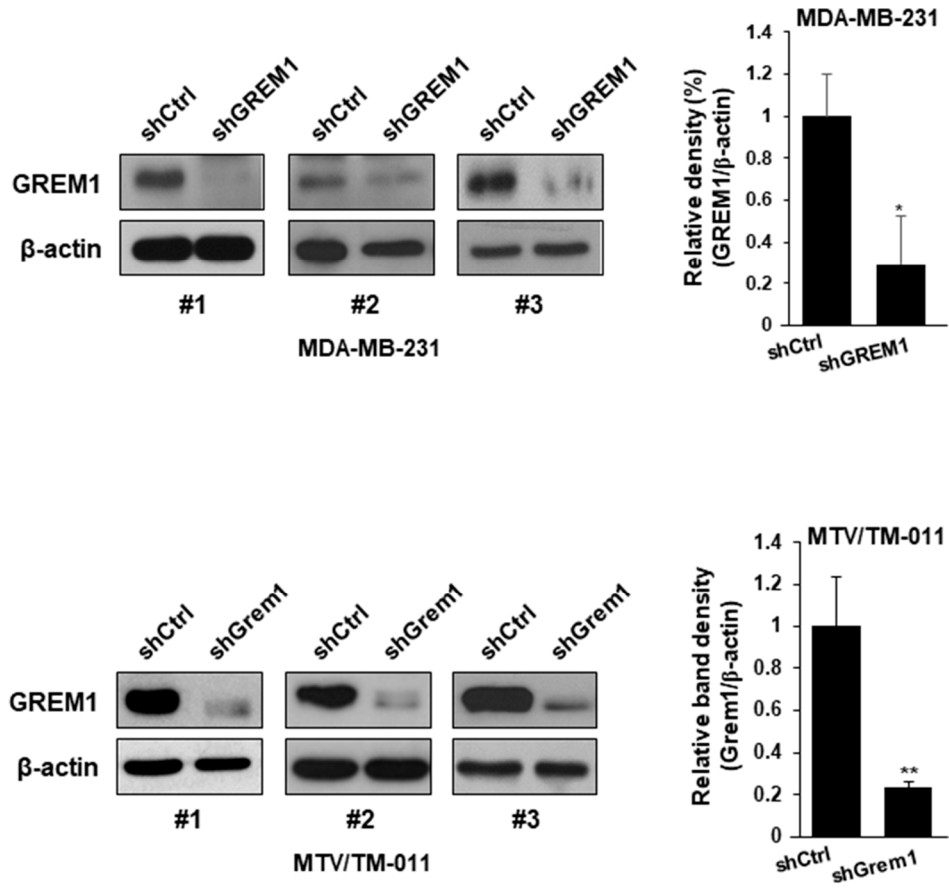
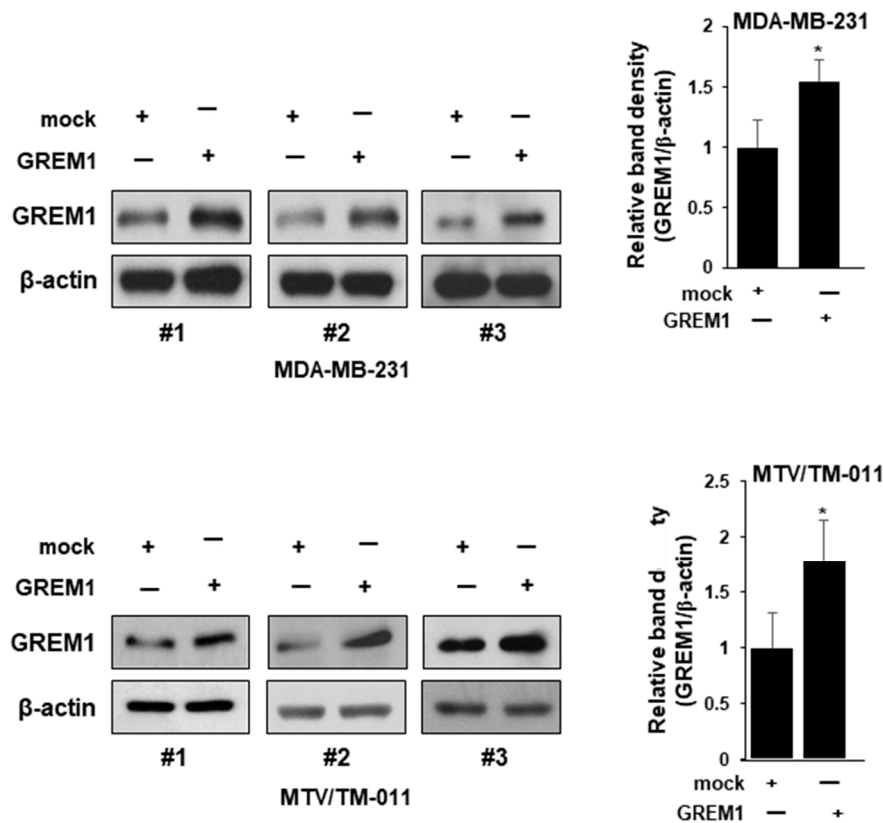


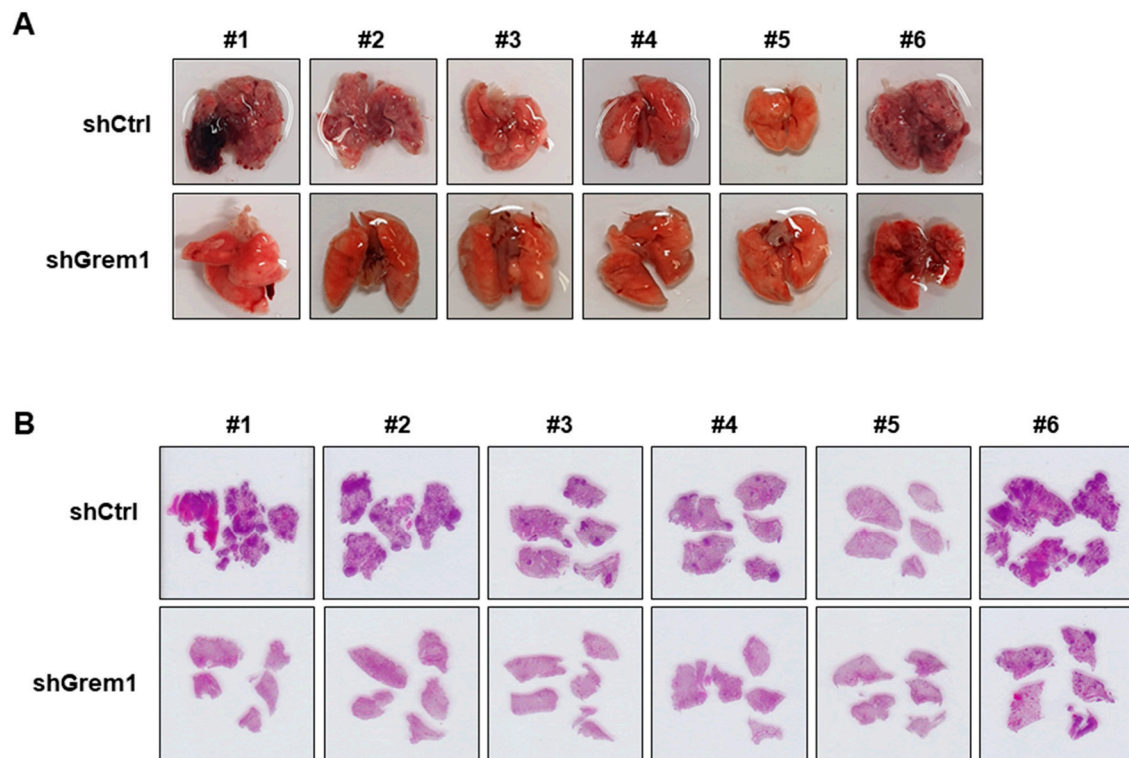
Supplementary Figure Legends

**A**

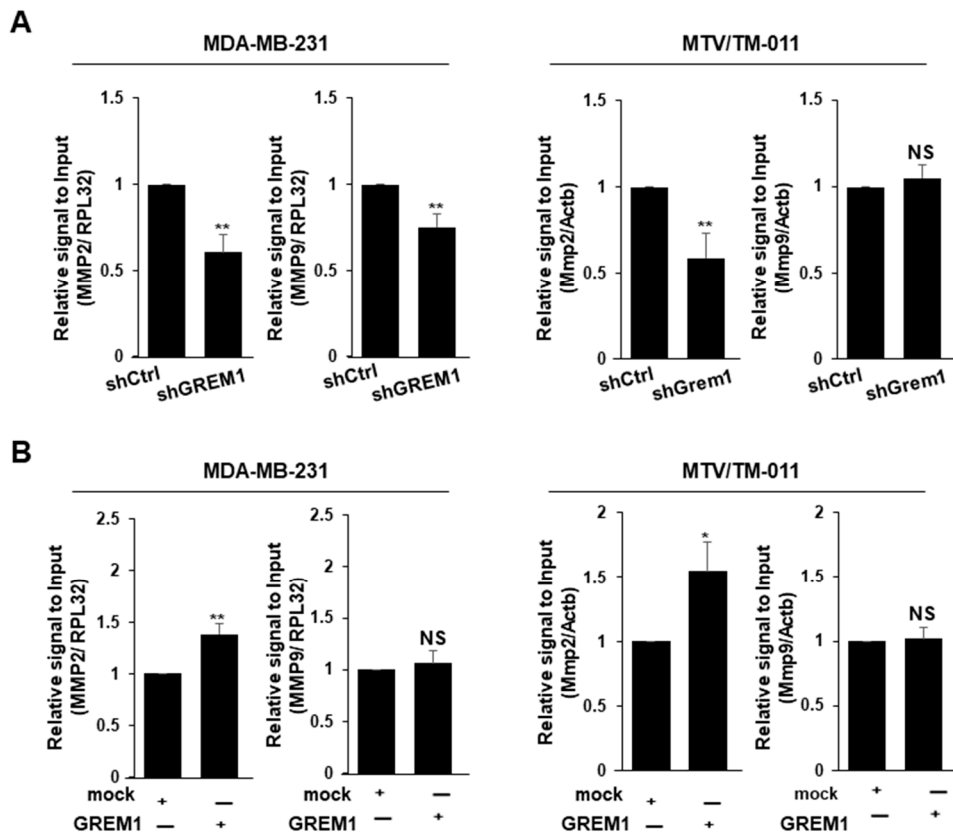


**B**

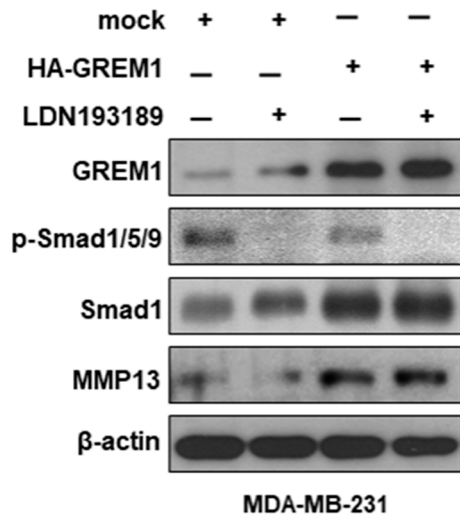
**Supplementary Fig. 1. The expression of GREM1 was verified in each indicated cell line. (A)** The cell lysates of MDA-MB-231 and MTV/TM-011 cell lines stably expressing shCtrl or shGREM1 were immunoblotted with GREM1 antibody. **(B)** MDA-MB-231 and MTV/TM-011 cells were transfected with each plasmid expressing mock or GREM1 for 48 h and the lysates were immunoblotted with GREM1 antibody. All quantitative values were obtained using three sets of independently separated samples on different days. Data are presented as the mean  $\pm$  SD of three independent experiments. Two-sided t test. \*,  $p < 0.05$ ; \*\*,  $p < 0.01$ .



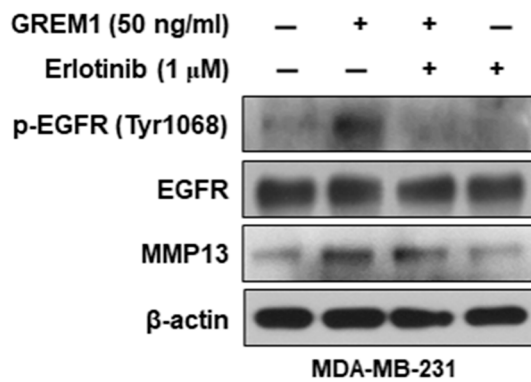
**Supplementary Fig. 2. GREM1 knockdown suppressed lung metastasis of breast cancer cells. (A)** Images of lung metastasis nodules formed by the injection of MTV/TM-011 cells expressing shCtrl or shGrem1 into the mouse mammary fat pads (n = 6/ each group). **(B)** Whole lung tissues were stained by H&E (n = 6/ each group).



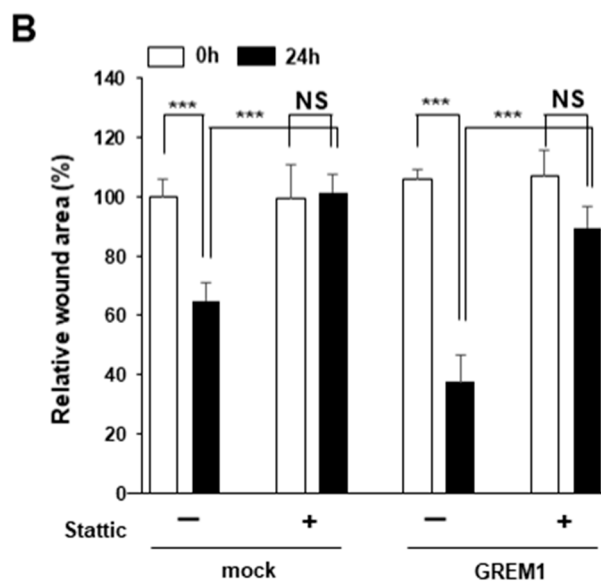
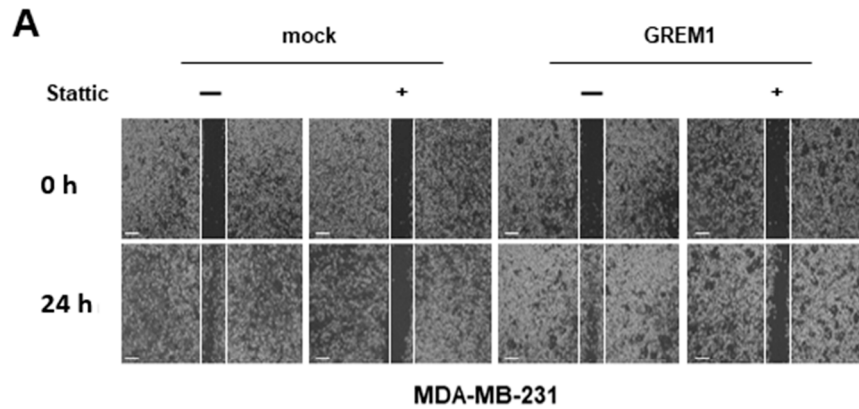
**Supplementary Fig. 3. GREM1 regulated the expression of MMP2 or MMP9. (A)** Relative mRNA levels of *MMP2* and *MMP9* in GREM1-depleted breast cancer cells. **(B)** Relative mRNA levels of *MMP2* and *MMP9* in GREM1-overexpressing breast cancer cells. MDA-MB-231 and MTV/TM-011 cells were transfected with each indicated plasmid for 48 h, and mRNA levels of genes were quantitated by qPCR analysis. Data are presented as the mean  $\pm$  SD of three independent experiments. Two-sided t test. \*,  $p < 0.05$ ; \*\*,  $p < 0.01$ ; NS, not significant.

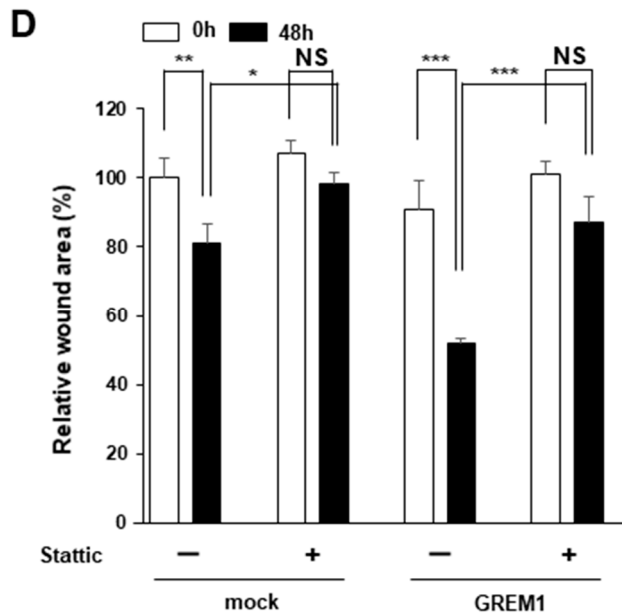
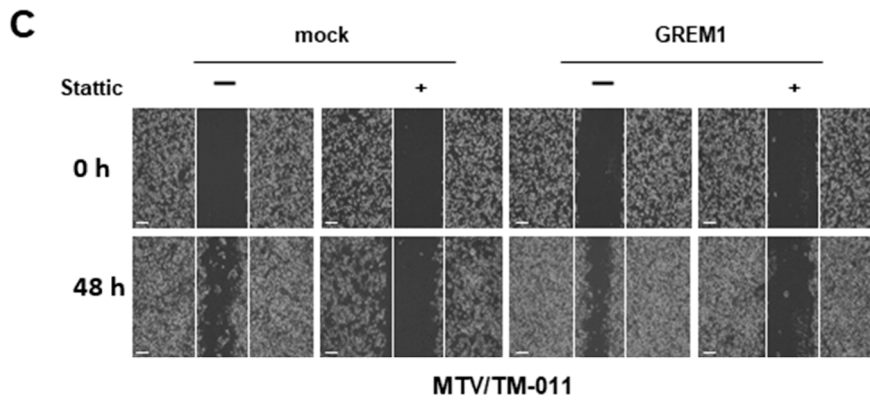


**Supplementary Fig. 4. The expression of MMP13, which was increased by GREM1, was not affected by treatment of LDN193189, a BMP pathway inhibitor.** MDA-MB-231 cells were transfected with each plasmid expressing mock or GREM1 for 24 h and then incubated in the absence or presence of LDN193189 (1  $\mu$ M) for another 24 h. The protein lysates were immunoblotted with each indicated antibody.



**Supplementary Fig. 5. The expression of MMP13, which was increased by GREM1, was not reduced by treatment of erlotinib, an EGFR tyrosine kinase inhibitor.** MDA-MB-231 cells were pretreated with erlotinib (1  $\mu$ M) for 24 h and then incubated with GREM1 (50 ng/ml) for another 30 min (pEGFR and EGFR) or 48 h (MMP13). The protein lysates were subjected to immunoblot analysis.





**Supplementary Fig. 6. The enhanced migratory capacity of GREM1-overexpressing cells was inhibited by Static treatment. (A-D)** MDA-MB-231 and MTV/TM-011 cells were transiently transfected with plasmid overexpressing mock or GREM1 for 48 h and seeded again in culture-inserts with medium containing Static (5  $\mu$ M). Images of wound sites were captured at 0 h (control) and the indicated time periods of incubation using an inverted microscope (4 x magnification). Scale bar = 200  $\mu$ m. Each wound area was determined using Image J software. Data are presented using triplicate wells per group and statistical significance was determined by one-way ANOVA. \*,  $p < 0.05$ ; \*\*,  $p < 0.01$ ; \*\*\*,  $p < 0.001$ ; NS, not significant.

**Supplementary Table 1. qRT-PCR primer sequences**

<b>Primers (Human)</b>	<b>Forward</b>	<b>Reverse</b>
GREM1	TCATCAACCGCTTCTGTTACG	GGCTGTAGTTCAGGGCAGTT
MMP2	GATACCCCTTTGACGGTAAGGA	CCTTCTCCCAAGGTCCATAGC
MMP3	CGGTTCCGCCTGTCTCAAG	CGCCAAAAGTGCCTGTCTT
MMP9	AGACCTGGGCAGATTCCAAAC	CGGCAAGTCTTCCGAGTAGT
MMP11	GGGTGTACGACGGTGAAAAG	GTGGAAACGCCAGTAGTCCC
MMP13	CCAGACTTCACGATGGCATTG	GGCATCTCCTCCATAATTTGGC
MMP19	GCAATGTGGCTCCCTTGAC	TCAGTCCAGAACTCGTCTTCG
RPL32	TTAAGCGTAACTGGCGGAAAC	AAACATTGTGAGCGATCTCGG

<b>Primers (Mouse)</b>	<b>Forward</b>	<b>Reverse</b>
Grem1	GGGACCCTACTGCCAACAG	TTTGCACCAATCTCGCTTCAG
Mmp2	ACCTGAACACTTTCTATGGCTG	CTTCCGCATGGTCTCGATG
Mmp3	TTAAAGACAGGCACTTTTGGCG	CCCTCGTATAGCCCAGAACT
Mmp9	CTGGACAGCCAGACACTAAAG	CTCGCGGCAAGTCTTCAGAG
Mmp11	CCGGAGAGTCACCGTCATC	GCAGGACTAGGGACCCAATG
Mmp13	CTTCTTCTTGTTGAGCTGGACTC	CTGTGGAGGTCACTGTAGACT
Mmp19	CTGTGGCTGGCATTCTTACTT	GGGCAGTCCAGATGCTTCC
Actb	GGCTGTATTCCCCTCCATCG	CCAGTTGGTAACAATGCCATGT