

Supplemental Table 5. Human quantitative RT-PCR data. Folds are AdCMV-*FOXM1b* versus AdCMV- β -gal treatments (N=4-5).

	Fold	Standard Error	p-value
Cyclin A1	2.84	0.66	0.0213
Cyclin A2	3.22	0.97	0.0392
Cyclin B1	2.43	0.51	0.0215
Cyclin B2	2.48	0.54	0.0346
Cyclin D1	1.04	0.20	0.9131
Cyclin D2	0.78	0.11	0.1369
Cyclin D3	0.89	0.14	0.4023
Cyclin E1	5.68	1.58	0.0170
Cyclin E2	1.97	0.24	0.0115
Cdk1	6.14	1.90	0.0210
Cdk2	1.85	0.28	0.0235
Cdk4	0.89	0.14	0.4176
Cdk6	0.90	0.13	0.4343
p21	0.96	0.33	0.5733
p27	0.85	0.12	0.2851
p57	0.93	0.31	0.5134
p15	0.89	0.24	0.4450
p16	0.84	0.14	0.3158
p18	1.42	0.32	0.2563
p19	1.46	0.19	0.0787
Skp2	0.91	0.12	0.4220
KIS	0.99	0.16	0.7695
Cks1b	1.41	0.10	0.0140
Cdc25a	3.89	1.03	0.0136
PCNA	2.35	0.43	0.0262
Cdc25b	2.02	0.37	0.0455
Cdc25c	15.07	8.62	0.0178
Cks2	4.30	0.84	0.0068
birc5	2.97	0.78	0.0408
Aurkb	4.85	1.82	0.0210
Plk1	2.28	0.55	0.0452
Cdca8	3.22	0.31	0.0015
Cenpa	1.76	0.74	0.0451
Cenpb	0.98	0.12	0.7472
Cenpe	6.22	1.48	0.0065
Cenpf	2.66	0.69	0.0264
Cdc20	4.40	1.16	0.0126
Cdca2	3.00	0.94	0.0451
Nek2	3.51	0.74	0.0160
Kif20a	3.28	0.37	0.0023
Mki67	1.55	0.28	0.1958