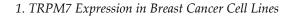
Supplementary Materials



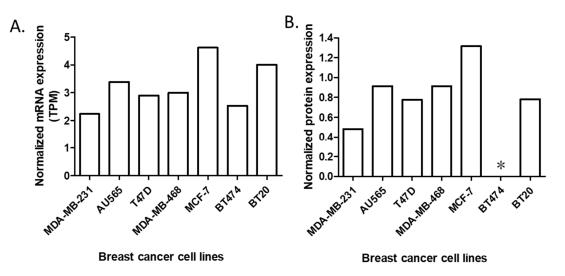


Figure S1. TRPM7 expression in breast cancer cell lines. **A.** The mRNA expression. The mRNA expression data (RNA seq) was downloaded and extracted from the Cancer Cell Line Encyclopedia (CCLE) [1]. **B.** The protein expression of TRPM7. The protein proteomics expression data was downloaded and extracted from the Quantitative Proteomics of the Cancer Cell Line Encyclopedia [2].

2. Additional Fluorescence Quench Data

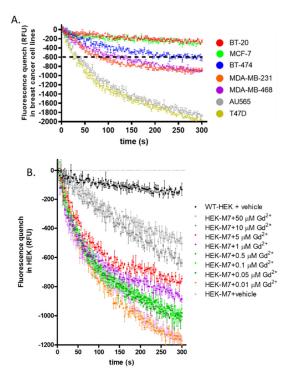


Figure S2. Fluorescence quench assay (n = 3). **A.** Fluorescence quench in Fura-2 loaded breast cancer cell lines (n = 3). The methods were described in the main text of the paper, n = 3. Cell lines with quench amounts larger than 600 relative fluorescence units (RFU) were used in the TRPM7 functional study. **B.** Effect of Gd²⁺ on TRPM7. WT-HEK was used as negative control. [Gd²⁺] \ge 0.05 µM concentration-dependently decreased the influx of Mn²⁺ in HEK-M7.

3. Effect of TRM7 Knockout on Migration of MDA-MB-231.

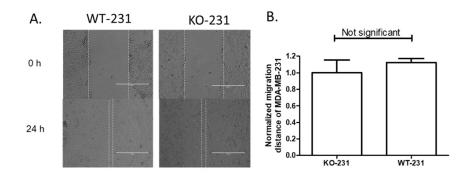


Figure S3. The effect of TRPM7 knockout on cell migration of breast cancer cell line MDA-MB-231. A. Representative image from wound healing assay. B. The data was normalized by the division of the average migration distance of KO-231. 24 hours after the creation of the wound, there was no significant difference in migration distance between KO-231 and WT-231 (p > 0.05).

References

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