## SUPPLEMENTARY DATA

Supplemental Experimental Procedures:

Primers used (all are against mouse, unless otherwise indicated):

qPCR primers:

hes1 NM 008235.2 (5'-CACAGAAAGTCATCAAAGCC and 5'-TGCTTGACAGTCATTTCCAG), notch1 NM 008714.3 (5'-CTACAGAAGGTTACACAG and 5'-CAGAGGTAGGAGTTGTCACG), p53 NM 001127233.1 (5'-TGAACCGCCGACCTATCCTTA and GGCACAAACACGAACCTCAAA), twist1 NM 011658.2 (5'-AATTCACAAGAATCAGGGCGTGGG and 5'-TCTATCAGAATGCAGAGGTGTGGG), snail NM 011427.2 (5'-TCCAAACCCACTCGGATGTGAAGA and 5'-TTGGTGCTTGTGGAGCAAGGACAT), snai2 NM 011415.2 (5'-CACATTCGAACCCACACATTGGCT and 5'-TGTGCCCTCAGGTTTGATCTGTCT), zeb1 NM 011546.3 (5'-CAGTGTTCCATGTTTAAGAGCA and 5'-GTCTTTCATCCTGGTTTCCG), klf4 NM 010637.3 (5'-CATTATCAAGAGTCTATGCCA and 5'-CACAGTGGTAAGGTTTCTCG), cdh1 NM 009864.2 (5'-GCCAAGTACATCCTCTATTCTC and 5'-GCAACGAATCCCTCAAAGAC), gapdh NM 008084.2 (5'-CGTGGAGTCTACTGGCGTCTTCAC and 5'-CGGGGGATGATGAGCCTTTTGGC), brcal NM 009764.3 (5'-CCAAAGAAGTAATGACCGTG and 5'-GCTAACTATCCACTTTCCTCC), gata-3NM\_008091.3 (5'-ACGAATCCAGCACAGAAGG and 5'-ATGTCCCTGCTCTCCTTG), ck14 NM 016958.1 (5'-TCTTCAGCAAGACAGAGGAG and 5'-CTCCAGGTTATTCTCCAGGG), p63 NM 001127259.1 (5'-GTTCAATGAGGGACAGATTGC and 5'-GAATTCAGTGCCAACCTGTG), Human brca1 NM 007294.3 (5'-CCCAGAAGAATTTATGCTCGT and 5'-CATTGACCACATCTCCTCTG), Human gapdhBC096440.1 (5'-AAGGTGAAGGTCGGAGTCAA and 5'-AATGAAGGGGTCATTGATGG). ChIP primers:

*p21* NC\_000006.11 (5'-GATTTCCTTTCTATCAGCCC and GTCACAAGATACATACCACCT), *gapdh* NC\_000012.11 (5'-GCCAAAGACAGAAGCCAGGA and 5'-CAGGATAGGACTCAGGGAATACAG), *notch1* NC\_000068.7 (5'-GTGACCGTGGAACGTCTA 5'-CTGTCCTAGGGCTCCAC)

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**Supplementary Figure S1: Effect of JNK2 on mammary epithelial cells. A.** Representative images of adult (8 wk-old) ducts probed for JNK2 expression; **B-C.** Images of p63<sup>+</sup> basal cells and ER<sup>+</sup> luminal cells in adult ducts (n = 3); **D.** *Jnk2wt* and *jnk2ko* mammary epithelial cells were isolated and grown in 3D culture. Acinar diameter was measured at indicated time points. Linear regression analysis was performed, \*\*\*p < 0.0001; **E-F.** Quantification of Ki67<sup>+</sup> and cleaved caspase 3<sup>+</sup> cells grown in 3D culture. Nonparametric *t*-test was performed, \*\*p < 0.001.





Supplementary Figure S2: Representative images of JNK2 effects on mammary cell differentiation and Notch activity. A. Representative staining where arrows indicate p63+ cells in 3D culture; B-C. Representative staining of ER<sup>+</sup> mammary cells in 3D culture; D. Representative staining of Notch1<sup>ICD</sup> in TEBs.



**Supplementary Figure S3: JNK2 effects on gene expression in** *p53ko* **tumors or tumor cell lines.** A. RNA from tumors was analyzed for Notch1 expression using qPCR. A nonparametric, two-tailed *t*-test was used to detect statistical differences between two groups; **B.** *p53ko;jnk2ko* GFP and GFP-JNK2 expressing cells were transfected with Notch1 promoter-luciferase constructs and analyzed for promoter activity. A 1 Way ANOVA with post-hoc *t*-test was performed; **C.** Gene Set Enrichment Analysis (GSEA) of *p53ko;jnk2ko* tumors; **D.** Genes significantly elevated with false discovery rate (FDR) of < 5% in the ATM/BRCA1 pathway.



**Supplementary Figure S4: JNK2 promotes EMT in the p53ko tumor model.** A. Expression of mesenchymal/stem (red) and epithelial (blue) markers in *p53ko* tumors (n = 8), **B.** Fold changes observed in microarray analysis of EMT/stem and differentiation markers in *p53ko;jnk2ko* GFP and GFP-JNK2 cell lines; **C.** Expression of EMT/stem-related proteins in *p53ko;jnk2ko* GFP and GFP-JNK2 cell lines by western blot; **D.** Expression of JNK1 in *p53ko;jnk2ko* GFP and GFP-JNK2 cell lines by western blot. A nonparametric, two-tailed *t*-test was also performed for A. \*p < 0.05.





Supplementary Figure S5: JNK2 promotes vimentin expression and inhibits e-cadherin in the *p53ko;jnk2ko* cell line. A. Adherent cells were stained using vimentin and e-cadherin primary antibodies; B. Adherent cells grown in culture.