

# Supplementary materials

## **A humanized RANKL transgenic mouse model of progestin-induced mammary carcinogenesis for evaluation of novel therapeutics**

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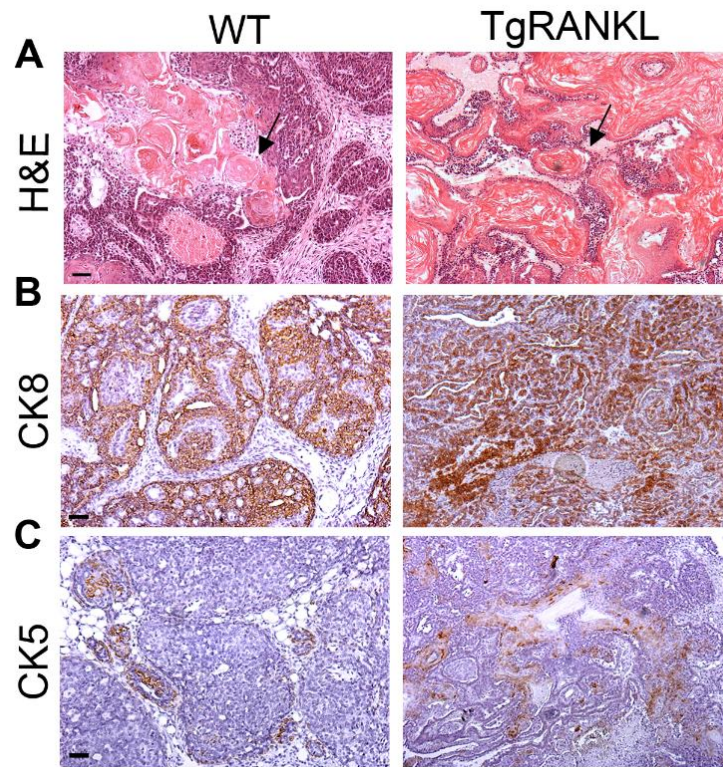
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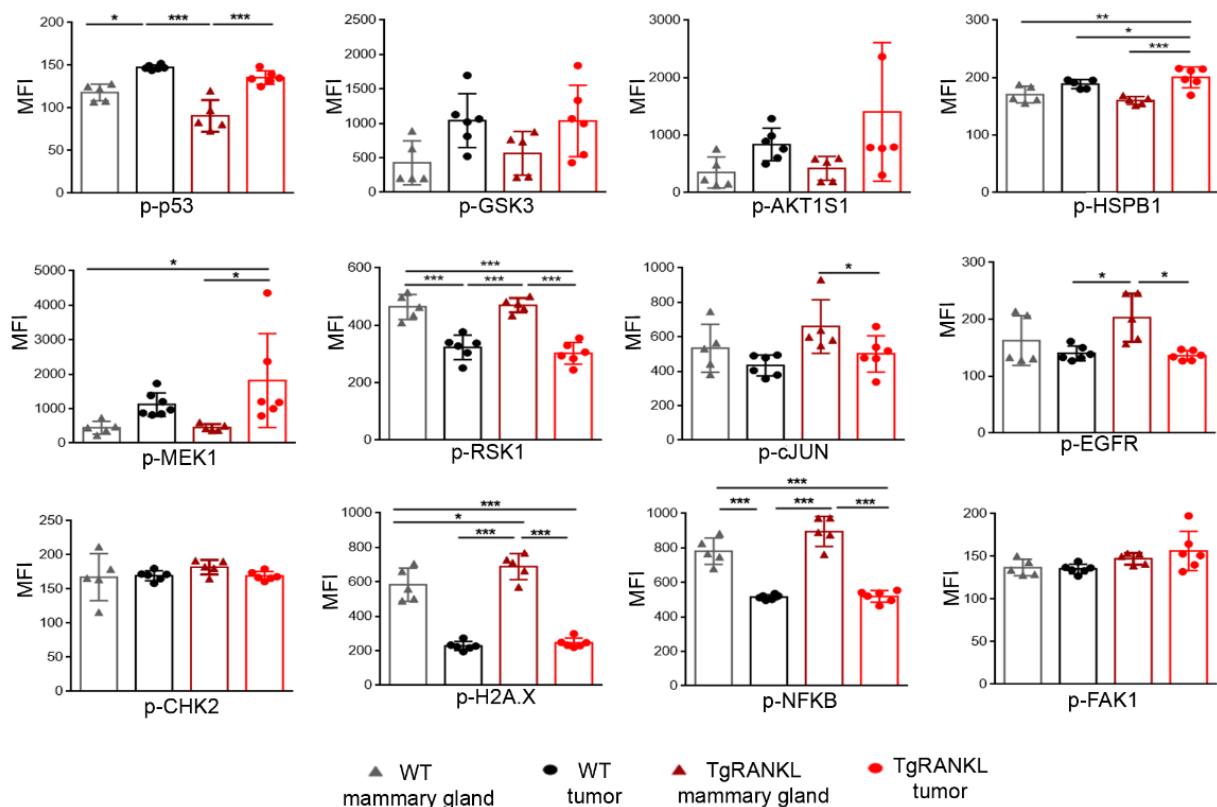
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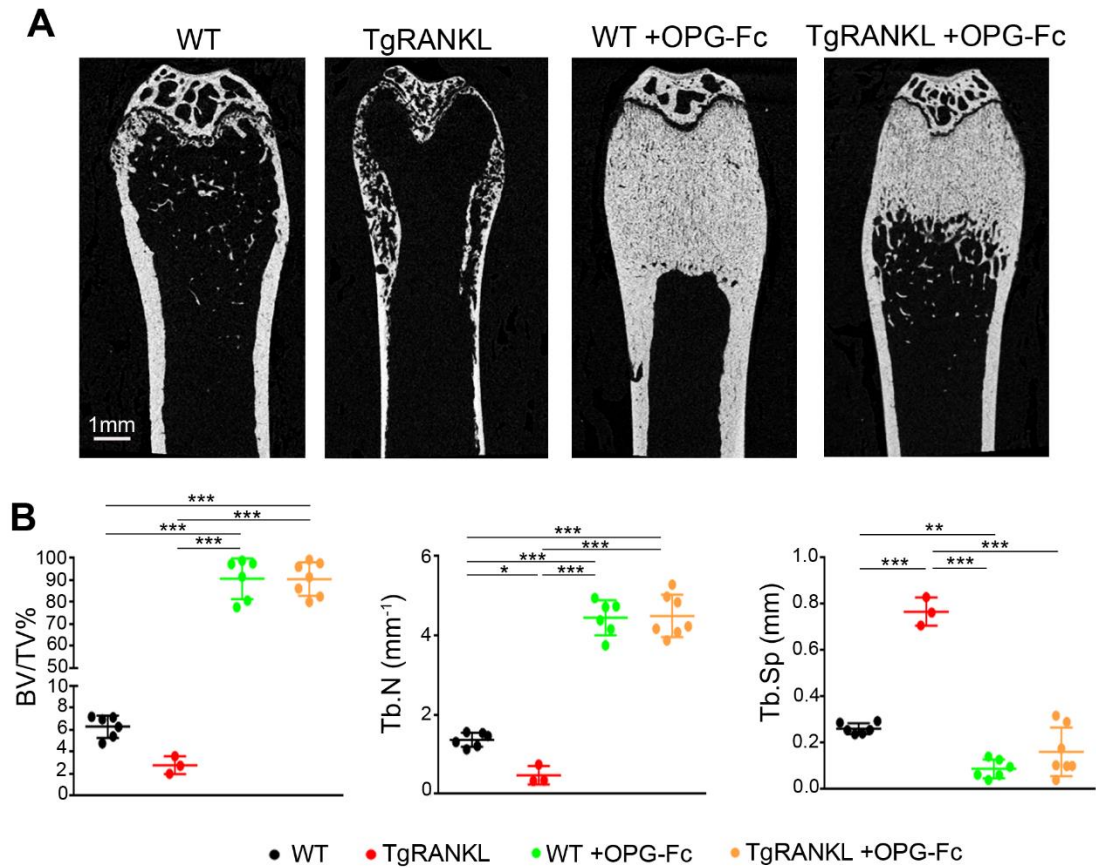
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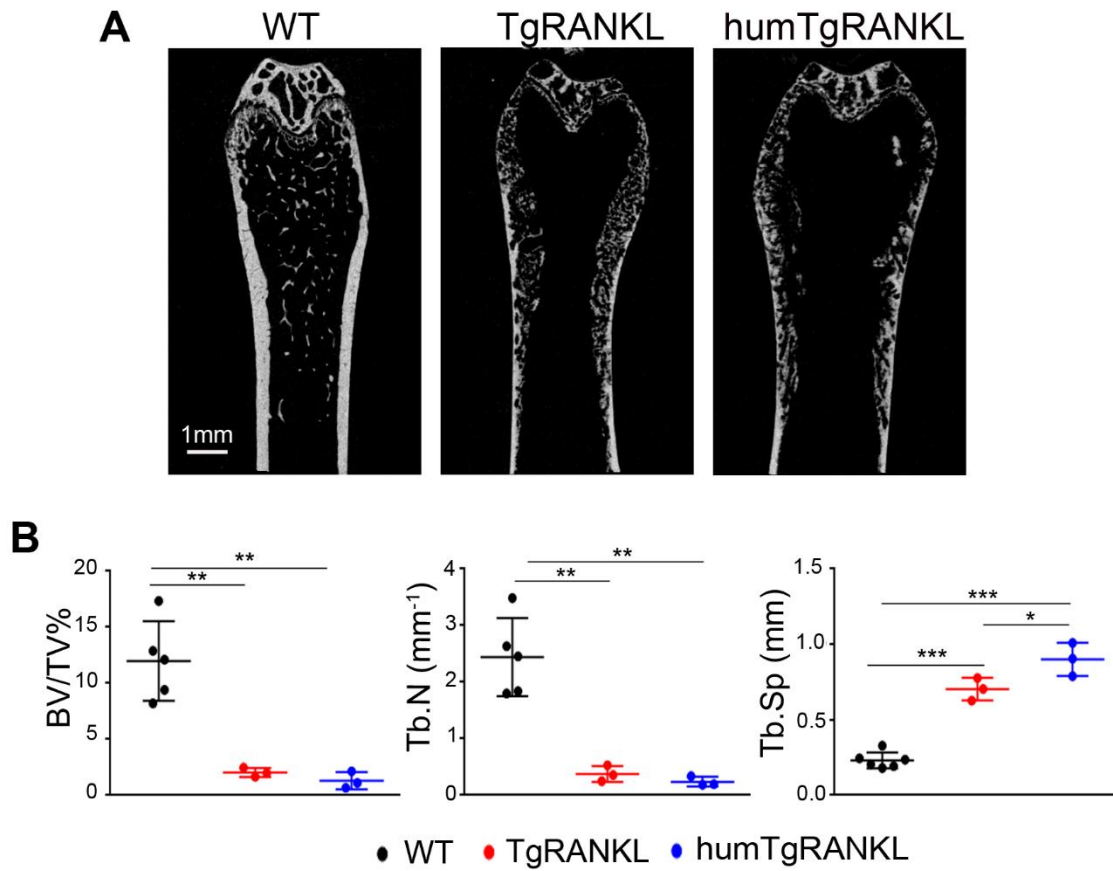
**Figure S1. Histological analysis of MPA/DMBA-induced mammary tumors in WT and TgRANKL mice.** Representative histological sections of mammary tumors isolated from WT and TgRANKL mice stained with **(A)** hematoxylin and eosin, **(B)** Cytokeratin 8, and **(C)** Cytokeratin 5 (n=3/group). Keratin pearls are indicated by arrows. Scale bar, 80µm.



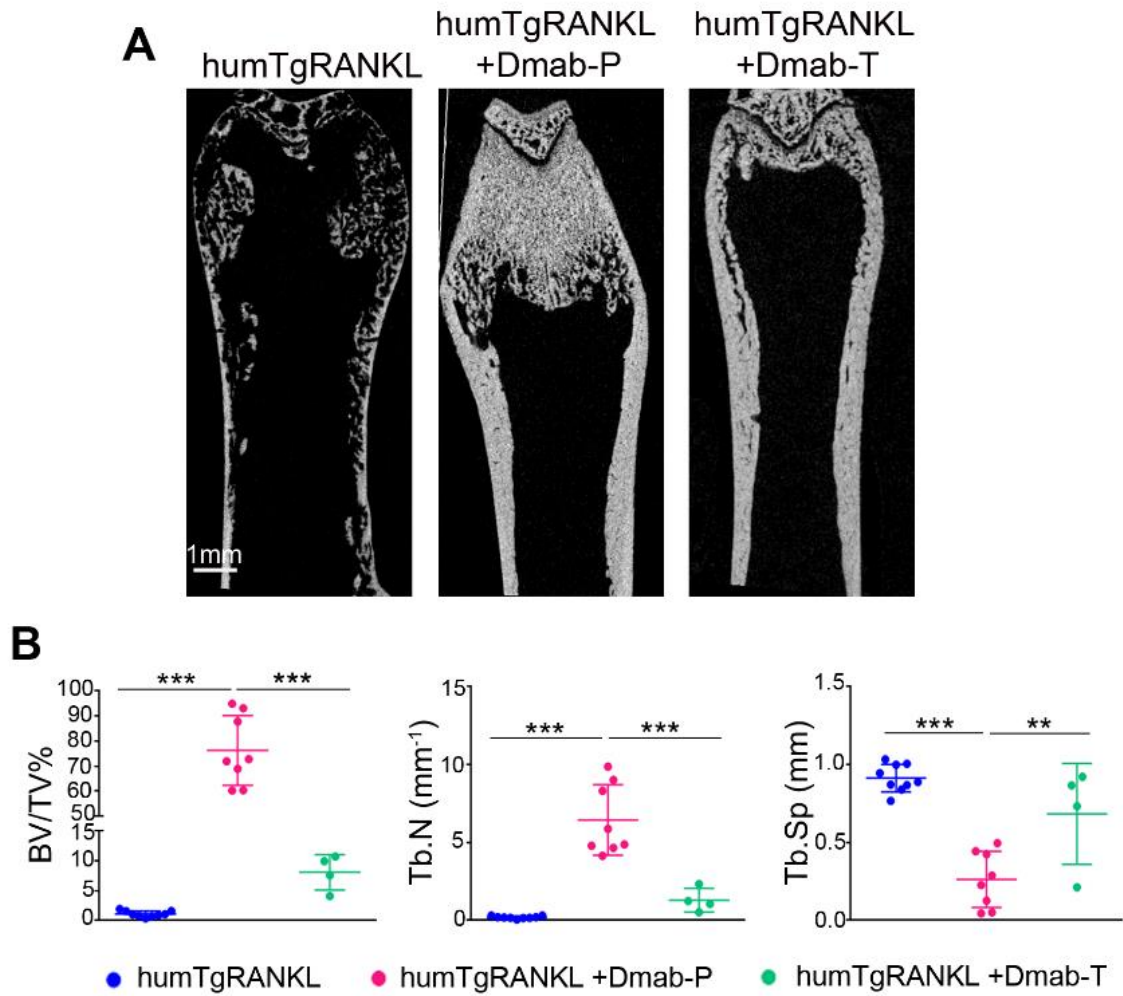
**Figure S2. Additional multiplex ELISA analysis for phospho-proteins.** p-p53, p-GSK3, p-AKT1S1, p-HSPB1, p-MEK1, p-RSK1, p-cJUN, p-EGFR, p-CHK2, p-H2A.X, p-NFkB and p-FAK1 were measured in mammary glands and tumors from WT and TgRANKL mice (n=5=6/group). Comparison was performed with one-way ANOVA and Tukey's post hoc test (\* p<0.05, \*\* p<0.01, \*\*\* p<0.001).



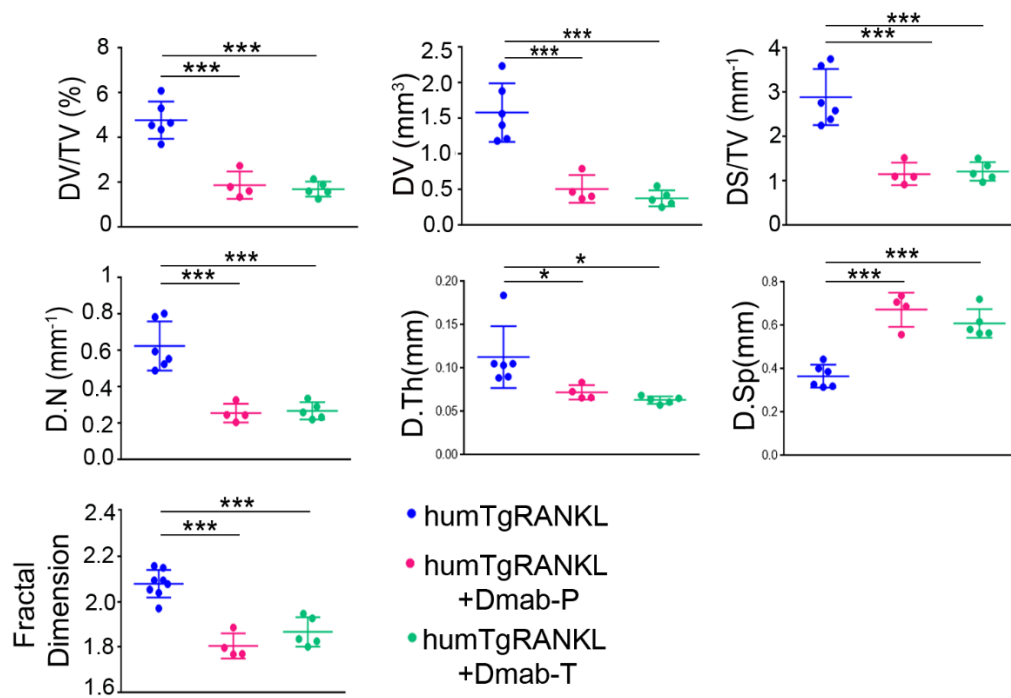
**Figure S3. OPG-Fc treatment prevented bone resorption in WT and TgRANKL mice. (A)** Representative microCT 2D images and **(B)** quantitative analysis for BV/TV%, Tb.N, and Tb.Sp in the trabecular bone of femurs from MPA/DMBA-treated WT and TgRANKL mice with or without OPG-Fc treatment (n = 3-7/group). Data are shown as mean  $\pm$  SD. One-Way ANOVA was performed for statistical analysis (\*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001).



**Figure S4. Humanized TgRANKL mice developed an osteoporotic phenotype similar to TgRANKL mice.** (A) Representative microCT 2D images and (B) quantitative analysis for BV/TV%, Tb.N, and Tb.Sp in the trabecular bone of femurs from WT, TgRANKL and humTgRANKL mice (n = 3-5/ group). Data are shown as mean ± SD. One-Way ANOVA was performed for statistical analysis (\*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001).



**Figure S5. Effect of either prophylactic or therapeutic treatment of humTgRANKL mice with denosumab in bone architecture.** (A) Representative microCT 2D images and (B) quantitative analysis for BV/TV%, Tb.N, and Tb.Sp in the trabecular bone of femurs from MPA/DMBA-treated humTgRANKL mice without or with Dmab treatment either prophylactically (Dmab-P) or therapeutically (Dmab-T) (n = 4-9 group). Data are shown as mean  $\pm$  SD. One-Way ANOVA was performed for statistical analysis (\*\* $p$  < 0.01, \*\*\* $p$  < 0.001).



**Figure S6. Denosumab restores MPA/DMBA-induced mammary gland density in humTgRANKL mice.** Quantitative microCT analysis of mammary gland epithelium from humTgRANKL mice either untreated or treated with Dmab prophylactically or therapeutically (n=4-6 mice/group) DV/TV%: Ductal volume/Tissue volume %, DV (mm<sup>3</sup>): Ductal volume, DS/TV (mm<sup>-1</sup>): Ductal Surface/Tissue Volume, Volume, D.N (mm): Ductal number, D.Th (mm): Ductal Thickness, D.Sp (mm): Ductal Separation, Fractal Dimension. Data are shown as mean  $\pm$  SD. One-Way ANOVA was performed for statistical analysis (\* p<0.05, \*\*\* p<0.001).

**Table S1.** Primer sequences used in qPCR.

Target genes	Forward (5'-3')	Reverse (5'-3')
<i>B2M</i>	ACATCAAGAAGGTGGTGAAGCAGG	AGTTGCTGTTGAAGTCGCAGGAGA
<i>mu+huRankl</i>	ACCTGTACGCCAACATTTGC	CTTGGG ATTTTGATGCTGGT
<i>huRankl</i>	ACGCGTATTTACAGCCAGTG	5CCCGTAATTGCTCCAATCTG
<i>muRankl</i>	TGTACTTTTCGAGCGCAGATG	AGGCTTGTTTCATCCTCCTG
<i>muOpg</i>	TGATGTATGCCCTCAAGCAC	TTGTGAAGCTGTGCAGGAAC
<i>muRank</i>	TCTTATGTTGGGGTCCATCC	AATAAGCTTAGCCCCGAACC
<i>muPr</i>	CTCCGGGACCGAACAGAGT	ACAACAACCCTTTGGTAGCAG
<i>muEra</i>	AATGAAATGGGTGCTTCAGG	ATAGATCATGGGCGGTTTCAG
<i>muCcmd1</i>	GCGTACCCTGACACCAATCTC	CTCCTCTTCGCACTTCTGCTC
<i>muLgr5</i>	CCTACTCGAAGACTTACCCAGT	GCATTGGGGTGAATGATAGCA
<i>muLgr4</i>	TACAACCTGGCTGGTAACGACC	TTGAGTTCTTTCAACCCAGACAA
<i>muSox2</i>	AAAGGGTTCTTGCTGGGTTT	AGACCACGAAAACGGTCTTG
<i>muSox9</i>	CGGAACAGACTCACATCTCTCC	CTTGACACGTCGGTTTTTG
<i>muSlug</i>	TCAACGCCTCCAAGAAGCCCA	ATAGGGCTGTATGCTCCCGAGGT



**Table S2.** Protein assays of the custom-developed 19-plex assay panel and their phosphorylation residues.

	<b>Protein-assay abbreviation</b>	<b>Protein name</b>	<b>Phosphorylation Residue</b>
1	Smad3	Mothers against decapentaplegic homolog 3	S423/S425
2	p53	Cellular tumour antigen p53	S15
3	AKTS1	Proline-rich AKT1 substrate 1	T246
4	GSK3	Glycogen synthase kinase-3 alpha/beta	S21/9
5	AKT1	RAC-alpha serine/threonine-protein kinase	S473
6	HSP27 (HSPB1)	Heat shock protein beta-1	S78/S82
7	p38	Mitogen-activated protein kinase 14/11	T180/Y182
8	MEK1	Dual specificity mitogen-activated protein kinase kinase 1	S217/S221
9	RSK1	Ribosomal protein S6 kinase alpha-1	S380
10	CREB1	Cyclic AMP-responsive element-binding protein 1	S133
11	cJUN	Transcription factor Jun	S63
12	EGFR	Epidermal growth factor receptor	Y1068
13	PTN11	Tyrosine-protein phosphatase non-receptor type 11	Y542
14	CHK2	Serine/threonine-protein kinase Chk2	T68
15	H2A.X	Histone H2AX	S139
16	NF-KB	Nuclear factor kappa-light-chain-enhancer of activated B cells	S536
17	STAT3	Signal transducer and activator of transcription 3	Y705
18	ERK1	Mitogen-activated protein kinase 3	T202/Y204
19	FAK1	Focal adhesion kinase 1	Y397