

Figure S1

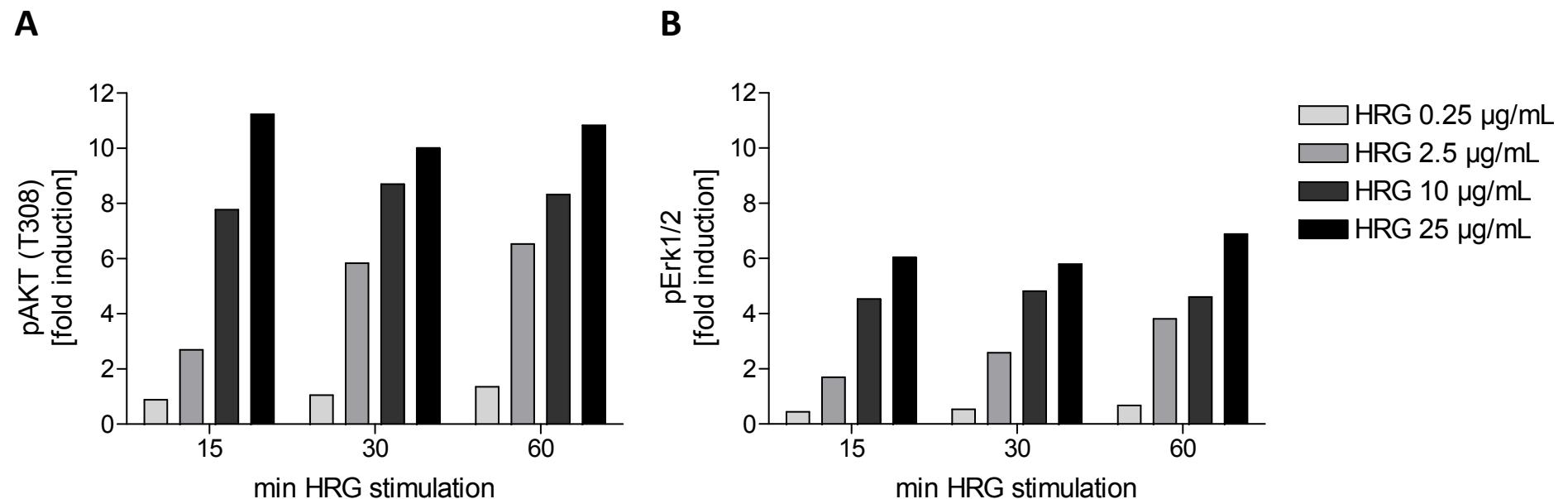


Figure S2

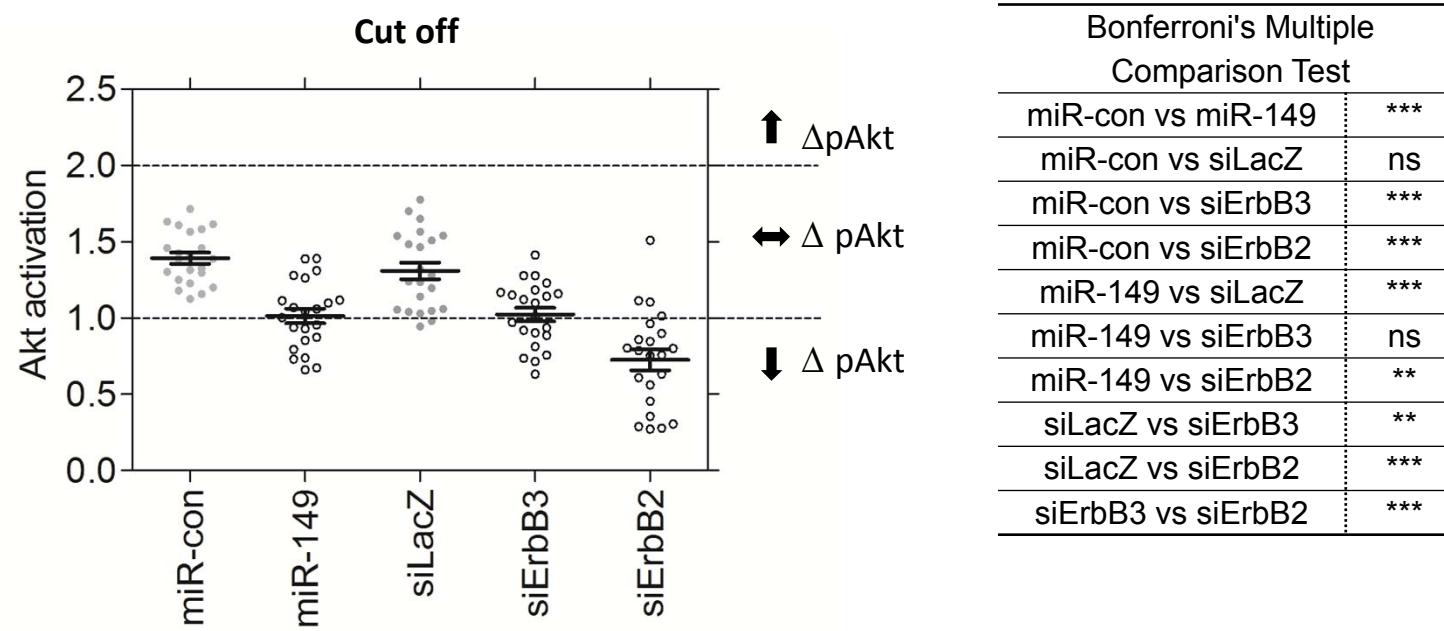


Figure S3

	miR-19a	miR-148b	miR-149	miR-155	miR-204	miR-302c	miR-326	miR-520a-3p	miR-520d-3p
SOS1	X	X			X	X	X	X	X
RPS6KA5	X	X			X	X		X	X
PDE4D	X	X	X	X	X	X		X	X
RAP1A	X	X	X	X		X		X	X
mTORC2	X	X		X	X		X		
EGFR		X	X	X	X	X		X	X
APBB1IP				X	X	X	X	X	X
ADAM17		X	X		X	X	X	X	X
CPA3	X	X		X		X		X	X
FRS2	X	X		X	X	X		X	X
CDKN1B		X	X	X		X		X	X
CALD1	X	X	X		X	X		X	X
ERBB3	X	X	X	X	X	X	X	X	X
PIK3	X	X	X	X	X	X		X	X

Figure S4

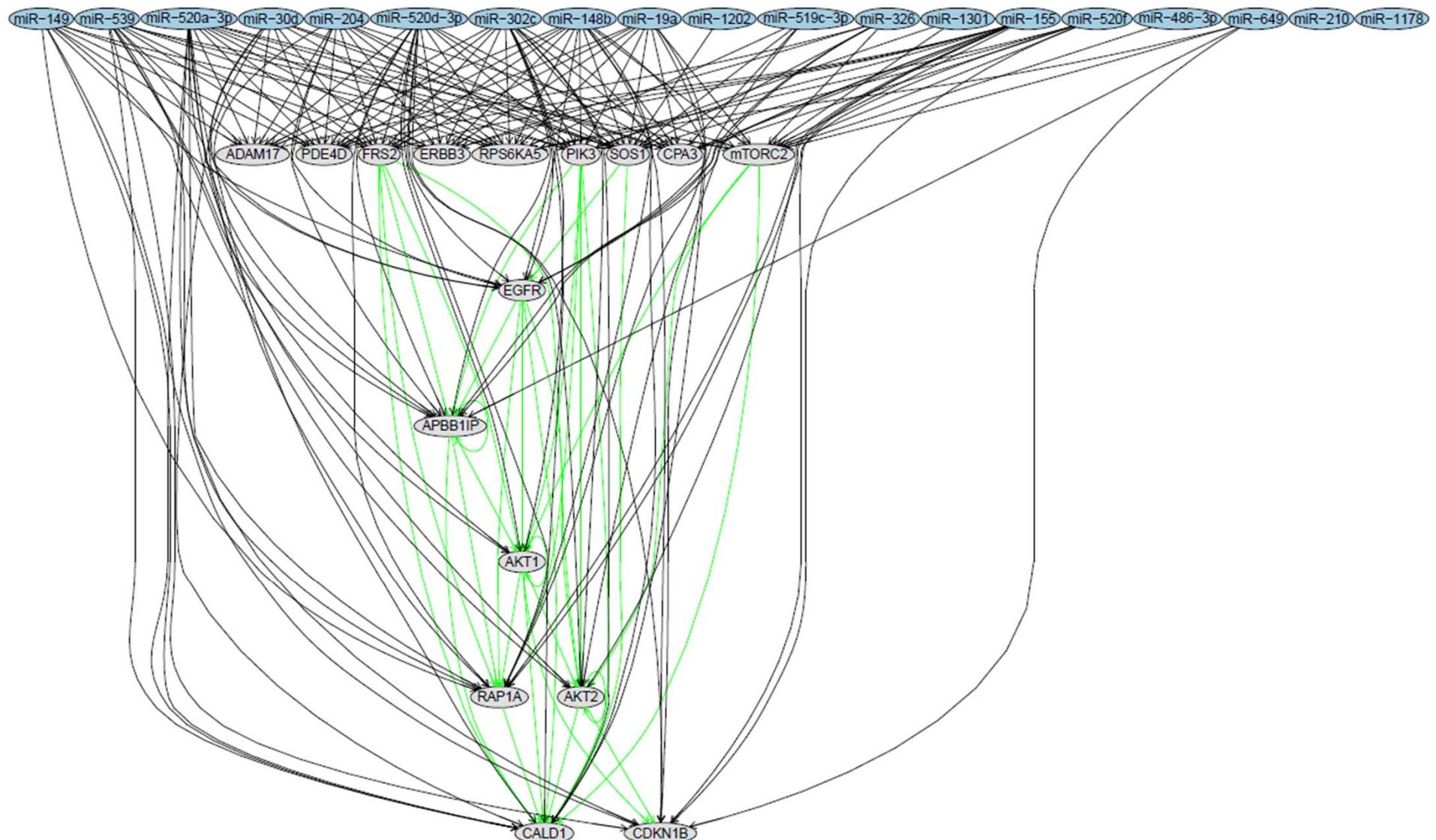


Figure S5

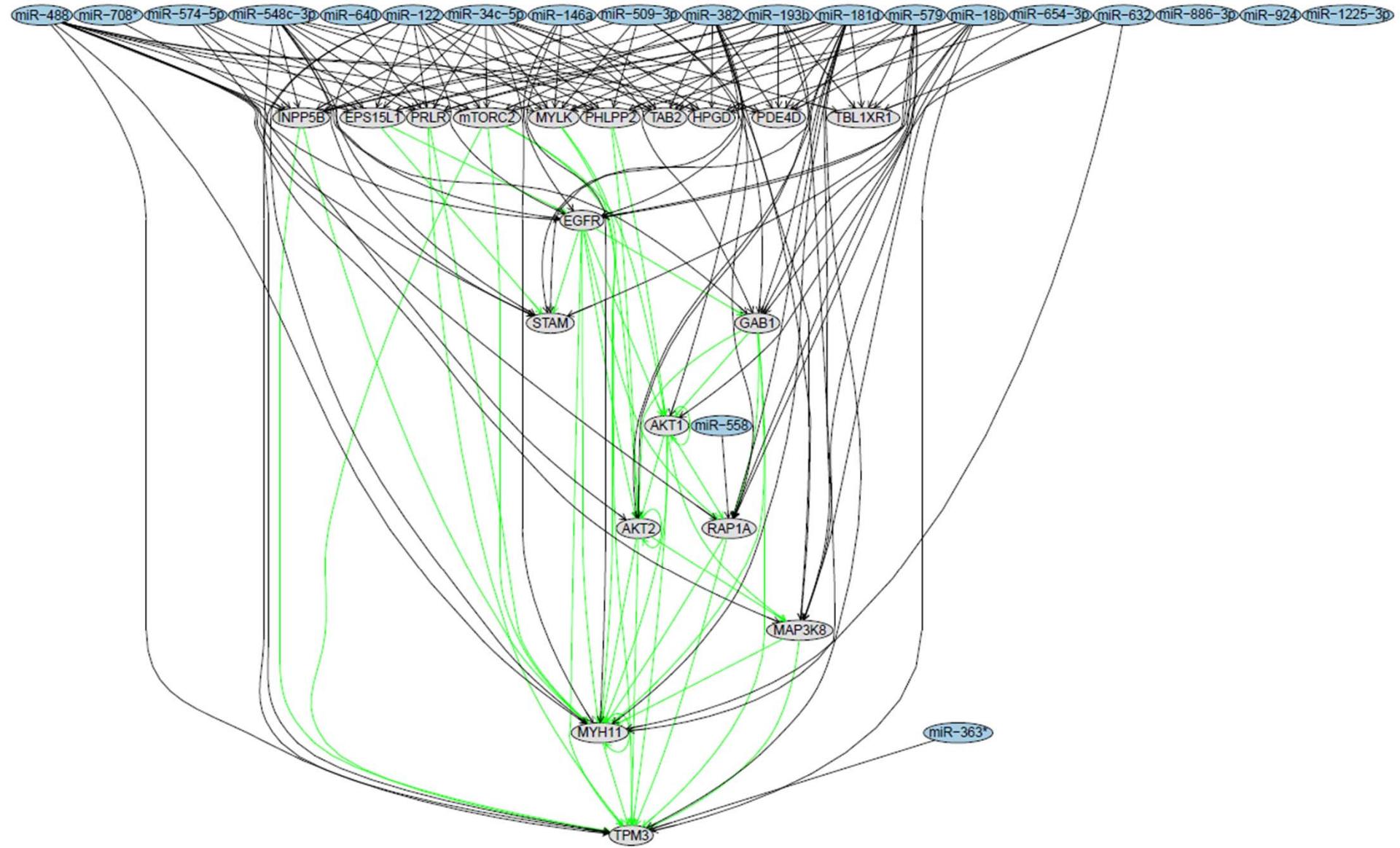


Figure S6

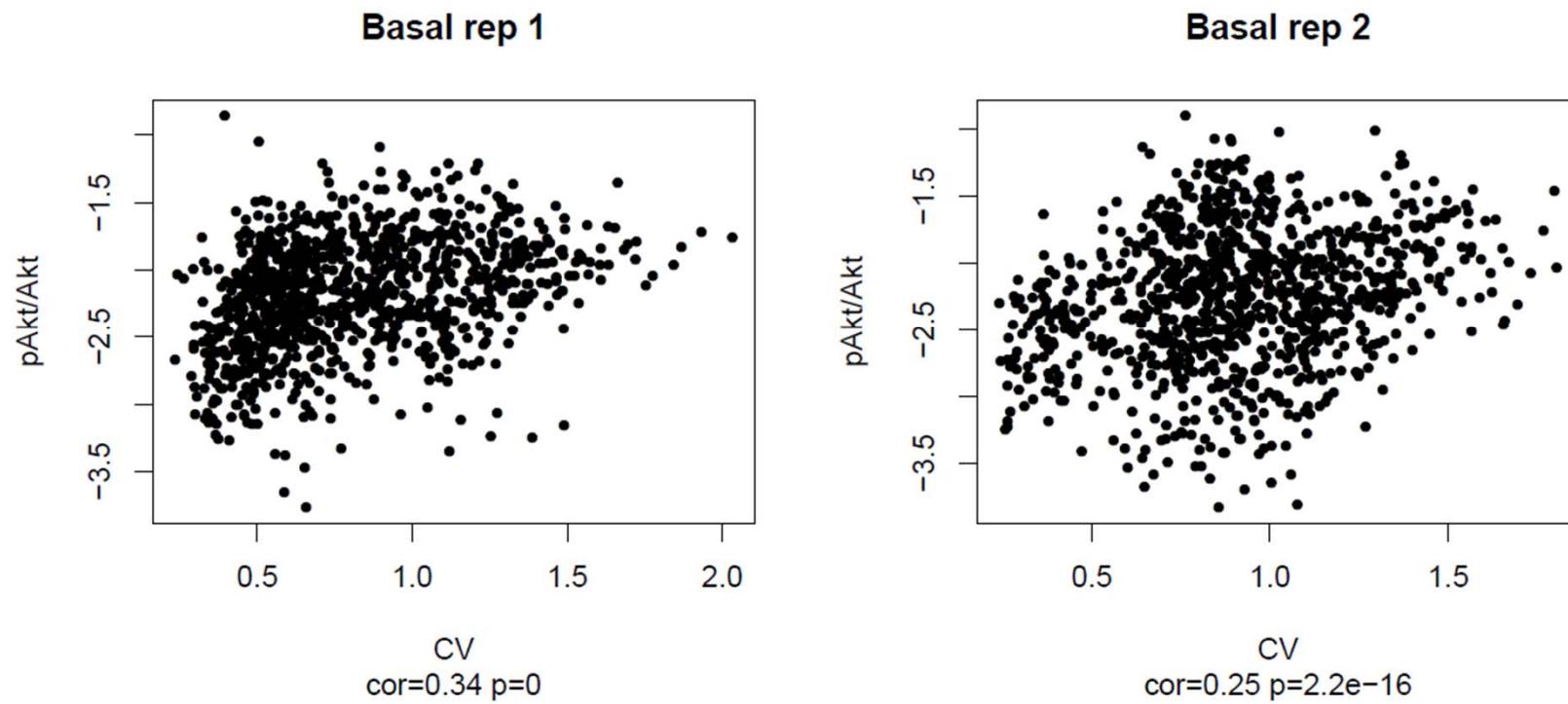


Figure S7

pGL3-ErbB3 3'UTR (Scott et al., 2007)

hsa-miR-520a-3p / ERBB3 alignment

3' ugucaGGUUUCCUUCGUGAAa 5' hsa-miR-520a-3p
500: 5' uacuaUCAUAAUUCAGCACUUa 3' ERBB3

3' ugucaggguuuuccuUCGUGAAa 5' hsa-miR-520a-3p
650: 5' augccuguaaucucAGCACUUu 3' ERBB3

ErbB3 mRNA (NM_001982)

hsa-miR-148b / ERBB3 alignment

3' uguuucAAG-AC-ACUACGUGACu 5' hsa-miR-148b
5366:5' cacuguUUCUUGUUUUUGCACUGa 3' ERBB3

hsa-miR-326 / ERBB3 alignment

3' gaccUCC-UUCCCGGGUCUCC 5' hsa-miR-326
4279:5' cccaAGGCUAUAUGCCCAGAGa 3' ERBB3

Figure S8

