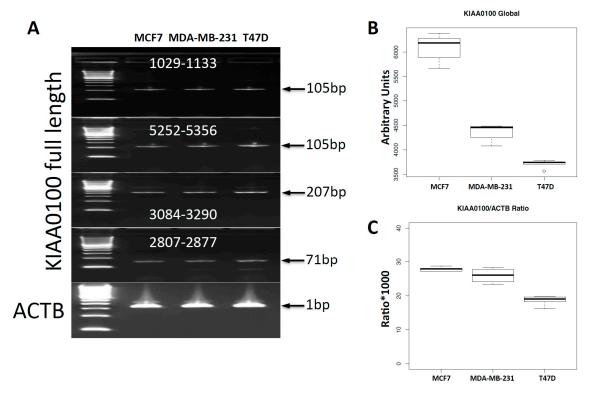
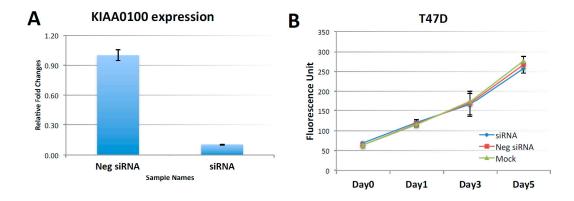
## Supplementary Materials: KIAA0100 Modulates Cancer Cell Aggression Behavior of MDA-MB-231 through Microtubule and Heat Shock Proteins

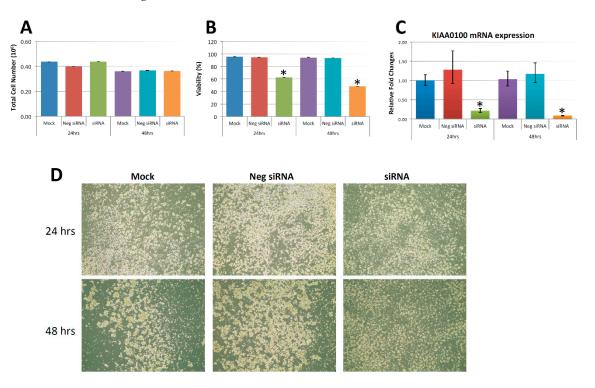
Zhenyu Zhong, Vaishali Pannu, Matthew Rosenow, Adam Stark and David Spetzler



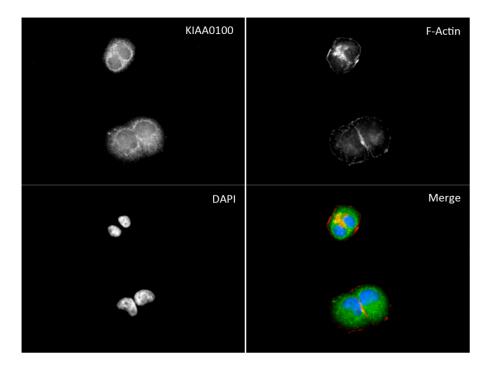
**Figure S1.** Expression of full length KIAA0100 in three breast cancer cell lines. (**A**) RT-PCR of full length KIAA0100 by targeting different regions of the mRNA on MCF7, MDA-MB-231. 4 sets of primers targeted to different region of the full length KIAA0100 mRNA was used. All three cell lines were confirmed to express full length KIAA0100; (**B**) semi-quantitative mass spectrometry analysis of KIAA0100 in the three different cell lines, normalized globally. Expression of KIAA0100 in MCF7> MDA-MB-231 > T47D, however, they are at similar range within 2 fold difference; (**C**) semi-quantitative mass spectrometry analysis of KIAA0100 in the three different cell lines, normalized by β-actin. Expression of KIAA0100 in MCF7 and MDA-MB-231 were similar while T47D was slightly lower, but they are all at similar level range with minor difference.



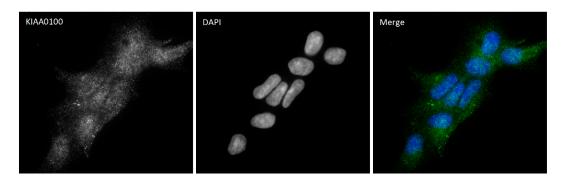
**Figure S2** Silencing expression of KIAA0100 in breast cancer cell T47D. T47D cells were transfected with KIAA0100 siRNA in a forward transfection manner, qPCR was used to examine the reduction of KIAA0100 mRNA 24 hours after the treatment. Cell proliferation/growth was examined. **(A)** RT-qPCR showed the efficiency of silencing the expression of KIAA0100 in T47D. 24 hours after the treatment, mRNA level of KIAA0100 was reduced >80%.; **(B)** Anchorag-dependent cell proliferation/growth was examined for up to 5 days after treatment, no significant difference between the Mock, neg siRNA control treated cell and the KIAA0100 siRNA treated cells.



**Figure S3.** Silencing expression of KIAA0100 in breast cancer cell MDA-MB-231 by siRNA targeted to exon 17. **(A)** Total cell number in 24 and 48 hours after treatment; **(B)** Cell viability in 24 and 48 hours after the treatment; **(C)** KIAA0100 RNA expression level in 24 and 48 hours after the treatment by qPCR; **(D)** Cell aggregation in 24 and 48 hours after the treatment under microscope (\* p < 0.05). Images were acquired at 40x.



**Figure S4.** Immunofluorescence of native KIAA0100 protein in breast cancer cell T47D. KIAA0100 was co-immunostained with anti-KIAA010 antibody and Phalloidin-564 as well as DAPI. Native KIAA0100 stained as in the cytoplasm as speckles, the pattern is similar to that from MDA-MB-231. Images were acquired at 400x.



**Figure S5.** Immunofluorescence of native KIAA0100 protein in HEK293. KIAA0100 was immunostained with anti-KIAA010 antibody and DAPI. Native KIAA0100 stained as in the cytoplasm as speckles, the pattern of KIAA0100 is also similar to that from MDA-MB-231 as well as T47D. Images were acquired at 400x.



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