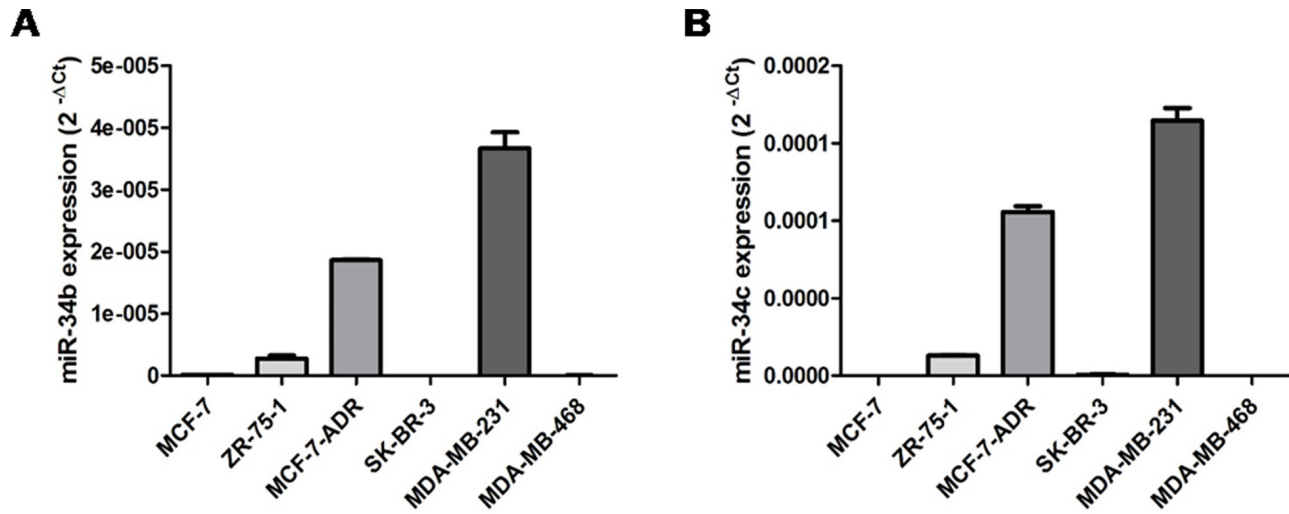
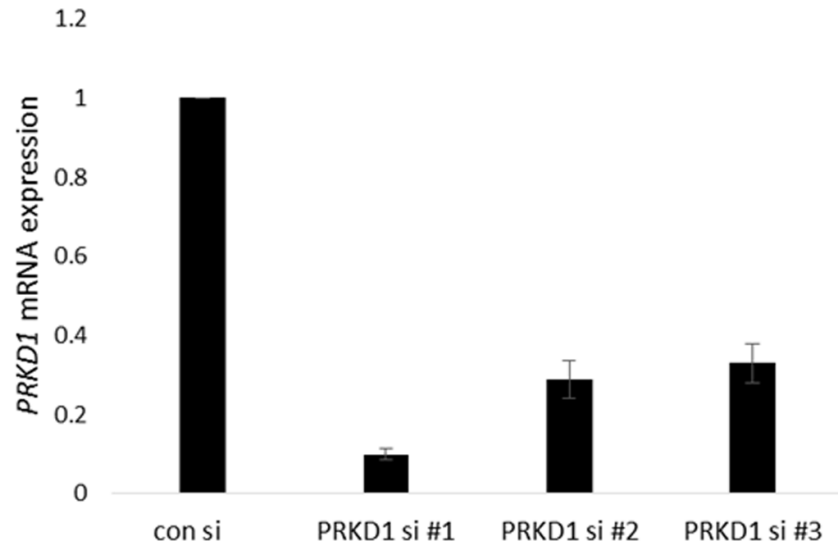


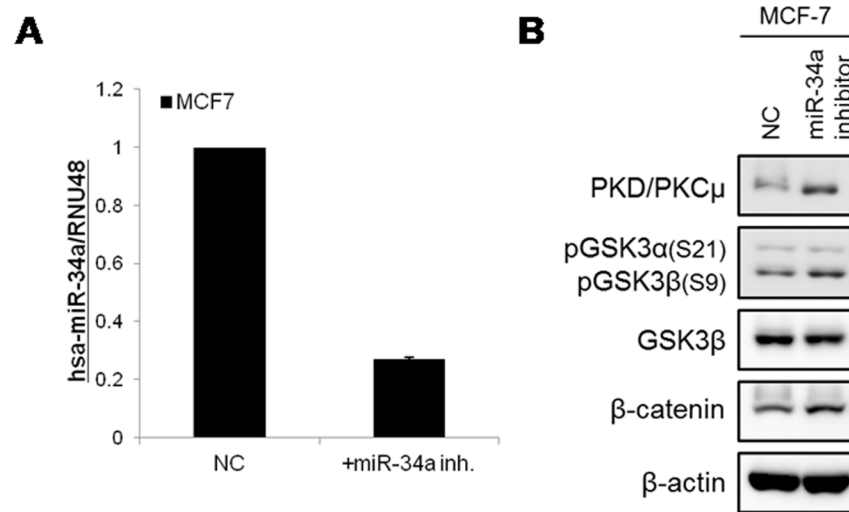
SUPPLEMENTARY FIGURES



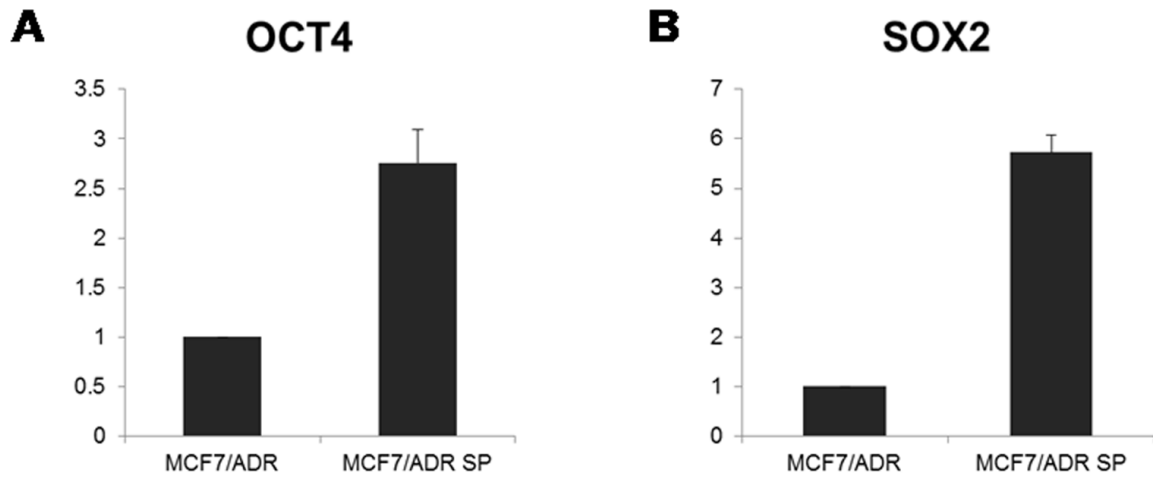
Supplementary Figure S1: No correlation with *PRKD1* expression and miR-34b or miR-34c expression. A-B. Expression levels of miR-34b and miR-34c in various breast cancer cell lines were confirmed by qRT-PCR.



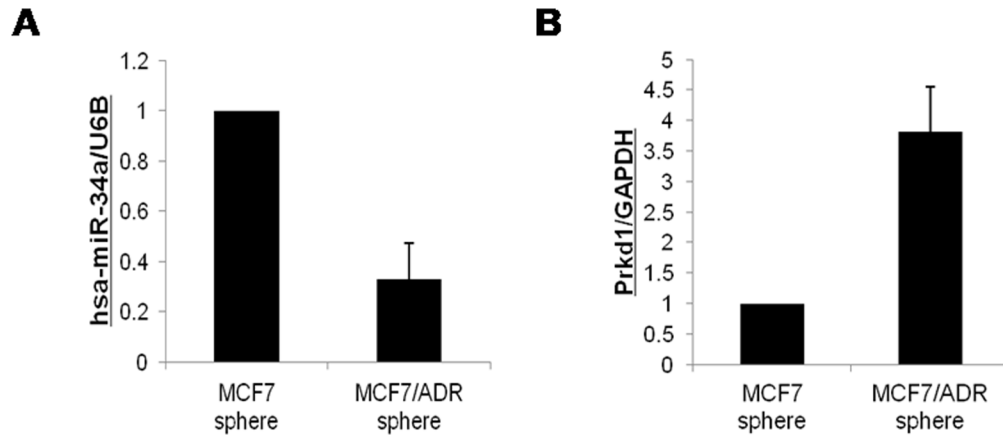
Supplementary Figure S2: Comparison of three different *PRKD1* siRNAs' efficiency.



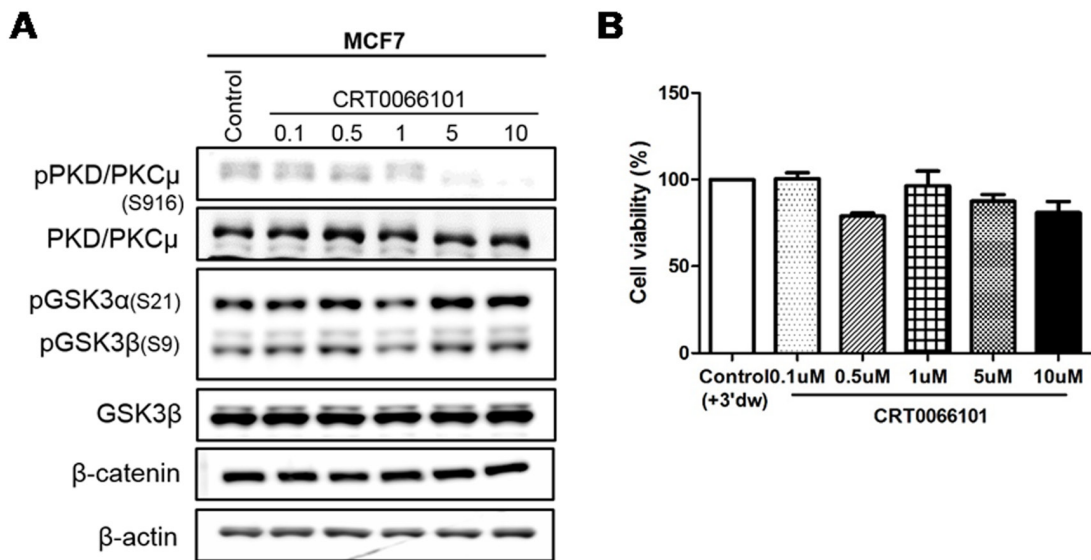
Supplementary Figure S3: Effects of suppressed miR-34a expression in MCF-7 cells. A. Following transfection with miR-34a inhibitors in MCF-7 cells, the expression level of miR-34a were confirmed by qRT-PCR and B. GSK3/ β -catenin signaling were determined by western blot analysis. β -actin was used as the loading control.



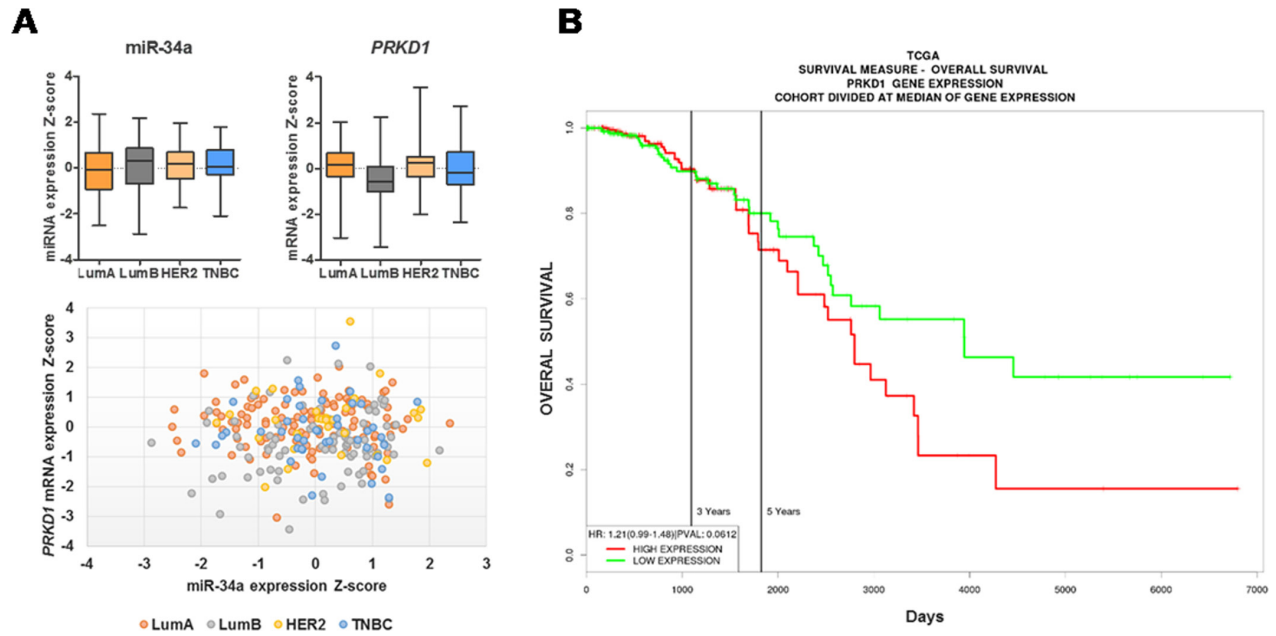
Supplementary Figure S4: The expression level of cancer stemness markers in MCF-7-ADR cells. A, B. Expression levels of *OCT4* and *SOX2* were checked by qRT-PCR.



Supplementary Figure S5: The expression level of miR-34a and *PRKD1* in MCF-7 and MCF-7-ADR mammospheres. A. miR-34a was highly expressed in MCF-7 cells in sphere status, and B. *PRKD1* was highly expressed in MCF-7-ADR mammospheres.



Supplementary Figure S6: No alteration of GSK3/ β -catenin signaling and cell viability by treatment of CRT0066101 in MCF-7 cells. **A.** Western blot analysis revealed the effects of CRT0066101 in MCF-7 cells. **B.** WST-8 assay following CRT0066101 treatment (0.1–5 μ M) in MCF-7 cells.



Supplementary Figure S7: Upregulated *PRKD1* expression in correlation with poor prognosis in breast cancer patients.
A. No correlation between miR-34a and *PRKD1* expression in diverse cell lines **B.** Overall survival according to level of *PRKD1* expression in TCGA clinical data sets.