Pathway	AKT	Beta-Catenin	E2F1	E2F2	E2F3	E2F4	EGFR	Myc	p53	p63	Ras	RB ko	RhoA	Src	TGFB	TNFA
Genes	140	50	79	102	48	56	407	233	182	230	120	338	100	63	423	84
Metagenes	3	3	2	2	2	2	2	2	2	2	3	2	2	2	2	2
# Burn-in for MCMC	1000	1000	1000	1000	2000	1000	1000	2000	1000	1000	2000	1000	1000	1000	2000	2000
# Iterations for MCMC	5000	5000	5000	5000	5000	3000	5000	5000	5000	5000	5000	5000	5000	5000	5000	5000
# Skips for MCMC	1	2	1	1	2	1	2	2	1	1	2	1	1	2	2	2
# Percent credible interval	95	95	95	95	95	95	95	95	95	95	95	95	95	95	95	95
# Date format: 0==unlogged , 1==logged	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
# Quantile-normalize arrays: 0==NO , 1==YES	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0
# Standardize genes: 0==NO , 1==YES	0	0	0	0	0	0	1	0	0	1	0	0	1	0	0	0
# Version: 1==train+valid , 2==train alone	1	0	1	2	1	0	1	2	2	1	1	1	2	2	0	2
# Do cross validation: 0==NO , 1==YES	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
# Make plots: 0==NO , 1==YES	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
Binreg Software	2	1	2	2	2	1	2	2	2	2	2	2	2	2	1	2