

Supplemental Table 4. BTBR quantitative RT-PCR data. Folds are AdCMV-*FOXM1b* versus AdCMV- $\beta$ -gal treatments (N=4).

	<b>Fold</b>	<b>Standard Error</b>	<b>p-value</b>
Cyclin A2	7.41	0.42	0.0001
Cyclin B1	4.21	0.73	0.0041
Cyclin B2	3.70	1.30	0.1173
Cyclin D1	0.92	0.12	0.4884
Cyclin D2	0.77	0.07	0.0612
Cyclin D3	1.19	0.19	0.4904
Cyclin E1	14.23	2.53	0.0008
Cyclin E2	5.63	0.50	0.0004
Cdk1	3.62	0.54	0.0041
Cdk2	5.34	1.72	0.0193
Cdk4	1.15	0.21	0.6215
Cdk6	1.43	0.30	0.2712
p21	1.47	0.40	0.7383
p27	1.34	0.45	0.7640
p57	1.42	0.44	0.5989
p15	1.09	0.39	0.8434
p16	4.19	1.74	0.5315
p18	1.13	0.09	0.2374
p19	1.80	0.77	0.6712
Skp2	2.18	0.35	0.0200
KIS	0.74	0.12	0.1288
Cks1b	2.04	0.33	0.0269
Cdc25a	1.34	0.20	0.2327
PCNA	2.62	0.39	0.0144
Cdc25b	1.43	0.19	0.1334
Cdc25c	20.72	10.32	0.0067
Cks2	14.07	3.04	0.0014
Birc5	29.34	8.67	0.0066
Aurkb	14.73	6.27	0.0157
Plk1	5.37	1.14	0.0039
Cdca8	11.75	1.81	0.0005
Cenpa	1.24	0.37	0.7631
Cenpb	0.96	0.15	0.6362
Cenpe	5.74	2.25	0.0276
Cenpf	6.17	1.98	0.0373
Cdc20	7.53	1.61	0.0023
Cdca2	3.92	0.75	0.0075
Nek2	11.66	3.38	0.0035
Kif20a	4.49	0.30	0.0002
Mki67	5.11	0.45	0.0004