

1 **Impact of Human Cytomegalovirus on Glioblastoma Cell Viability and**  
2 **Chemotherapy Treatment**

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19

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24 Abbreviations: BCNU, Carmustine (bis-chloroethylnitrosourea); BER, base excision  
25 repair; cmvIL10, HCMV interleukin 10 homolog; DMSO, dimethyl sulfoxide; DSBs,

26 DNA double-strand breaks; EtOH, ethanol; FACS, fluorescence-activated cell sorting;  
27 GBM, glioblastoma multiforme; HCMV, Human cytomegalovirus; hpi, hours post  
28 infection; MGMT, O6-methylguanine DNA methyltransferase; MTT, 3-(4,5-  
29 dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; STAT3, signal transducer and  
30 activator of transcription 3; TMZ, Temozolomide; PBS, phosphate-buffered saline; PI,  
31 propidium iodide.

32 **ABSTRACT**

33

34 The relationship between the Human Cytomegalovirus (HCMV) and tumors has been  
35 extensively investigated, mainly in glioblastoma multiforme (GBM), a malignant tumor  
36 of the central nervous system with low overall survival. Several reports demonstrated  
37 the presence of HCMV in GBM, although typically restricted to a low number of cells,  
38 and studies indicated that viral proteins have the ability to dysregulate cellular  
39 processes and increase tumor malignancy. Treatment of GBM involves the use of the  
40 chemotherapeutic agents Temozolomide (TMZ) and Carmustine (BCNU), which lead  
41 to attachment of adducts to the DNA backbone, causing errors during replication and  
42 consequent cell death. It is known that HCMV infection can modulate DNA repair  
43 pathways, but the effects the virus may exhibit during chemotherapy are unknown.  
44 Here we approach this question by analyzing HCMV infection and viral protein  
45 accumulation in GBM cell lines with different genotypes and their response to TMZ and  
46 BCNU in the presence of the virus. We demonstrate that A172, TP365MG and  
47 U251MG GBM cells are efficiently infected by both low passage (TB40E) and high  
48 passage (AD169) HCMV strains. However, the GBM cell lines vary widely in  
49 permissiveness to viral gene expression and exhibit very different patterns of  
50 immediate early, early and late protein accumulation. HCMV reduces viability of  
51 permissive GBM cells in a multiplicity-dependent manner both in the absence and  
52 presence of TMZ or BNCU. In sum, we demonstrate that GBM cell lines are equally  
53 susceptible but differentially permissive to infection by both low and high passage  
54 strains of HCMV. This observation not only indicates that viral replication is largely  
55 controlled by cellular factors in this system, but also provides a possible explanation  
56 for why viral gene products are found in only a subset of cells in GBM tumors.

57 Furthermore, we conclude that the virus does not confer increased resistance to  
58 chemotherapeutic drugs in various GBM cell lines but instead reduces tumor cell  
59 viability. These results highlight that the oncomodulatory potential of HCMV is not  
60 limited to cancer promoting activities but also comprises adverse effects on tumor cell  
61 proliferation or survival.

62

## 63 INTRODUCTION

64

65 Human cytomegalovirus (HCMV) is a member of the herpesvirus family and is  
66 present in the majority of people worldwide, including almost 100% of the adult  
67 population in developing countries [1]. Congenital infection by HCMV is the most  
68 common cause of severe neurological impairments including hearing loss, optic  
69 atrophy and microcephaly in many parts of the world [1]. The virus is also a life-  
70 threatening pathogen in immunocompromised individuals such as transplant recipients  
71 as well as patients with AIDS or cancer [2].

72 HCMV has been detected in many tumor types [3–12] , and the relationship  
73 between the virus and cancer has been investigated most extensively in glioblastoma  
74 multiforme (GBM) [13–16]. GBM is the most aggressive primary brain tumor with a  
75 median survival of 15 months and an unknown etiology [17]. GBM is characterized by  
76 the presence of a heterogeneous population of cells that are infiltrative, angiogenic and  
77 poorly metastatic [18]. Various reports demonstrated the presence of HCMV DNA,  
78 transcripts and/or proteins in GBM samples (reviewed in [16]). In addition, a number  
79 of studies showed that viral proteins can modulate the phenotype of GBM cells leading  
80 to cell proliferation [19], cell immortalization [20], cell migration [21, 22], stem cell  
81 maintenance [23, 24], angiogenesis [22, 25–27] and immunomodulation [28]. Although  
82 HCMV is not considered to be oncogenic, it is believed that viral proteins may increase  
83 cellular malignancy and contribute to tumor progression, a phenotype called  
84 oncomodulation [29]. In fact, studies suggest that extracellular factors released during  
85 HCMV infection can cause immunosuppression of the tumor microenvironment  
86 (reviewed in [30] ), but the virus may also be more directly involved in the malignant  
87 phenotype of tumor cells (reviewed in [31]).

88           The standard GBM treatment is surgical resection followed by radiotherapy and  
89 chemotherapy. The cytotoxic treatment with Temozolomide (TMZ) and Carmustine  
90 (bis-chloroethylnitrosourea, BCNU) inserts alkyl adducts into the DNA backbone,  
91 causing errors during DNA replication due the formation of numerous DNA double-  
92 strand breaks (DSBs) not repaired by the cellular machinery. This lack in DNA repair  
93 ultimately leads to cell death [32]. Despite the standard treatment, less than 5% of  
94 GBM patients live longer than 5 years after diagnosis (reviewed in [33]), mainly  
95 because of the acquired and intrinsic drug resistance (reviewed in [34]). The main  
96 mechanism of chemotherapy resistance is mediated by O6-methylguanine DNA  
97 methyltransferase (MGMT), a suicide repair protein that removes DNA adducts formed  
98 at the O6-position of guanidine in response to alkylating agents, such as TMZ. MGMT  
99 expression is associated with poor patient response to alkylating therapies. Other  
100 molecular mechanisms and proteins involved in cell resistance to DNA damage are  
101 base excision repair (BER), ataxia-telangiectasia mutated (ATM) and ataxia  
102 telangiectasia and Rad3-related (ATR) (reviewed in [34]).

103           Studies demonstrated that HCMV upregulates the host BER machinery [35],  
104 activates the DNA damage response [36], and significantly reduces cyclobutane  
105 pyrimidine dimers in UV-exposed viral DNA [37], indicating a viral role in inducing DNA  
106 repair pathways that may limit DNA damage. On the other hand, HCMV was also found  
107 to cause DNA damage. For example, the HCMV protein UL76 was demonstrated to  
108 induce chromosomal aberrations [38], and the virus has been associated with specific  
109 damage at several loci in chromosome 1, 1q21, 1q23 and 1q42 [39, 40]. These  
110 findings lead to the speculation that the virus might either increase or decrease cell  
111 resistance to DNA damaging chemotherapeutic drugs.

112           To investigate these possibilities, we analyzed infection by low and high  
113 passage strains of HCMV in different GBM cell lines and their response to  
114 chemotherapy treatment in the presence and absence of the virus.

115 **RESULTS**

116

117 **GBM cell lines are generally susceptible to HCMV infection but not equally**  
118 **permissive to viral replication**

119

120 We and others have demonstrated that HCMV DNA and proteins are present in  
121 GBM tumors [14, 15]. However, few cells in the tumor bulk seem to contain viral DNA,  
122 transcripts and/or proteins [14], indicating that HCMV infection or replication is not  
123 equally efficient in all tumor cells. We therefore sought to investigate viral infection and  
124 replication using GBM cell lines and HCMV strains with different genetic backgrounds.

125 The GBM cell lines A172, TP365MG and U251MG were exposed to a low  
126 passage (TB40E) and a high passage (AD169) HCMV strain at MOI 3, and the  
127 presence of intracellular pp65 protein, a major component of HCMV particles, was  
128 evaluated at 2 h post infection (hpi) by confocal immunofluorescence microscopy. The  
129 pp65 associated with the input virus was detected in the vast majority (>90%) of cells  
130 in all infections, indicating no significant difference in virus entry among the different  
131 cell lines or HCMV strains (Fig. 1).

132 To examine the extent and temporal pattern of HCMV protein accumulation  
133 following infection, viral protein markers for the immediate-early (IE1-72kDa), early  
134 (pp52 also known as pUL44) and late (pp28) stages of viral replication were analyzed  
135 at different times post infection by immunofluorescence. Very different patterns of  
136 expression were observed in the three cell lines (Fig. 2 and 3). Controls are shown in  
137 Figures S1 and S2. The IE1-72kDa protein was detectable in only a small subset  
138 (<20%) of A172 cells at 16 hpi, and the proportion of cells staining positive for this  
139 protein further decreased over time (30–96 hpi). Neither pp52 nor pp28 proteins were  
140 detected at any time point evaluated in this cell line. In contrast, IE1-72kDa was

141 observed in 40% to >70% of TP365MG and U251MG cells at 16 hpi, and up to almost  
142 100% of these cells stained positive for IE1-72kDa at later times post infection.  
143 However, pp52 and pp28 protein accumulation differed markedly in the two cell lines.  
144 While most of the TP365MG cells had accumulated pp52 and pp28 by 96 hpi, the two  
145 proteins were detectable in only a small proportion of U251MG cells at this time.

146 Together the results show that all GBM cell lines tested are highly and equally  
147 susceptible to low and high passage strains of HCMV. However, the temporal pattern  
148 of immediate-early, early and late viral protein accumulation varies widely between the  
149 cell lines suggesting major differences in permissivity to HCMV replication determined  
150 by the cellular genetic background.

151

## 152 **HCMV replication reduces the number of viable GBM cells but does not increase** 153 **tumor cell resistance to TMZ or BCNU**

154

155 Previous studies have shown that HCMV can augment or inhibit cellular DNA  
156 damage responses. Therefore, we aimed to evaluate the effect of HCMV infection on  
157 chemotherapy treatment of GBM cell lines with the DNA damaging drugs TMZ or  
158 BCNU.

159 Initially, viability of non-infected A172, TP365MG and U251MG cells in the  
160 presence of 25, 50 and 100  $\mu$ M TMZ or BCNU for 96, 120 and 144 h was assessed by  
161 MTT assay. Compared to controls, both drugs caused a marked decrease of in the  
162 number of viable cells in a concentration-and time-dependent manner in all three cell  
163 lines (Fig. 4). However, the timing and extent by which cell viability was reduced varied  
164 with different combinations of drugs and cell lines. Further analyses were performed  
165 with drug concentrations that decrease cell viability by at least 30%, namely 50  $\mu$ M

166 TMZ or BCNU for A172 and TP365MG cells and 75  $\mu$ M TMZ or 35  $\mu$ M BCNU for  
167 U251MG cells.

168 Subsequently, GBM cells were exposed to HCMV AD169 or TB40E at different  
169 MOIs (0.12, 0.6 and 3) or mock infection overnight and treated with TMZ, BCNU or  
170 solvent (DMSO or EtOH, respectively) for 96 h. Infection with either virus strain lead to  
171 a marked decline in the number of TP365MG and U251MG viable cells, even in the  
172 absence of drug. The reduction in the number of viable TP365MG cells was MOI-  
173 dependent for both AD169 and TB40E. However, only AD169 but not TB40E reduced  
174 the number of viable U251MG cells in an MOI-dependent fashion. As expected, TMZ  
175 and BCNU reduced cell viability in all cell lines in the absence of virus.

176 In U251MG cells, high MOI infection with AD169 and treatment with BCNU  
177 appeared to work additively in reducing the number of viable cells. Similar additive  
178 effects were less pronounced or absent upon treating AD169-infected U251MG cells  
179 with TMZ or infecting these cells with TB40E. Likewise, BCNU and TMZ did no exert  
180 significant additive effects when combined with AD169 or TB40E infection of TP365MG  
181 cells, most likely due to the pronounced reduction in the number of viable cells caused  
182 by HCMV alone in this system. Finally, there was no significant impact of HCMV on  
183 A172 cell viability, and no obvious additive effects between the viruses and drugs were  
184 observed in these cells (Fig. 5).

185 These results show that HCMV infection adversely affects viability of permissive  
186 GBM cells, but does not increase chemoresistance of the cell lines tested. Instead,  
187 HCMV may even increase susceptibility to BCNU under certain conditions.

188

189 **Intracellular presence of HCMV is required to reduce GBM cell viability**

190

191 It has been demonstrated that HCMV infection leads to the production and  
192 secretion of numerous cytokines some of which may adversely affect cell viability [41].  
193 For example, the HCMV interleukin 10 homolog (cmvIL10) has been shown to mediate  
194 local tumor immunosuppression and activation of signal transducer and activator of  
195 transcription 3 (STAT3), a transcription factor associated with chemoresistance in  
196 ovarian cancer [42]. Our previous cell viability analyses were performed in *de novo*  
197 infected cells, suggesting that viral gene expression or replication caused the  
198 substantial reduction in the number of viable cells. However, this approach may mask  
199 possible indirect effects the virus may exert on the viability of uninfected cells by  
200 causing the release of soluble factors.

201 To test the possibility that soluble factors released from infected cells contribute  
202 to reduction in the number of viable cells even in the absence of infection A172,  
203 TP365MG and U251MG cells were inoculated with supernatants from MRC-5 cells  
204 infected with HCMV AD169 or TB40E for 48 h, and treated with the two  
205 chemotherapeutic drugs for 96 h. Expression of cmvIL10 in infected MRC-5 cells was  
206 confirmed by Western blotting (data not shown). Notably, only A172 cells treated with  
207 supernatants from AD169- or TB40E-infected cells and treated with TMZ exhibited a  
208 significant decrease in number of viable cells. This phenotype was not observed with  
209 any other combination of cells, viruses and drugs (Fig. 6).

210 These results confirm that reduced number of viable GBM cells caused by  
211 HCMV infection largely depends on the presence of the virus in the affected cells and  
212 likely results from viral gene expression and replication.

213

214 **HCMV infection reduces GBM cell proliferation and induces tumor cell death**

215

216 To further explore the impact of HCMV on GBM cell viability, we determined the  
217 amount of cell death in U251MG cells by fluorescence-activated cell sorting (FACS).  
218 U251MG cells were infected at MOI 3 overnight, and treated with low concentrations  
219 (25  $\mu$ M) of TMZ or BCNU for 96 h. U251MG cells were chosen for this experiment due  
220 to the less pronounced decrease in cell viability upon viral infection compared to  
221 TP365MG cells, and the low drug concentration was used to limit cell debris during the  
222 analysis. IE1-72kDa was used as a marker for viral infection. Only a small proportion  
223 (<10%) of cells were positive for both propidium iodide (PI) and IE1-72kDa (Fig. 7a),  
224 suggesting that most of the infected cells were still viable. Further cell death analysis  
225 by trypan blue exclusion assay performed in TP365MG and U251MG cells confirmed  
226 a reduction in infected cell numbers when compared with controls. However, the  
227 percentage of viable cells in this assay proved to be higher than that obtained in the  
228 metabolic assays (Fig. 7b and Fig. 5).

229 These findings indicate that the reduced number of viable cells observed in  
230 HCMV-infected GBM cell lines is largely due to decreased cell proliferation, and that  
231 cell death makes a minor contribution to the overall effect.

232 **DISCUSSION**

233

234 Glioblastoma is a highly heterogeneous tumor, with genetic diversity existing  
235 both among patients and within individual tumors [43]. The intratumoral genetic  
236 heterogeneity can result in many phenotypes including populations of drug resistant  
237 subclones that are the main cause of treatment failure. HCMV is an infectious agent  
238 detected in GBM samples worldwide, although usually only in a subset of tumor cells  
239 [14]. There are also studies that failed to detect HCMV in GBM [44–46]. The causes  
240 underlying the presence or absence of HCMV in GBM cells and the consequences  
241 resulting from infection are not well understood. It has been demonstrated that HCMV  
242 can mediate DNA damage, but the virus can also induce DNA repair pathways [36,  
243 37], phenotypes that could interfere in GBM chemotherapeutic treatment.

244 Our results demonstrate that three different GBM cell lines (A172, TP365MG  
245 and U251MG) are equally susceptible to HCMV infection. This conclusion is based on  
246 the intracellular presence of the major viral tegument component (pp65) in a similar  
247 number of cells at the beginning of infection. However, the permissivity of the cell lines  
248 to the temporal accumulation of HCMV proteins differs widely, again in a manner  
249 independent of the virus strain. While A172 cells undergo an infection that seems to  
250 be silenced or aborted at the immediate-early stage, TP365MG and U251MG cells are  
251 permissive to viral early and late gene expression. However, infection progresses more  
252 rapidly in TP365MG compared to U251MG cells. High passage (AD169) and low  
253 passage (TB40E) HCMV strains showed very similar phenotypes in terms of both  
254 infectivity and viral protein accumulation in the tested cell lines. These results suggest  
255 that the permissivity of GBM cells to HCMV replication is determined by cellular rather  
256 than viral factors. In fact, the three cell lines under investigation have different  
257 backgrounds with respect to important genes including CDKN2A, PTEN, p16, and p53

258 [47–50]. Further studies are necessary to investigate the significance of mutations in  
259 these and other genes for permissivity to HCMV gene expression and replication in  
260 both cell lines and tumor tissue. Although we did not perform experiments in tumor  
261 tissue, the cell lines used in this work have been commonly used as experimental  
262 models in GBM research. Our results are compatible with the general view that HCMV  
263 is present in a limited number of cells in GBM tumor samples [14] implying that genetic  
264 differences between cells may result in different outcomes of infection.

265 Treatment of all three cell lines with TMZ or BCNU causes significant reduction  
266 in the number of viable cells in a dose-and time-dependent manner. In TP365MG cells  
267 both drugs caused a similar decrease in the number of viable cells. In contrast,  
268 U251MG cells treated with BCNU exhibited a decline in the number of viable cells of  
269 almost 100% at 144 h, a phenotype not observed with TMZ. It has been previously  
270 shown that BCNU induces cell death by necrosis and apoptosis in U138MG cells, a  
271 GBM cell line which carries mutant p53 and is therefore more susceptible to toxicity by  
272 BCNU than TMZ [51]. These findings are consistent with our own results, since  
273 U251MG cells carry mutant p53. However, the same phenotype is not observed in  
274 TP365MG cells which also have mutant p53, indicating that drug sensitivity is not  
275 exclusively linked to p53 status [52]. Notably, viral infection alone reduced the number  
276 of viable U251MG and TP365MG cells, which are both permissive to HCMV early and  
277 late gene expression, in a MOI-dependent manner. This phenotype was particularly  
278 pronounced in TP365MG cells, most likely due to their high levels of HCMV gene  
279 expression. No major effects were observed in A172 cells where viral protein  
280 accumulation is limited to immediate-early.

281 The concomitant presence of HCMV and TMZ or BCNU caused a further  
282 reduction in the number of viable cells. Interestingly, the reduction in the number of

283 viable TP365MG cells was MOI-dependent for both AD169 and TB40. However, only  
284 AD169 but not TB40E reduced the number of viable U251MG cells in an MOI-  
285 dependent fashion. This difference could reflect the genetic differences in the virus  
286 strains used in this study, such as the genes from the UL-b' region of the genome,  
287 which has been deleted from AD169 but are present in TB40E [53].

288 Overall, our results indicate that HCMV replication occurs differentially in GBM  
289 cell lines depending on their genetic background and that the adverse effects HCMV  
290 confers on GBM cells is likely caused mainly by reduction of cell proliferation, as shown  
291 by FACS and trypan exclusion assay, although cell death also contributes.

292 The effects HCMV may exert during chemotherapy treatment of GBM was  
293 previously investigated in several cell types including astrocytoma cells, breast cancer  
294 cells, neuroblastoma cells, peripheral blood lymphocytes and smooth muscle cells, but  
295 the results are controversial. HCMV-infected astrocytoma cells showed a decrease in  
296 apoptosis when treated with cisplatin, a phenotype also observed in etoposide-treated  
297 MDA-MB-231 breast cancer cells [54–57]. Likewise, UKF-NB-2A-AD169 cells, a  
298 neuroblastoma cell line persistently infected with HCMV, were more resistant to  
299 cisplatin-induced cell death [58]. Conversely, treatment with ganciclovir re-established  
300 sensitivity to chemotherapy in neuroblastoma and astrocytoma cell lines persistently  
301 infected with HCMV [56, 58]. HCMV has also been reported to increase resistance of  
302 glioma cancer stem cells to TMZ [59]. On the other hand, HCMV has been shown to  
303 increase genetic damage induced by bleomycin in peripheral blood lymphocytes [60,  
304 61]. Interestingly, HCMV IE1-72kDa seems to stimulate while IE2-86kDa may  
305 suppress doxorubicin-induced apoptosis in smooth muscle cells [62]. We demonstrate  
306 here that HCMV does not increase resistance in three GBM cell lines. Instead, the virus  
307 causes a reduction in TP365MG and U251MG cell proliferation or survival both in the

308 absence and presence of the tested drugs. These findings highlight that HCMV not  
309 only exhibits tumor promoting potential but may also exert adverse oncomodulatory  
310 effects depending on tumor cell permissivity to viral replication. In tumor cells  
311 permissive to viral early and late gene expression, the virus may be oncostatic or  
312 oncolytic while the tumor promoting activities linked to viral immediate-early proteins  
313 may predominate in cells with restricted viral replication.

314         Many studies have been conducted to understand the relationship between  
315 HCMV and GBM, but we are still far from understanding the role the virus may have in  
316 tumor pathogenesis. Our results indicate that HCMV does not increase GBM cell  
317 resistance to TMZ and BCNU *in vitro*. However, further studies need to be done to  
318 examine whether the virus behaves similarly in other tumor cell lines and in *in vivo*.

319 **METHODS**

320

321 **Cell culture and virus production**

322

323 Human glioblastoma cell lines A172, TP365MG and U251MG were provided by  
324 Dr. Markus Riemenschneider, University of Regensburg. MRC-5 human embryonic  
325 lung fibroblast cells were obtained from the American Type Culture Collection. Cells  
326 were maintained in Dulbecco's Modified Eagle's Medium (DMEM) with 10% fetal  
327 bovine serum, 100 U/ml penicillin, and 100 µg/ml streptomycin (Thermo Fisher  
328 Scientific) at 37°C in a humidified atmosphere with 5% CO<sub>2</sub>.

329 HCMV BAD<sub>in</sub>UL99GFP [63] was produced by electroporation, and AD169 and  
330 TB40E were produced by nucleofection of MRC-5 cells with the corresponding  
331 bacterial artificial chromosome clones (provided by Dr. Thomas Shenk, Princeton  
332 University). Cell supernatants containing virus were collected after complete cell lysis  
333 and cleared of cell debris by low-speed centrifugation. Cell-free virus stocks were  
334 stored at -80°C. For virus titration, serial dilutions of stocks were plated on MRC-5 cells.  
335 At 24 hpi, cells were fixed with 4% formaldehyde, permeabilized with 0.1% Triton X-  
336 100, and blocked with 3% BSA. Cells were labeled using a hybridoma supernatant of  
337 a primary mouse antibody to IE1 (1B12) [64] diluted 1:5, and a secondary goat anti-  
338 mouse antibody conjugated to Alexa Fluor 594 (Abcam, ab150116) diluted 1:1000.  
339 Nuclei were stained with Hoechst 33342 (Abcam, ab145597). Virus titers were  
340 expressed as plaque forming units per ml.

341

342 **Infection and viral protein detection**

343

344 To determine viral protein levels during a time course, A172, TP365MG and  
345 U251MG cells were plated in low serum (1%) media at a density of 100.000 cells on  
346 glass coverslips in 24-well plates and infected with HCMV AD169 or TB40E at MOI 3  
347 overnight. After that, the inoculum was removed and replaced by fresh medium. Cells  
348 were maintained until 24, 48, 72, or 96 hpi (U251MG were also infected for 120 h) and  
349 subjected to standard or confocal epifluorescence microscopy. Briefly, cells were  
350 washed in PBS with 0.05% Tween 20 (PBS-T) and fixed with iced methanol for 15 min  
351 at -20°C. Methanol was then removed, coverslips dried, and cells blocked with 2% BSA  
352 in PBS-T for 1 h. Cells were incubated with primary antibodies mouse anti-pp65 (1:10),  
353 mouse anti-IE1-72kDa (1:200), mouse anti-pp52 (1:100) or mouse-anti pp28 (1:10)  
354 (provided by Dr. Thomas Shenk, Princeton University) for 1 h, washed and incubated  
355 with a secondary goat anti-mouse antibody conjugated to Alexa 594 (1:1000) (Abcam,  
356 ab150116). Controls included mock infected cells with HCMV specific primary  
357 antibodies and secondary antibody, mock infected cells with secondary antibody only,  
358 HCMV infected cells with secondary antibody only (Fig. S1). In addition  
359 immunofluorescence experiments were performed in mock and HCMV infected cells  
360 using a mouse monoclonal antibody against the VP1 protein of JC polyomavirus  
361 (Abcam, ab34756) and the secondary anti-mouse conjugate from the above  
362 experiments (Fig. S2).

363 Coverslips were mounted with 4',6-diamidino-2-phenylindole (DAPI). For  
364 standard fluorescence microscopy, cells were observed using an EVOS FL Cell  
365 Imaging System (Thermo Fisher Scientific) and quantified with ImageJ software using  
366 the following formula: % infected cells = number of infected cells (red) / total number  
367 of cells (blue) × 100%. For confocal fluorescence microscopy, a series of z-images  
368 were recorded using a Leica TCS SP5 confocal microscope (Leica Microsystems).

369

370 **Cell viability and cell death assays**

371

372 TMZ (4-methyl-5-oxo-2,3,4,6,8-pentazabicyclo[4.3.0]nona-2,7,9-triene-9-  
373 carboxamide) and BCNU (1,3-bis(2-chloroethyl)-1-nitrosourea) were purchased from  
374 Sigma-Aldrich, and stock solutions (100 mM) were prepared by dissolving the drugs in  
375 dimethyl sulfoxide (DMSO) or ethanol (EtOH), respectively. Solutions were aliquoted  
376 and stored at -20°C. For cytotoxicity analysis, A172, TP365MG and U251MG cells  
377 (10.000 cells/well) were incubated in triplicates overnight in 24-well plates. On the next  
378 day, the culture supernatants were replaced with fresh media containing 25, 50 or 100  
379 µM TMZ or BCNU, and cells were incubated for 96, 120 and 144 h.

380 To determine the effects of HCMV during drug treatment, A172, TP365MG and  
381 U251MG cells were plated in medium with low serum (10.000 cells/well in 24-well  
382 plates), infected at MOI 3 overnight with AD169 or TB40E, and treated with  
383 chemotherapeutic drugs. A172 and TP365MG cell lines were treated with 50 µM BCNU  
384 or TMZ and U251MG cells were treated with 25 µM BCNU or 75 µM TMZ. In all cases,  
385 cells were incubated for 96 h and cell viability was analyzed by MTT (3-(4,5-  
386 dimethylthiazolyl-2)-2, 5-diphenyltetrazolium bromide) or Alamar Blue assays (Thermo  
387 Fisher Scientific). The MTT assay was performed as previously reported[65, 66].  
388 Briefly, after incubation with drugs the medium was removed, cells were washed twice  
389 with PBS, and 200 µL of a 0.5 mg/mL MTT solution in serum-free medium was added  
390 to each well. The cells were incubated for 4 h at 37°C, medium was removed, and  
391 formazan crystals were solubilized in 300 µl DMSO. The plate was shaken for 15 min  
392 and absorbance was measured at 570 nm. For the Alamar Blue assay, 100 µL reagent  
393 was added directly to the culture media and incubated for 3 h. Fluorescence was

394 measured at 530 nm excitation and 560 nm emission. In both assays results were  
395 compared to controls considered 100% viable, where cells were seeded and cultivated  
396 as described but not exposed to drugs and/or virus. DMSO or EtOH were used as  
397 solvent controls for TMZ and BCNU, respectively, at concentrations below 0.1%  
398 (concentration non-toxic to the cells). For trypan blue staining, A172, TP365MG and  
399 U251MG cells were plated and treated as in the cell viability assays. After treatment,  
400 cells were trypsinized, and the cell suspension mixed with 0.4% trypan blue solution at  
401 a 1:1 ratio. Live and dead cells were counted using a Neubauer chamber. The  
402 percentage of living cells was measured using the following formula: % live cells =  
403 number of live cells / (number of live cells + number of dead cells) × 100%.

404 To determine the effects of the HCMV secretome during drug treatment, MRC-  
405 5 cells were infected with AD169 or TB40E at low MOI for 48 h. Cell supernatants were  
406 centrifuged and filtrated through 0.4 µm filters. A172, TP365MG and U251MG cells  
407 were plated overnight (10.000 cells/well), inoculated with 200 µl supernatant overnight,  
408 and treated with chemotherapeutic drugs as described above. After 96 h, cell viability  
409 was analyzed by MTT assay.

410

### 411 **Flow cytometry**

412

413 U251MG cells (10.000 cells/well) were infected with HCMV BAD<sub>in</sub>UL99GFP or  
414 mock-infected and treated with 25 µM TMZ or BCNU, incubated with PI for 15 min,  
415 trypsinized, and collected. After centrifugation and washing, cells were permeabilized  
416 with 0.1% Triton X-100, blocked with 3% BSA solution, and incubated overnight with  
417 anti-IE1 antibody (1:5) followed by goat anti-mouse Alexa 488 antibody (Abcam,

418 ab150113) (1:250) for 30 min. The cell pellet was re-suspended in PBS, and 10,000  
419 events were analyzed in a BD FACSCalibur flow cytometer.

420

## 421 **Statistical analysis**

422

423 All experiments were performed at least twice (biological duplicates) in technical  
424 triplicates. Statistical analysis was performed using two-way ANOVA (GraphPad  
425 Prism).  $P < 0.05$  was considered statistically significant.

426

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430 The authors declare that there is no conflict of interest.

431

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628

629

## 630 **FIGURE LEGENDS**

631

632 **Fig. 1. Detection of viral pp65 protein in HCMV-infected GBM cells.** A172,  
633 TP365MG and U251MG cells were infected with HCMV AD169 or TB40E at MOI 3 for  
634 2 h. Confocal immunofluorescence microscopy was performed, and a series of z-  
635 images were recorded. The HCMV pp65 protein is shown in green, cell nuclei are  
636 shown in blue, and merge. Images were taken at magnification objective of 63x.

637

638 **Fig. 2. Expression of viral IE1-72kDa, pp52 and pp28 proteins in HCMV-infected**  
639 **GBM cells.** A172, TP365MG and U251MG cells were infected with HCMV AD169 or  
640 TB40E at MOI 3 for 16, 48 or 96 h and analyzed by immunofluorescence microscopy.  
641 HCMV proteins are shown in red, cell nuclei are shown in blue, and merge images is  
642 shown as well. Images were taken at magnification objective of 20x.

643

644 **Fig. 3. Quantification of viral IE1-72kDa, pp52 and pp28 protein accumulation in**  
645 **HCMV-infected GBM cells.** Immunofluorescence microscopy for HCMV immediately  
646 early (IE1-72kDa), early (pp52) and late (pp28) proteins was performed at different  
647 times post infection in A172, TP365MG and U251MG cells. Results are expressed as  
648 a percentage of infected cells divided by the number of total cells. At least 100 cells  
649 were counted for each immunofluorescence sample using ImageJ.

650

651 **Fig. 4. Cell viability of GBM cells after treatment with TMZ and BCNU.** The number  
652 of A172, TP365MG and U251MG viable cells was assessed by MTT assay after  
653 treatment with 25, 50 or 100  $\mu$ M TMZ or BCNU for 96, 120 and 144 h. Data represent  
654 the mean ( $\pm$  standard deviation, SD) of three independent experiments done in  
655 triplicate,  $p < 0.05$ .

656

657 **Fig. 5. Cell viability of GBM cells after HCMV infection at different MOIs and**  
658 **treatment with TMZ and BCNU.** Cell viability was assessed by MTT assay for A172  
659 cells and by Alamar Blue assay for TP365MG and U251MG cells after overnight  
660 infection with HCMV AD169 or TB40E at the indicated MOIs and treatment with  
661 chemotherapeutic drugs. A172 and TP365MG cells were treated with 50  $\mu$ M TMZ or  
662 BCNU for 96 h, and U251MG cells were treated with 75  $\mu$ M TMZ and 35  $\mu$ M BCNU for  
663 96 h. Data represent the mean ( $\pm$  standard deviation, SD) of three independent  
664 experiments, \*  $p < 0.05$ , \*\*  $p < 0.0036$ , \*\*\*  $p < 0.0009$ ; \*\*\*\*  $p < 0.0001$ .

665

666 **Fig. 6. Cell viability of GBM cells after incubation with supernatants from HCMV-**  
667 **infected cells and treatment with TMZ and BCNU.** A172, TP365MG and U251MG  
668 cell viability was assessed by MTT assay after overnight incubation with supernatants

669 of AD169- or TB40E-infected MRC-5 cells for 48 h and treatment with  
670 chemotherapeutic drugs for 96 h. A172 and TP365MG cells were treated with 50  $\mu$ M  
671 TMZ or BCNU, and U251MG cells were treated with 75  $\mu$ M TMZ or 35  $\mu$ M BCNU. Data  
672 represent the mean ( $\pm$  standard deviation, SD) of three independent experiments, \*\*  
673  $p < 0.0098$ .

674

675 **Fig 7. Cell death analysis of HCMV-infected GBM cells.** (A) FACS analysis of  
676 U251MG cells infected overnight with BAD $in$ UL99GFP at MOI 3 (HCMV) or mock-  
677 infected after treatment with 25  $\mu$ M TMZ or BCNU for 96 h. A monoclonal antibody  
678 against HCMV IE1-72kDa was used as a marker for infection and propidium iodide (PI)  
679 as a marker for cell death. (B) For trypan blue exclusion assay A172, TP365MG and  
680 U251MG cells were infected overnight at MOI 3 with HCMV AD169 or TB40E and  
681 treated with chemotherapeutic drugs for 96 h. A172 and TP365MG cells were treated  
682 with 50  $\mu$ M TMZ or BCNU, and U251MG cells were treated with 75  $\mu$ M TMZ or 35  $\mu$ M  
683 BCNU. After treatment, cells were trypsinized and the cell suspension mixed with 0.4%  
684 trypan blue solution at a 1:1 ratio. Living and dead cells were counted using a  
685 Neubauer chamber. Data represent the mean ( $\pm$  standard deviation, SD) of three  
686 independent experiments, \*  $p < 0.05$ , \*\*  $p < 0.0036$ , \*\*\* $p < 0.0009$ .

687

688

689 **Fig. S1. Control experiments for the analysis of viral protein expression.**

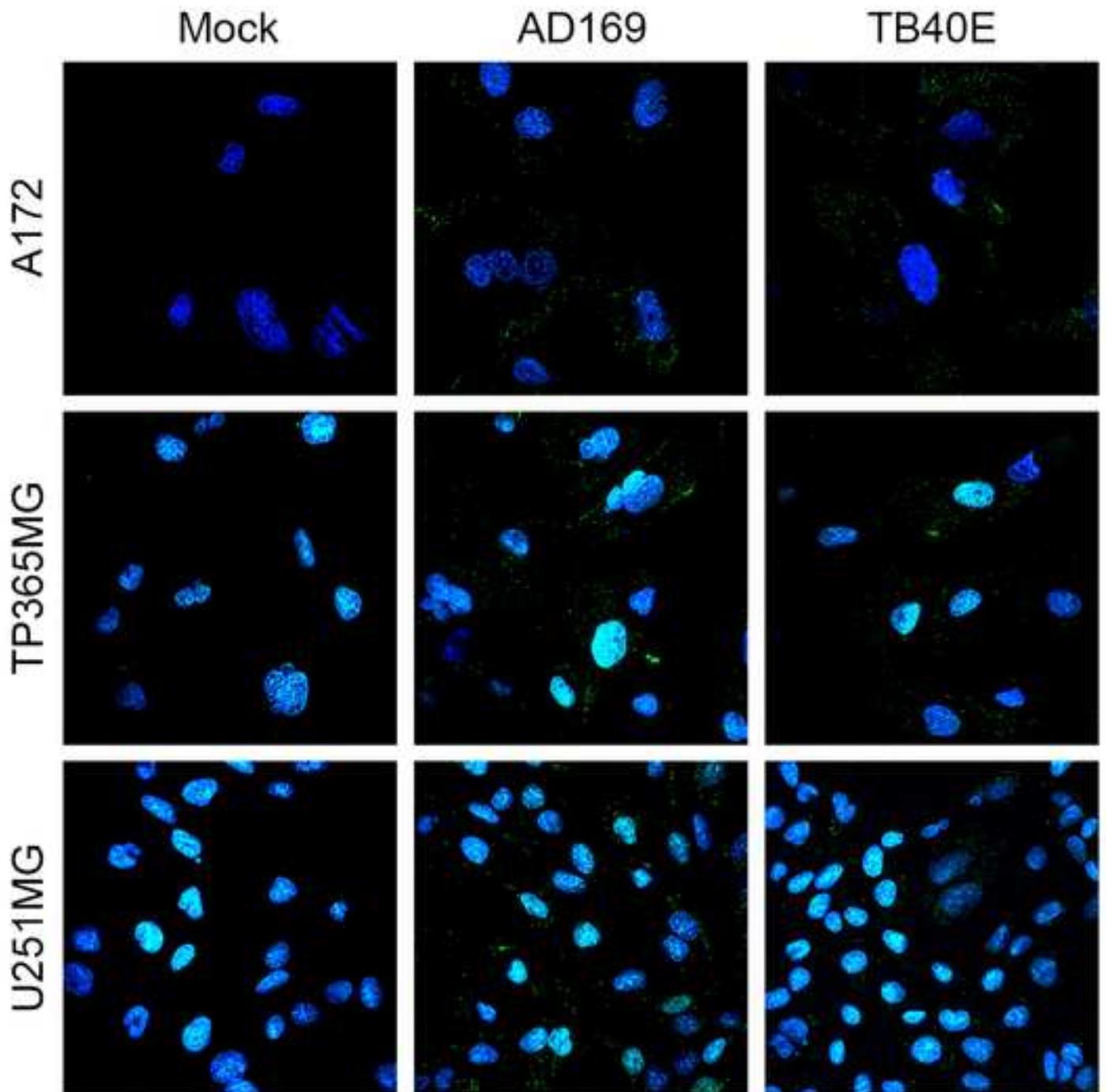
690 Immunofluorescence microscopy in mock-infected cells with HCMV-specific primary  
691 antibodies for HCMV proteins and secondary antibody; mock-infected cells with  
692 secondary antibody only and HCMV-infected cells with secondary antibody only.  
693 HCMV proteins are shown in red, cell nuclei are shown in blue. Images were taken at  
694 magnification objective of 20x.

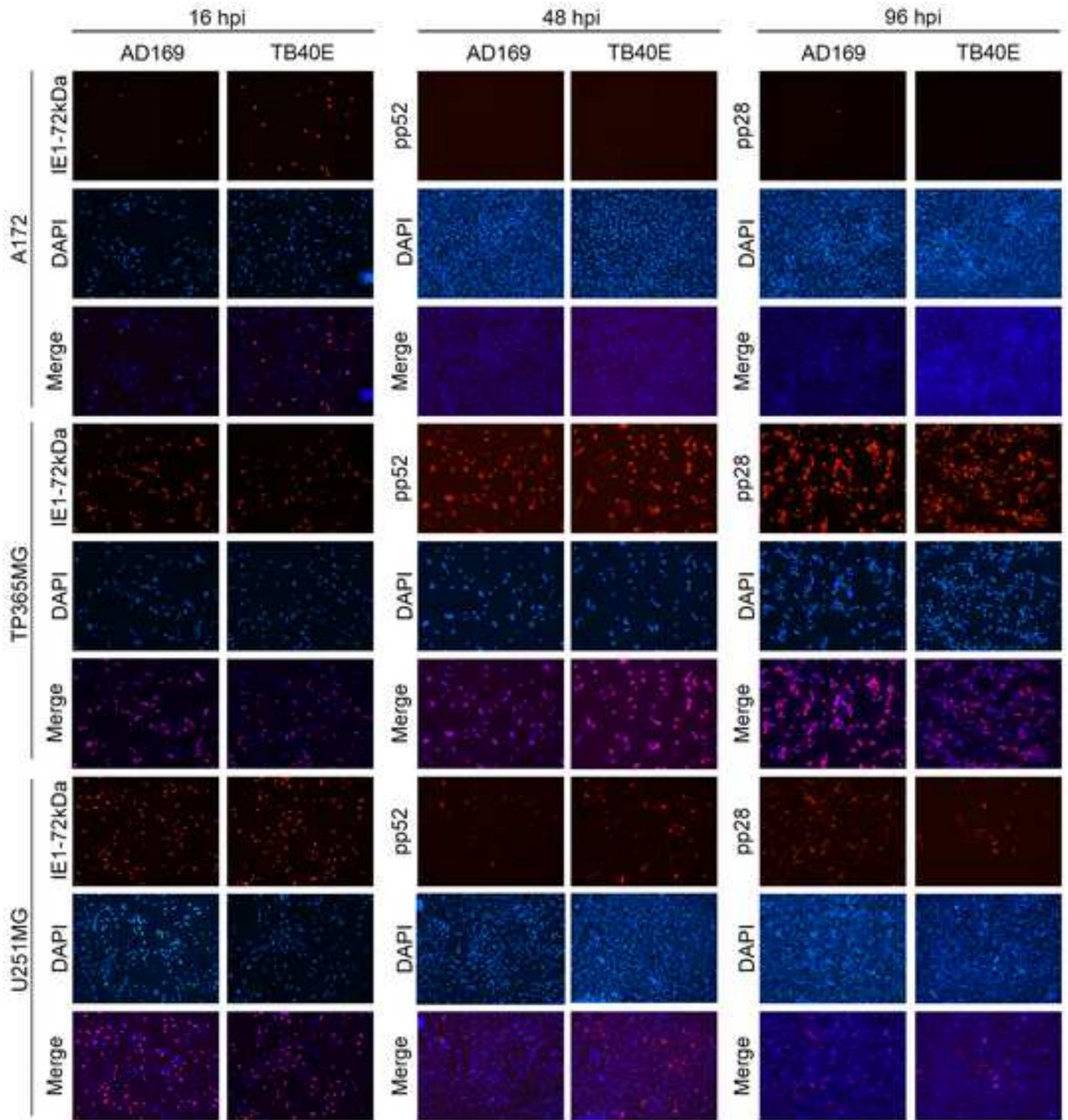
695

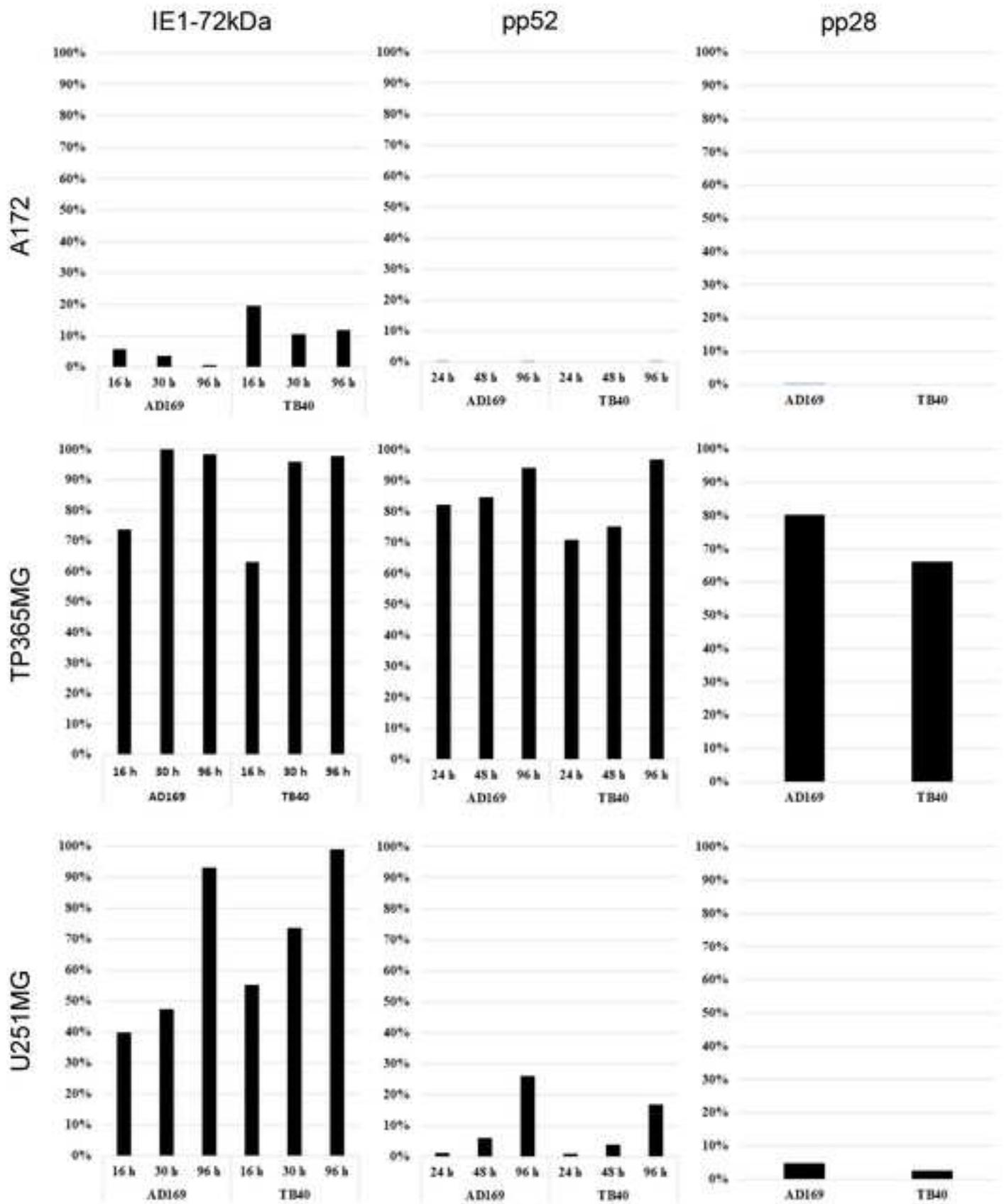
696 **Fig. S2. Control experiments for the analysis of HCMV protein expression.**

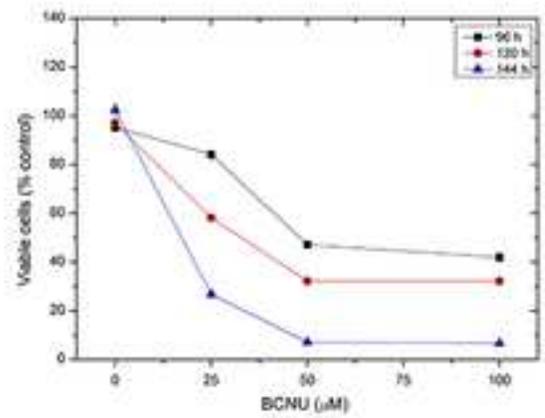
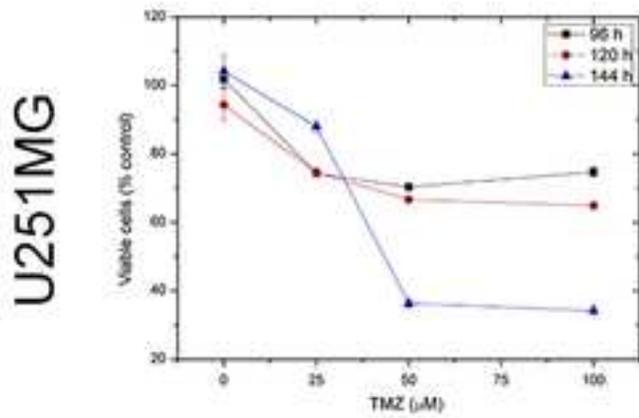
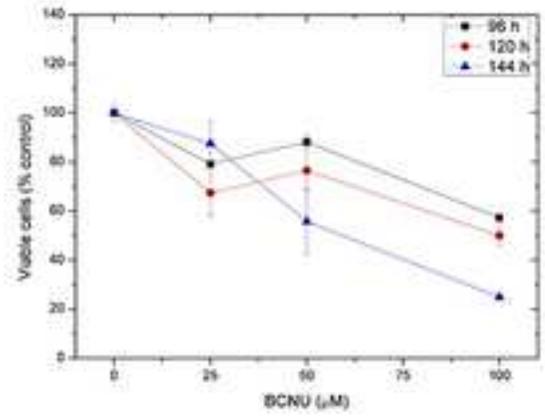
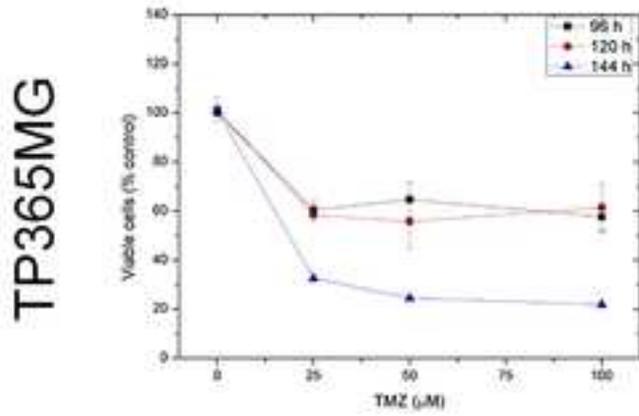
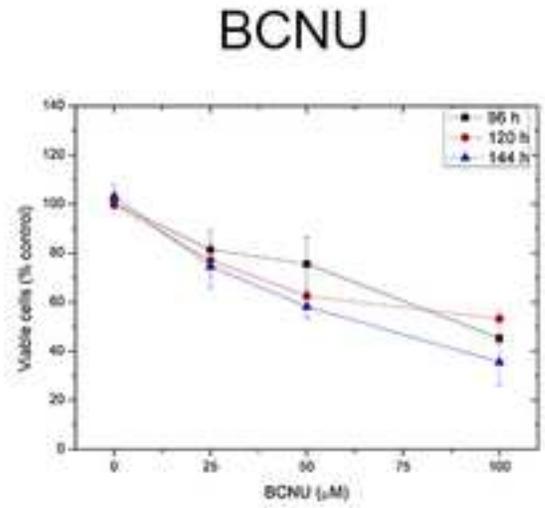
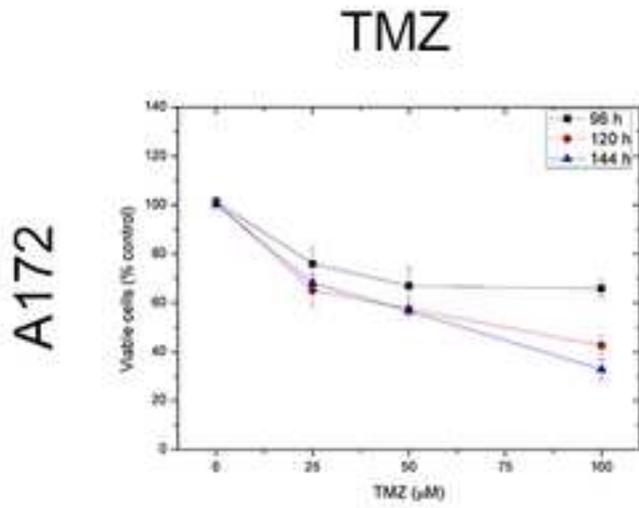
697 Immunofluorescence microscopy in mock-infected and HCMV infected cells with a  
698 mouse monoclonal antibody against the VP1 protein of JC polyomavirus and the  
699 secondary. IE1 staining in HCMV infected cells is shown as positive control. Positive  
700 control for HCMV infection. Viral proteins are shown in red, cell nuclei are shown in  
701 blue. Images were taken at magnification objective of 20x.

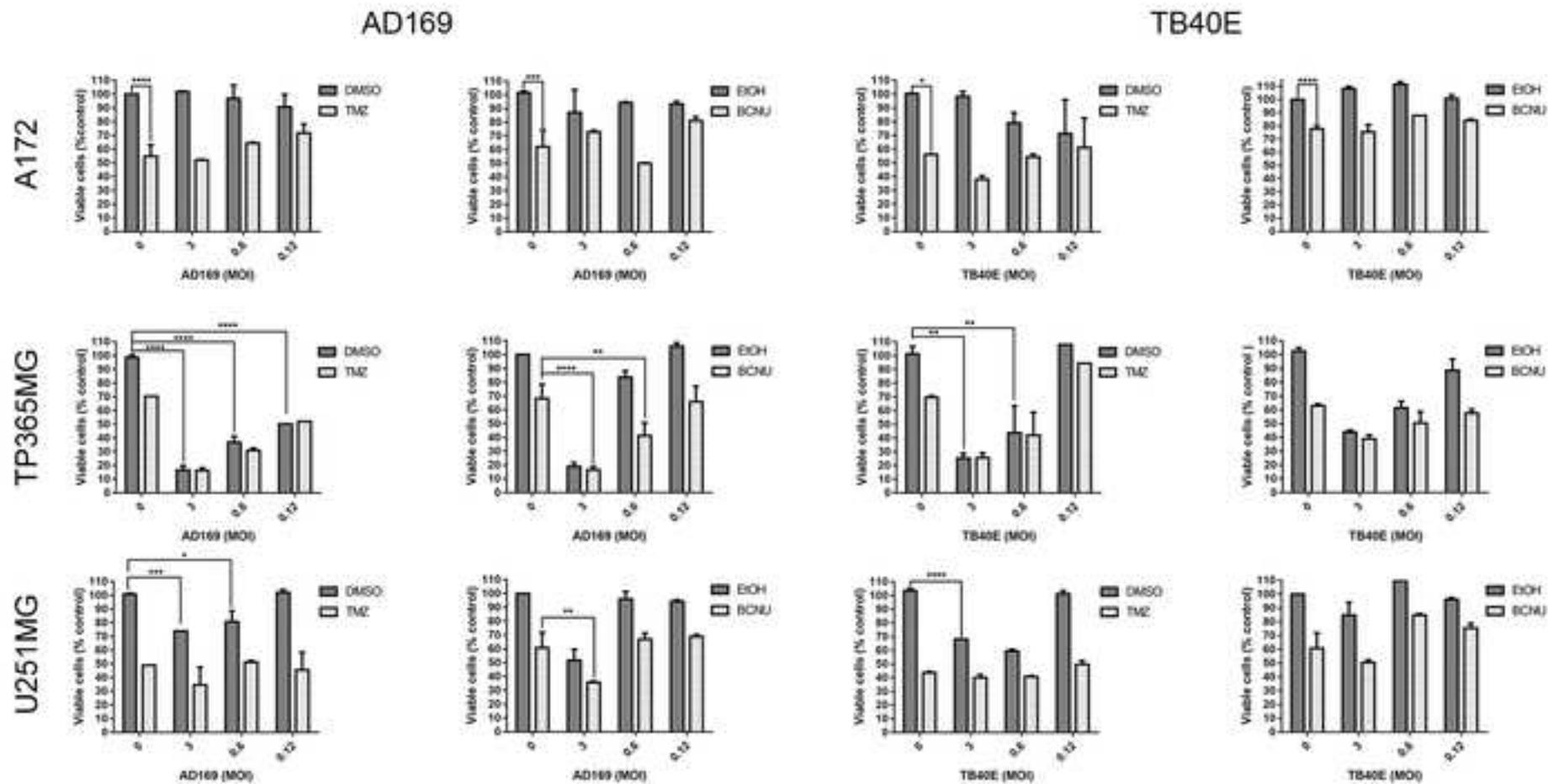
702

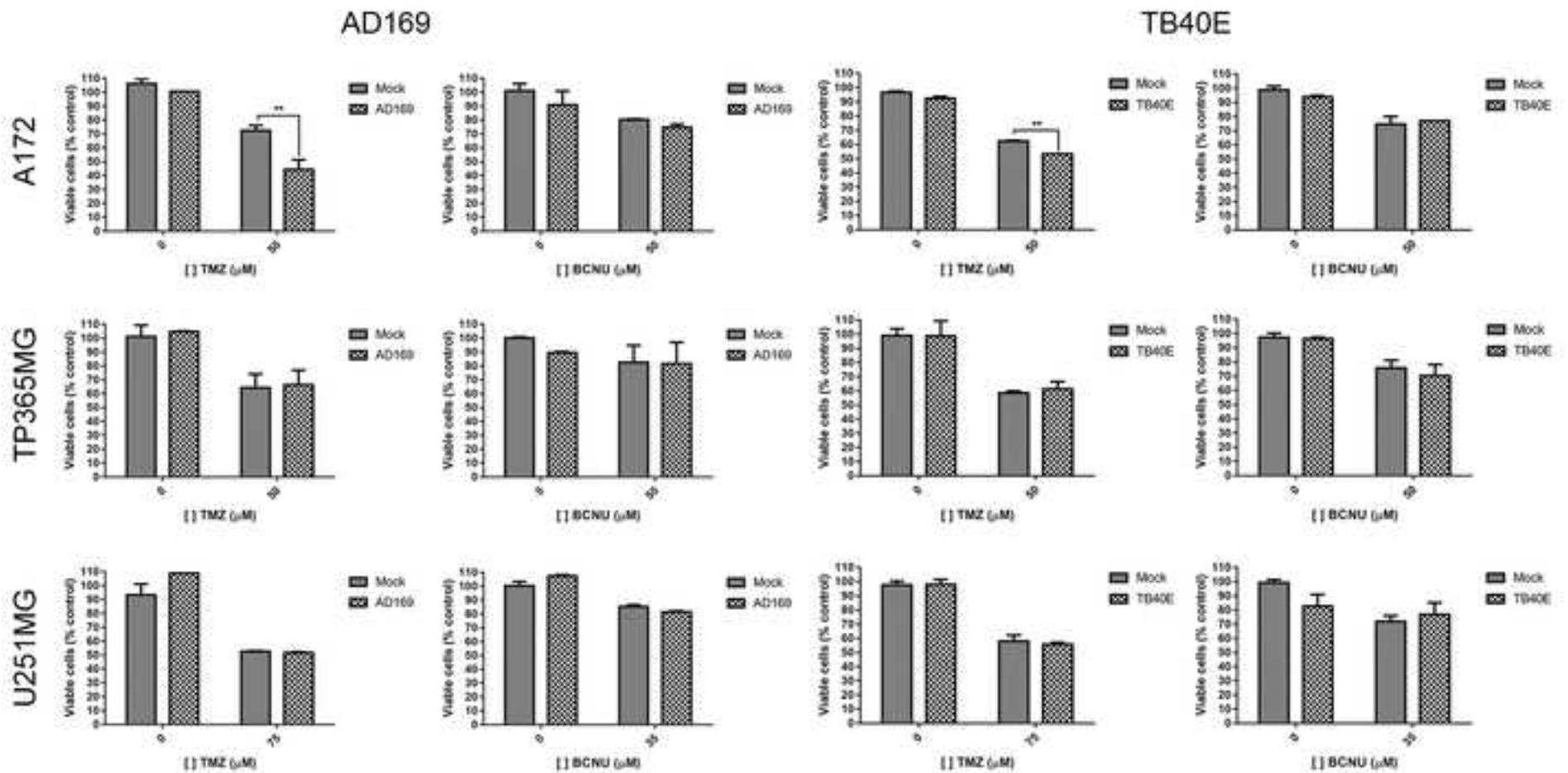




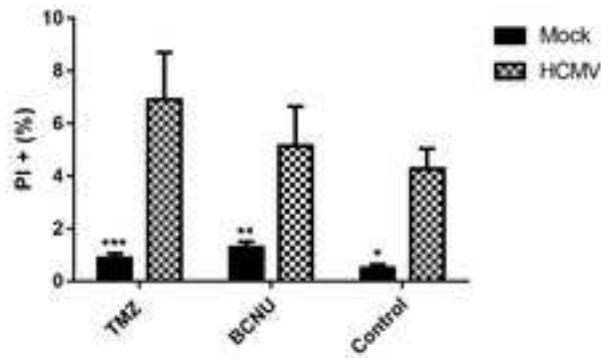




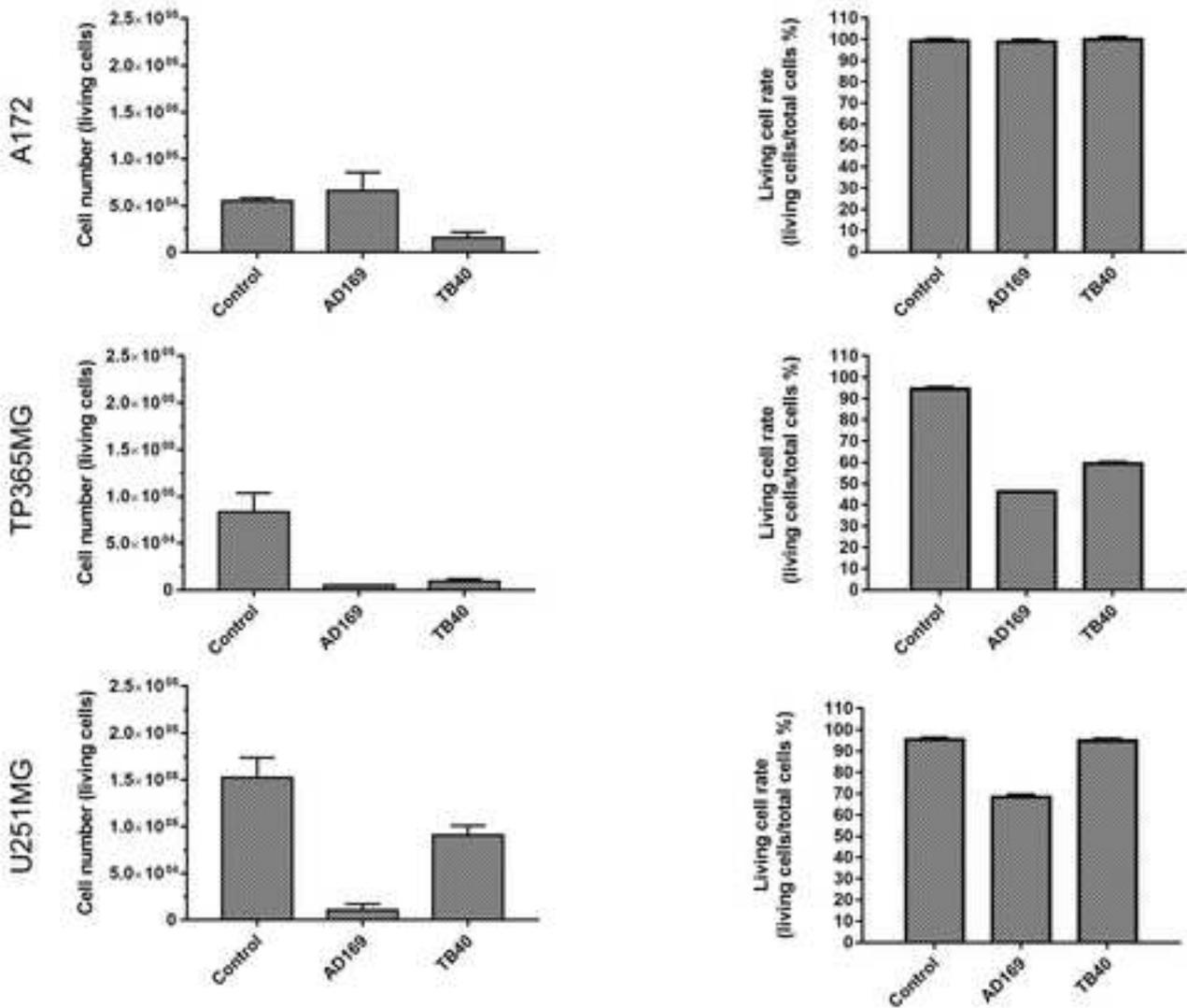


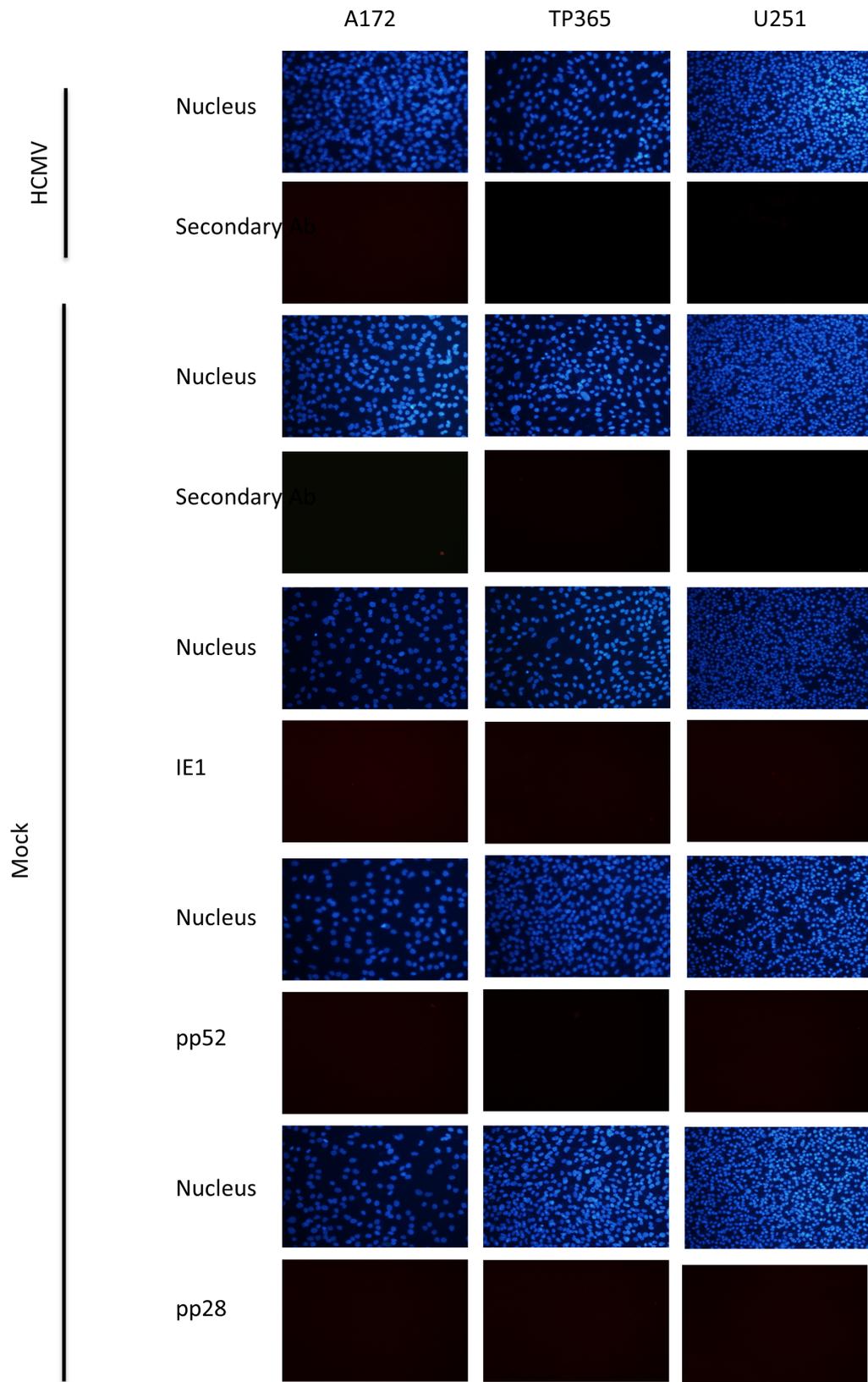


(a)



(b)

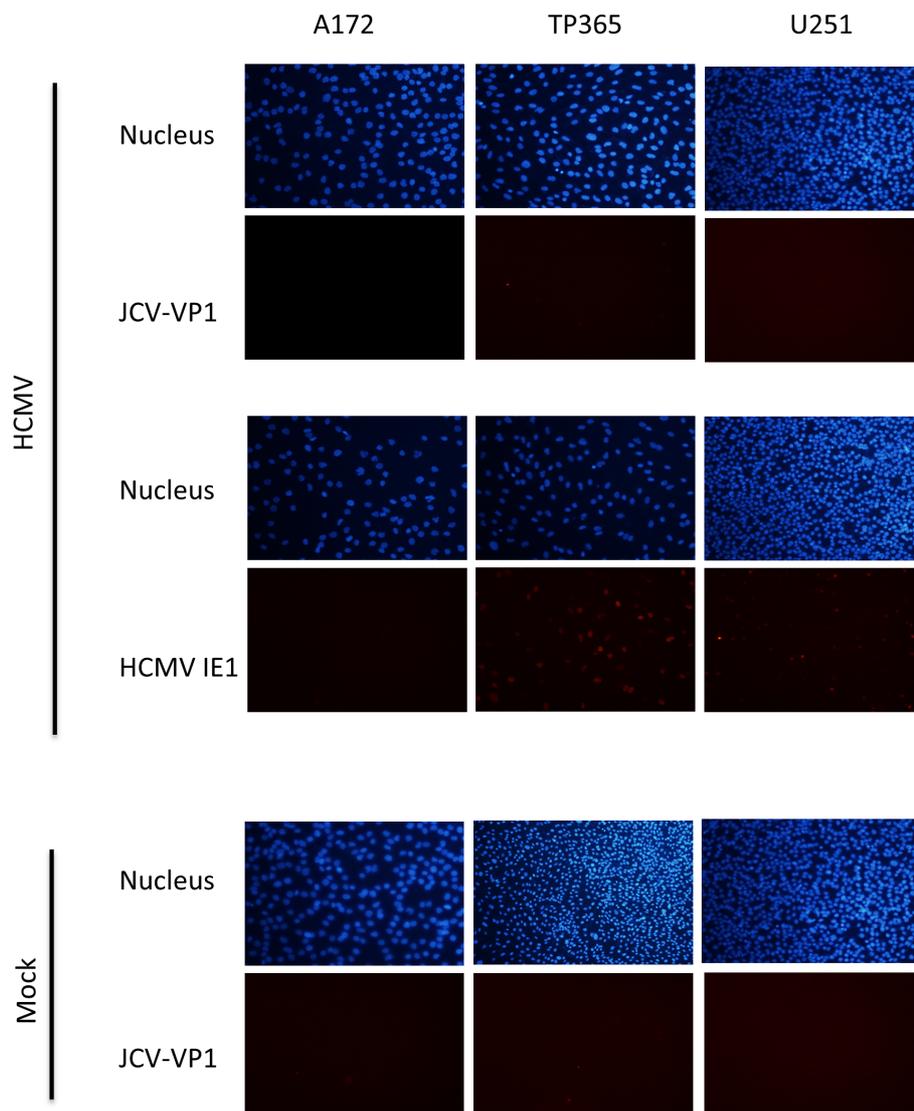




2 **Fig. S1. Control experiments for the analysis of viral protein expression.**

3 Immunofluorescence microscopy in mock-infected cells with HCMV-specific primary  
4 antibodies for HCMV proteins and secondary antibody; mock-infected cells with  
5 secondary antibody only and HCMV-infected cells with secondary antibody only.  
6 HCMV proteins are shown in red, cell nuclei are shown in blue. Images were taken at  
7 magnification objective of 20x.

8



10 **Fig. S2. Control experiments for the analysis of HCMV protein expression.**

11 Immunofluorescence microscopy in mock-infected and HCMV infected cells with a  
12 mouse monoclonal antibody against the VP1 protein of JC polyomavirus and the  
13 secondary. IE1 staining in HCMV infected cells is shown as positive control. Positive  
14 control for HCMV infection. Viral proteins are shown in red, cell nuclei are shown in  
15 blue. Images were taken at magnification objective of 20x.

16