Additional File 1. Normalized MAO-A mRNA counts from Nanostring nCounter analysis.

Cell Lines	Adherent	Tumorspheres
BT20	63.53	258.63
BT474	42.68	496.09
MCF-7	35.97	550.82
MDA-MB-361	118.65	320.39
MDA-MB-436	47.54	52.69
T47D	36.32	64.63
ZR75-1	39.66	71.93
MDA-MB-157	82.81	627.16
HCC1954	270.2	330.55
Brain	1566.64	

MAO-A abundance was determined from total RNA using Nanostring nCoutner and custom probe sets. mRNA read counts were normalized by subtracting negative probe counts using Nanostring nSolver software. Human Brain RNA was included as a positive control.