

Supplementary Table 5. Gene expression analysis of select pluripotency genes in CD44+/CD24– and CD44+/CD24+ MCF-7-ADR populations*

Assay identification	Gene	Fold change (95% CI)	P†
Hs00742896_s1	<i>POU5F1</i>	0.81 (0.61 to 1.00)	.29
Hs00602736_s1	<i>SOX2</i>	0.88 (0.16 to 1.60)	.31
Hs00162669_m1	<i>TERT</i>	0.74 (0.40 to 1.07)	.17
Hs01062014_m1	<i>NOTCH1</i>	0.89 (0.67 to 1.11)	.16
Hs00702808_s1	<i>LIN28</i>	1.01 (0.67 to 1.35)	.50
Hs02387400_g1	<i>NANOG</i>	0.77 (0.24 to 1.30)	.17
Hs00358836_m1	<i>KLF4</i>	0.89 (0.88 to 0.90)	.35

*The delta-delta cycle threshold method was used to determine the fold change of gene expression in CD44+/CD24– MCF-7-ADR cells compared with CD44+/CD24+ MCF-7-ADR cells. CI = confidence interval.

† Statistical significance was determined with unpaired Student *t* tests. *P* values less than .05 were considered statistically significant. All statistical tests were two-sided. Means and 95% CIs are reported.