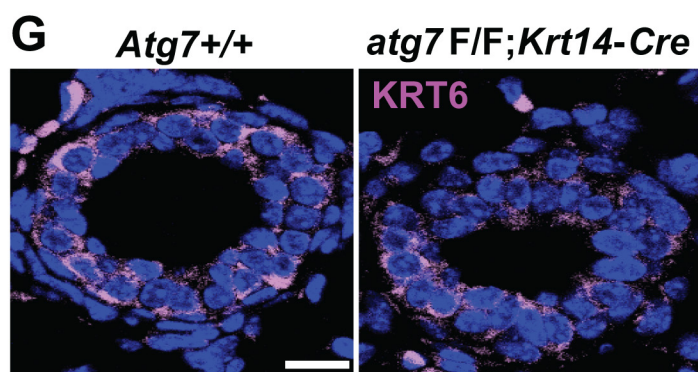
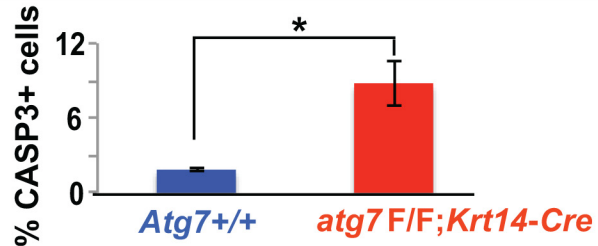
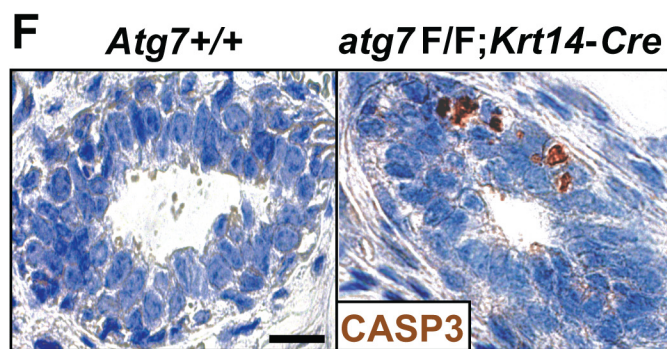
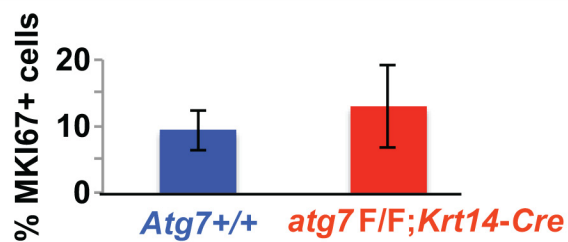
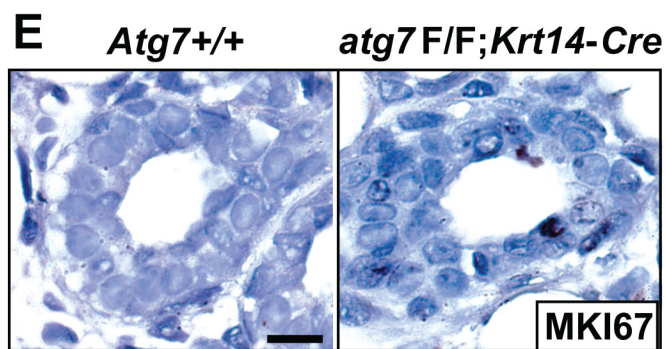
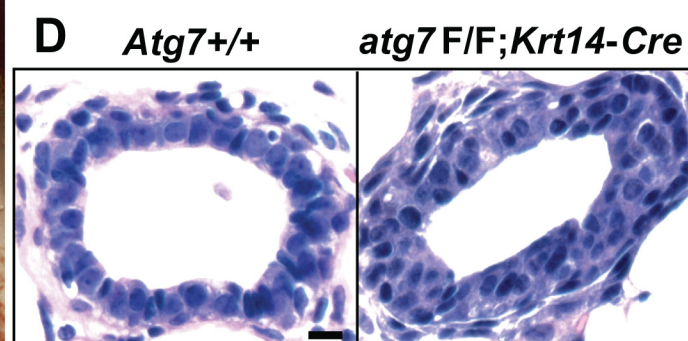
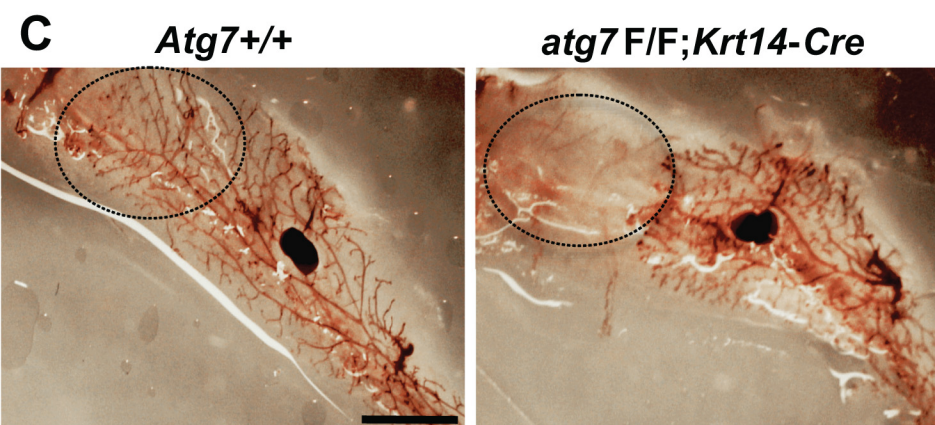
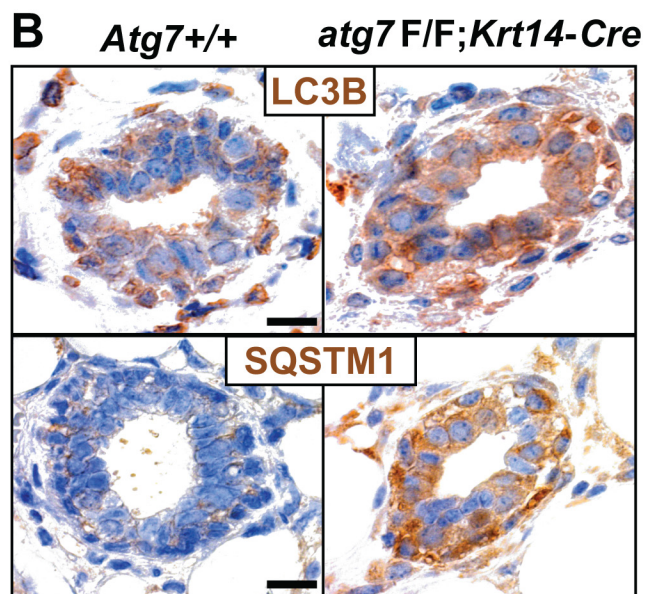
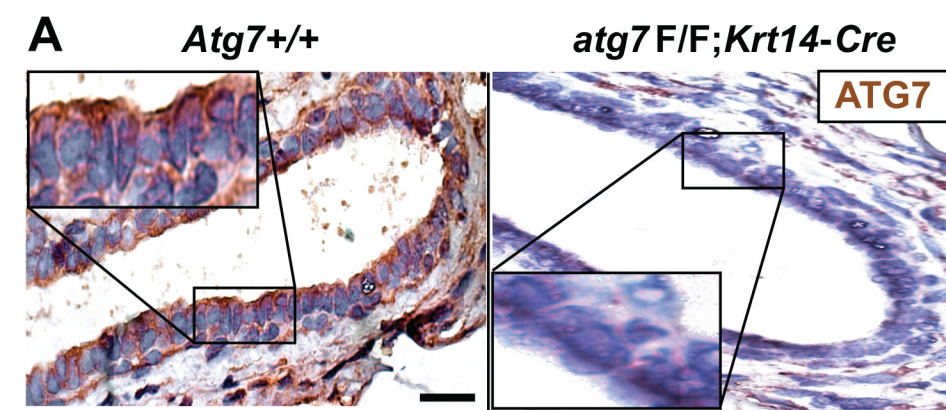
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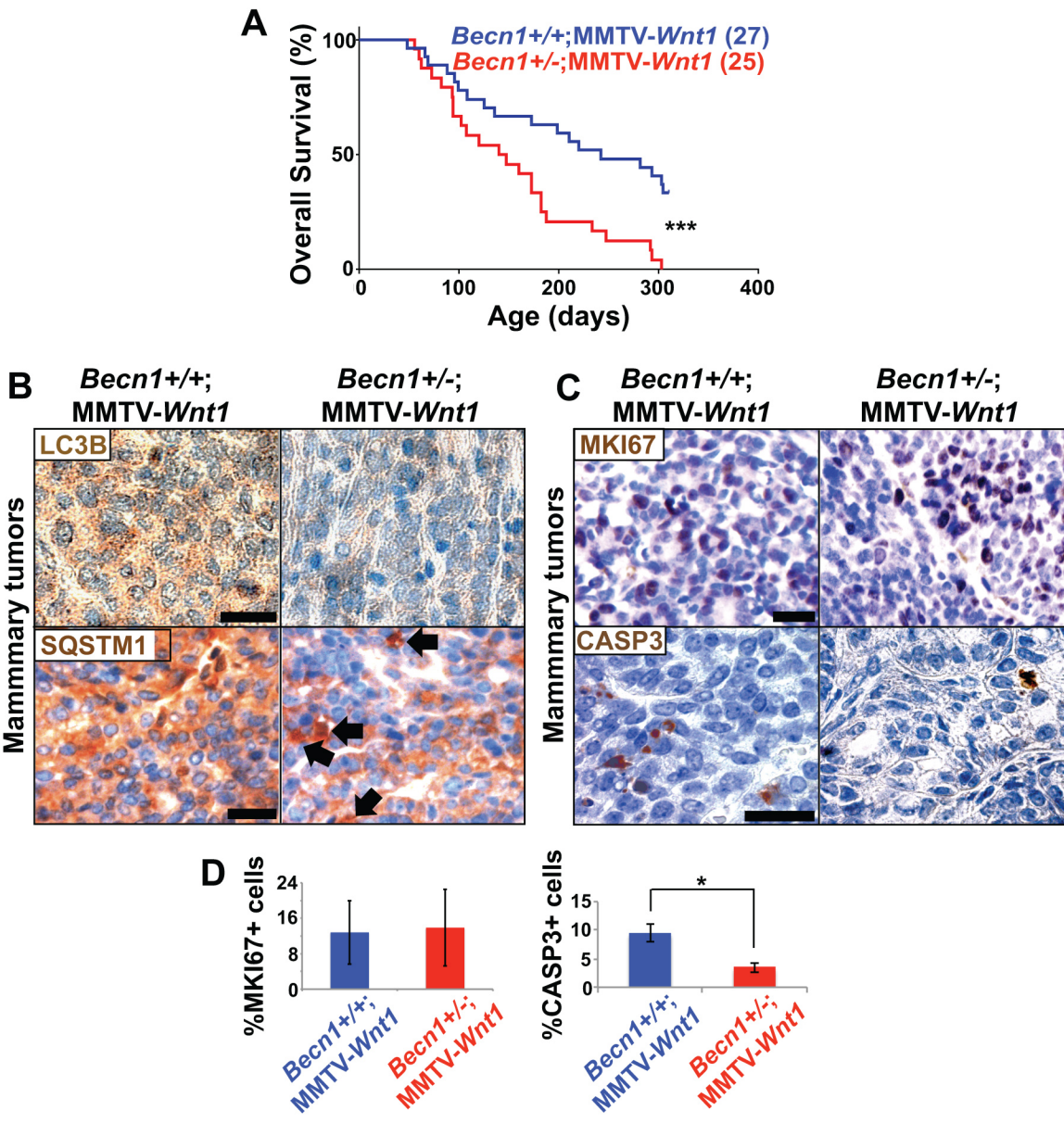
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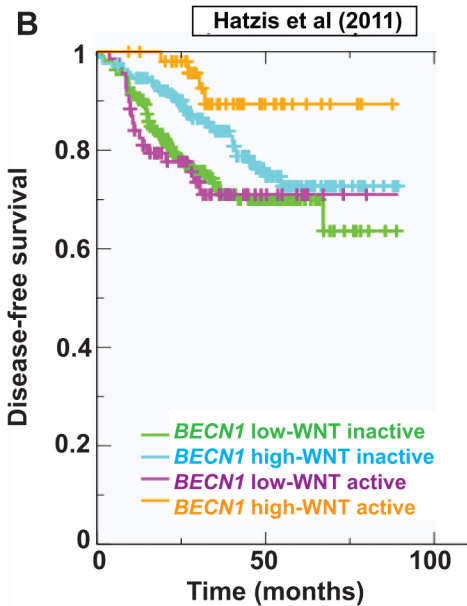
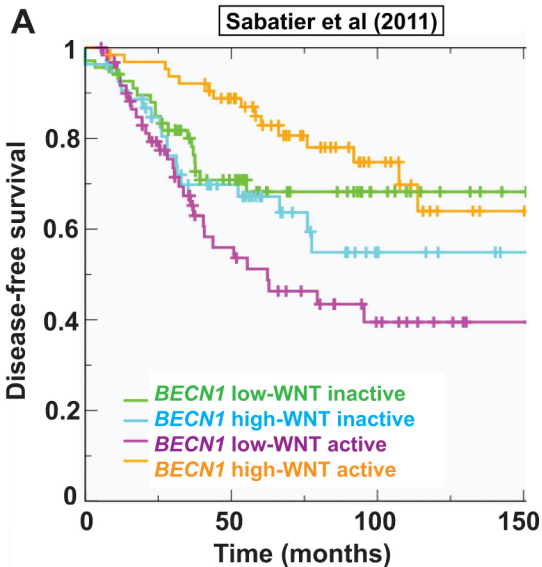


Figure S1. MGs of *Becn1*^{+/-} mice display increased proliferation and defective autophagy status. **(A, C, E and F)** Immunohistochemical analysis of MGs from cohoused *Becn1*^{+/+} (left) and *Becn1*^{+/-} (right) 5-wk old littermates. **(A)** Representative images and **(B)** quantification of MKI67⁺ cells in mammary ducts (left) and TEBs (right) shown as means \pm SDs (n=3 mammary gland specimens per genotype). **(C)** Representative images and **(D)** quantification of cleaved CASP3 in each sample in TEBs and ducts shown as means \pm SDs (n=3 mammary gland specimens per genotype). **(E)** Representative image of LC3B staining confirms decreased LC3B puncta in MGs from *Becn1*^{+/-} mice. **(F)** Representative image of SQSTM1 staining confirms accumulation of SQSTM1 in MGs from *Becn1*^{+/-} mice. **P* <0.05 and ***P* <0.01 by a two-tailed Student t test. Scale bar = 30 μ m for **(A,C,E,F)**.

Figure S2. Outgrowths from CD24⁺CD29^{hi} *Becn1*^{+/-} MEC transplantation exhibit increased MaSC activity and retain defective autophagy status. **(A and B)** MEC sorting experiments were performed 5 independent times. **(A)** FACS was performed on freshly isolated MECs stained with PI, LIN (CD31, CD45, and LY76), CD24, and CD29. Images are representative of *Becn1*^{+/+} (left) and *Becn1*^{+/-} (right) MEC distribution with gating for PI⁻LIN⁻ cells and based on CD24 and CD29 expression. No significant differences were seen in relative MEC subpopulations. **(B)** A representative histogram shows similar numbers of CD61⁺ cells between *Becn1*^{+/+} and *Becn1*^{+/-} CD24⁺CD29^{hi} MEC populations. **(C and D)** Circles represent fat pads and the black color represents the percentage of the

mammary fat pad that is filled in. **(C)** Increased repopulation frequency and mammary fat pad filling are seen following transplantation of 500 CD24⁺CD29^{hi} MECs isolated from 8 wk old *Becn1*^{+/-} mice, contralateral CD24⁺CD29^{hi} *Becn1*^{+/+} and *Becn1*^{+/-} MEC transplantations were performed. **(D)** Increased repopulation frequency and mammary fat pad filling are seen following transplantation of 500 CD24⁺CD29^{hi} MECs isolated from 9.5-wk old *Becn1*^{+/-} mice, contralateral CD24⁺CD29^{hi} *Becn1*^{+/+} and *Becn1*^{+/-} MEC transplantations were performed. **(E)** Representative image of LC3B staining in samples confirms decreased LC3B puncta in outgrowths from CD24⁺CD29^{hi} *Becn1*^{+/-} MEC transplantation. **(F)** Representative image of SQSTM1 staining confirms accumulation of SQSTM1 in outgrowths from CD24⁺CD29^{hi} *Becn1*^{+/-} MEC transplantation. Scale bar= 30 μ m for **(E,F)**.

Figure S3. FACS for CD24 and CD29 expression defines MEC populations with different autophagy levels. PILLIN⁻ MECs were isolated by FACS based on CD24 and CD29 expression, and examined by EM. \blacktriangle and * denote mitochondria and autophagosomes, respectively.

Figure S4. *Krt14*-driven conditional biallelic *Atg7* deletion confers autophagy deficiency in the mammary gland. **(A to H)** Evaluation of MGs from *Atg7*^{+/+} (left) and *atg7*^{F/F}; *Krt14-Cre* (right) mice. **(A)** Representative images of ATG7 staining confirms *Atg7* deletion in MGs from *atg7*^{F/F}; *Krt14-Cre* mice. **(B)** Representative images of SQSTM1 (top) and LC3B (bottom) staining. **(C)** Representative MG

whole mounts (WM) from 6.5-wk old mice reveal decreased mammary fat pad filling in MGs from *atg7^{F/F};Krt14-Cre* mice. **(D)** Representative images of hematoxylin and eosin (H&E) staining reveal similar ductal organization in MGs from *atg7^{F/F};Krt14-Cre* and *Atg7^{+/+}* mice. **(E)** Representative images and quantification of MKI67 staining. **(F)** Representative images and quantification of cleaved CASP3 staining. Representative images of **(G)** KRT6 (purple), and **(H)** KRT8 (green) and KRT14 (red) staining. Scale bar = 20 μ m for **(A,B,D to H)** and 5 mm for **(C)**.

Figure S5. Mammary tumors arising in *Becn1^{+/-};MMTV-Wnt1* mice exhibit autophagy-deficient status and decreased apoptosis. **(A)** Kaplan-Meier curves for overall survival in *Becn1^{+/+};MMTV-Wnt1* (n=27) and *Becn1^{+/-};MMTV-Wnt1* (n=25) mice demonstrates decreased overall survival in *Becn1^{+/-};MMTV-Wnt1* mice. **(B and C)** Analysis of mammary tumors arising in *Becn1^{+/+};MMTV-Wnt1* (left) and *Becn1^{+/-};MMTV-Wnt1* (right) mice. **(B)** Representative image of LC3B staining (top) and SQSTM1 aggregates annotated with arrow (bottom) in tumors from *Becn1^{+/-};MMTV-Wnt1* mice. **(C)** Representative image from MKI67 staining (top) and cleaved CASP3 staining (bottom). **(D)** Quantification of MKI67⁺ cells (left) and cleaved CASP3⁺ cells (right) determine proliferation and apoptosis rates, respectively, in tumors. Results are presented as means \pm SDs (n=3 mammary tumors per genotype) **P* < 0.05 determined by a two-tailed Student t test. Scale bar= 40 μ m for **(B,C)**.

Figure S6. Kaplan-Meier curves for breast tumors stratified for *BECN1* expression and WNT pathway activation. **(A)** Disease-free survival for patients with breast cancers annotated in the Sabatier cohort. **(B)** Disease-free survival for patients with ERBB2-negative breast cancers annotated in the Hatzis cohort.