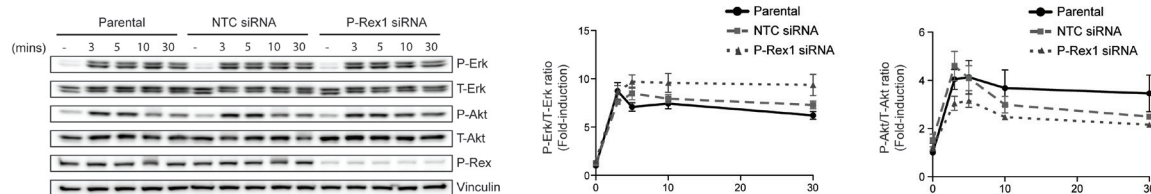


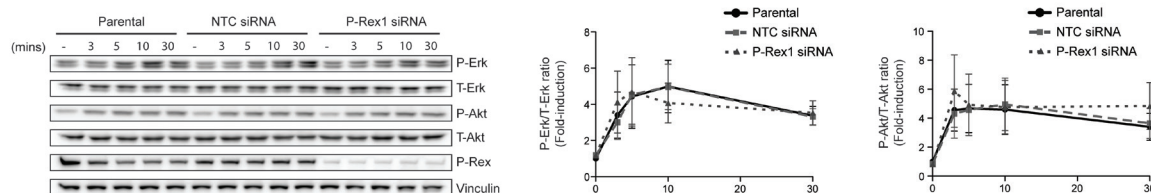
P-Rex1 is dispensable for Erk activation and mitogenesis in breast cancer

SUPPLEMENTARY MATERIALS

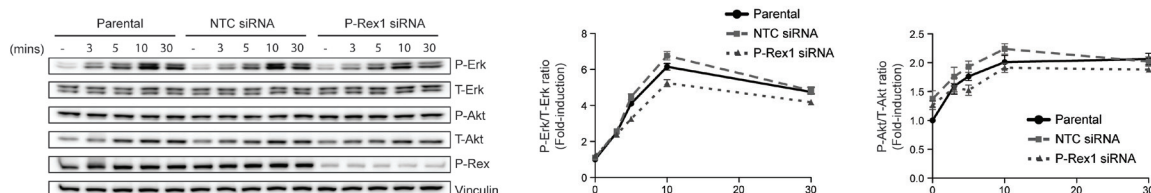
EGF



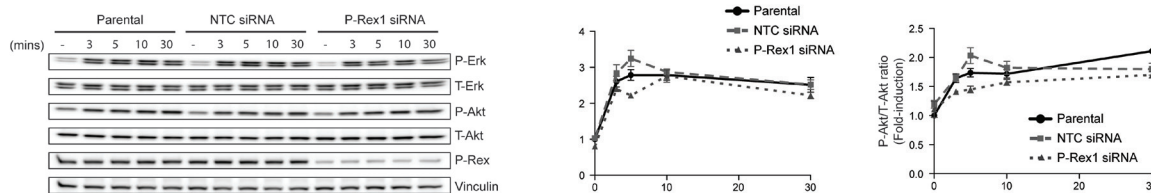
IGF-1



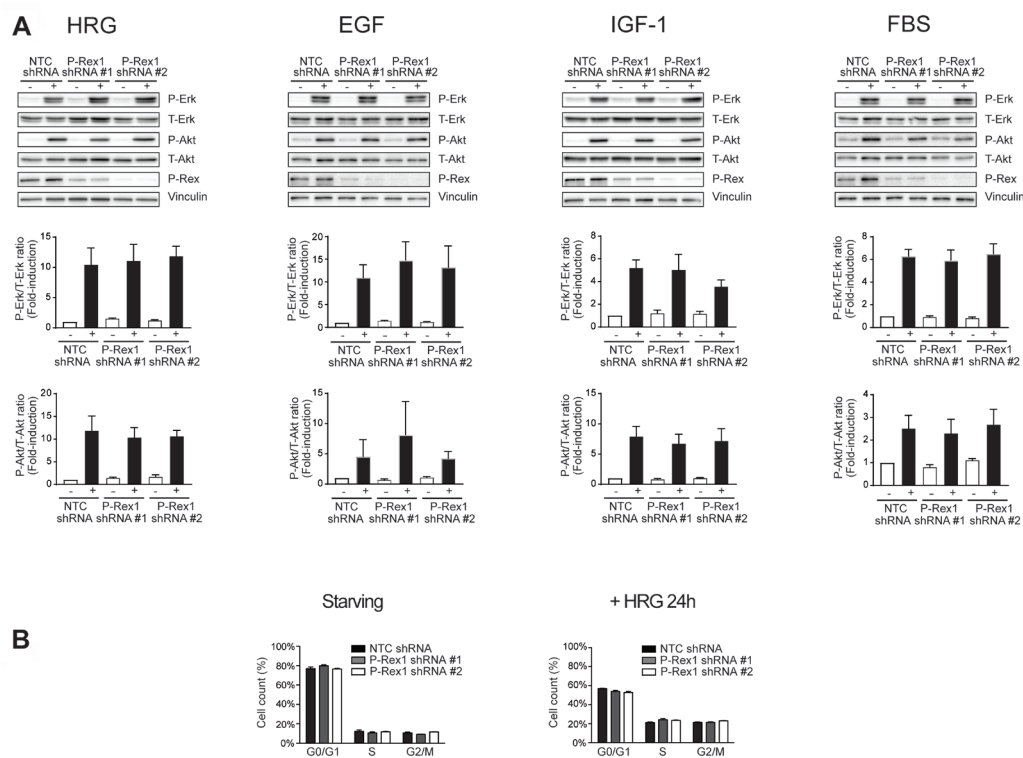
FBS



SDF-1



Supplementary Figure 1: Time-course of Erk and Akt activation in MCF-7 cells. Cells were transfected with P-Rex1 siRNA (pool) or non-target control (NTC) siRNA. After 16 h, cells were serum starved for 24 h and stimulated with EGF (100 ng/ml), IGF-I (100 ng/ml), FBS (10%), or SDF-1 (100 ng/ml) at different times. *Left panels*, representative experiments. *Middle and right panels*, densitometric analysis of phospho-Erk and phospho-Akt levels normalized to total levels. Results were expressed as fold-change relative to parental cells with vehicle stimulation. Data were expressed as mean ± S.E.M. of 3 independent experiments. No statistically significant differences were observed between parental, NTC and P-Rex1-depleted cells.



Supplementary Figure 2: Activation of Erk and Akt, and cell cycle analysis in MCF-7 cells subject to stable P-Rex1 silencing. P-Rex1-depleted MCF-7 cells lines were generated by infection with two different P-Rex1 shRNA lentiviruses (#1 and #2), followed by puromycin selection. As a control, a NTC shRNA lentivirus was used. **(A)** Cells were serum starved for 24 h and stimulated with HRG (20 ng/ml, 5 min), EGF (100 ng/ml, 2 min), IGF-I (100 ng/ml, 5 min) or FBS (10%, 10 min). Activation of Erk and Akt was determined by Western blot using phospho-specific antibodies. *Upper panels*, representative experiments. *Middle panels*, densitometric analysis of phospho-Erk, normalized to the total Erk. *Lower panels*, densitometric analysis of phospho-Akt, normalized to the total Akt. Results were expressed as fold-change relative to parental cells with vehicle stimulation. Data were expressed as mean \pm S.E.M. of 3 independent experiments. **(B)** Cells were serum starved for 24 h, stimulated with HRG (20 ng/ml) or vehicle for 24 h, and subject to cell cycle analysis by FACS. Graphs show the distribution of cells in the different phases of cell cycle. Data are expressed as mean \pm S.E.M. of 3 independent experiments.