

1 **A versatile reporter system to study cell-to-cell and cell-free bovine leukemia virus  
2 infection**

3 Florencia Rammauro<sup>1,2</sup>, Martín Fló<sup>1,2,‡</sup>, Federico Carrión<sup>1,‡</sup>, Claudia Ortega<sup>3</sup>, Francesca Di  
4 Nunzio<sup>4</sup>, Alexander Vallmitjana<sup>5</sup>, Otto Pritsch<sup>1,2,†</sup>, Natalia Olivero-Deibe<sup>1,\*</sup>

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6 <sup>1</sup>Laboratorio de Inmunovirología, Institut Pasteur de Montevideo, Montevideo, Uruguay

7 <sup>2</sup>Unidad académica de Inmunobiología, Facultad de Medicina, Universidad de la Repùblica,  
8 Montevideo, Uruguay

9 <sup>3</sup>Programa de Tecnología Molecular, Celular y Animal, Institut Pasteur de Montevideo,  
10 Montevideo, Uruguay

11 <sup>4</sup>Advanced Molecular Virology Unit, Institut Pasteur-Université Paris Cité, 75015 Paris,  
12 France

13 <sup>5</sup> Department of Biomedical Engineering, University of California Irvine, CA, USA

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15 \*Corresponding author: Natalia Olivero-Deibe, [nolivero@pasteur.edu.uy](mailto:nolivero@pasteur.edu.uy)

16 <sup>‡</sup> These authors have contributed equally to this work

17 <sup>†</sup> Deceased

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20 syncytium, cell free transmission

21

22 **ABSTRACT**

23

24 Bovine leukemia virus (BLV) is a B-lymphotropic oncogenic retrovirus of the genus  
25 *Deltaretrovirus* that infects dairy cattle worldwide and is the causative agent of enzootic  
26 bovine leukemia. BLV demonstrates remarkably low efficiency in infecting cells via free  
27 viral particles derived from infected B cells, as virions are rarely detected in the  
28 bloodstream of infected cattle. However, transmission efficacy significantly increases  
29 upon the establishment of direct cell-to-cell interactions. Syncytium formation assays are  
30 the main tool in the study of BLV infectivity. Although this traditional method is highly  
31 robust, the complexity of visually counting syncytia poses a significant technical  
32 challenge. Using lentiviral vectors, we generated a stable reporter cell line, in which the  
33 GFP reporter gene is under the control of the full-length BLV-LTR. We have  
34 demonstrated that the BLV-Tax protein and histone deacetylases inhibitors, such as VPA  
35 and TSA, can transactivate the BLV LTR. Upon co-culturing the reporter cell line with the  
36 BLV-infected cells, fluorescent syncytia can be visualized. By implementing automated  
37 scanning and image acquisition using a confocal microscope, together with the  
38 development of an analysis software, we can detect and measure single GFP cells and  
39 fluorescent multinucleated cells. In summary, our reporter cell line, combined with the  
40 development of analysis software, is a useful tool for understanding the role of cell fusion  
41 and cell-free mechanism of transmission in BLV infection.

## 42 INTRODUCTION

43 Bovine leukemia virus (BLV) is an oncogenic deltaretrovirus that infects cattle  
44 worldwide and causes Enzootic bovine leukosis (EBL). BLV causes persistent subclinical  
45 infection (60% of the infected animals are asymptomatic, about 30% develop persistent  
46 lymphocytosis, and about 5-10% develop an aggressive tumor pathology called  
47 lymphosarcoma, which frequently causes the death of these animals [1], [2]. BLV  
48 infection leads to high economic losses in dairy and beef industries, related to mortality  
49 caused by lymphosarcoma [3]; negative impact on production parameters [4],[5] ;  
50 immunological alteration and secondary infections [6]; and restriction on the international  
51 trade of live cattle, semen, and infected embryos [7], [3], [8].

52 In contrast to retroviruses like HIV, deltaretroviruses such as BLV or human T  
53 cell leukemia virus (HTLV) exhibit remarkably low efficiency in infecting cells via free  
54 viral particles derived from B or T lymphocytes, respectively, as these virions are seldom  
55 detected in the bloodstream of infected hosts [9] , [10]. Thus, efficient BLV transmission  
56 from a seropositive bovine to a susceptible counterpart primarily occurs via cell-  
57 containing fluids, such as blood, semen, milk, colostrum, saliva, or mucus, through  
58 horizontal or vertical routes [9].

59 Transmission efficacy substantially increases upon the establishment of direct  
60 cell-to-cell interactions. This mode of spread not only facilitates rapid viral dissemination  
61 but may also promote immune evasion and influence disease progression [11], [12], [13].  
62 Cell-cell fusion (syncytium formation) has been identified as an alternative pathway for  
63 cell-to-cell transmission [13, 14].

64 Due to the intrinsic difficulty of infecting cells with cell-free BLV virions,  
65 syncytium formation assays are the main tool in the study of BLV infectivity [15–17].  
66 This assay involves co-culturing an "acceptor" cell line, typically CC81 with a cell line  
67 persistently infected with BLV, such as FLKBLV, as the "donor". The syncytia  
68 (multinucleated cells) form by fusion between an infected and an uninfected cell due to  
69 co-culture [16, 18]. After co-cultivation, the cells are commonly stained with Giemsa, and  
70 the number of syncytia is visually counted. Although this traditional method is highly  
71 robust, the complexity of visually counting syncytia poses a significant technical  
72 challenge, especially in high-throughput screening assays.

73 Reporter cell lines capable of detecting infectious viruses have been developed  
74 utilizing a BLV long terminal repeat (LTR) as a promoter and GFP or Luciferase as the  
75 reporter gene. Infected cells can be monitored using fluorescence microscopy or  
76 luminescence. These reporter cells offer greater sensitivity and quantification compared  
77 to traditional methods [19–22].

78 The BLV LTR promoter bears many regulatory sites and is responsible for virus  
79 integration and replication. It comprises by three distinct regions: the U3, R, and U5. The  
80 U3 region, contains several critical cis-acting elements besides to the CAAT box, TATA  
81 box, and transcription start site [23, 24]. The primary regulatory elements are three copies  
82 of an imperfectly conserved 21-bp sequence called the Tax-responsive element (TxRE).  
83 The TxREs are essential for the promoter's responsiveness to the Tax transactivator  
84 protein. The BLV LTR has been suggested as a gene promoter under Tax activation in  
85 mammalian cells [19]. A glucocorticoid-responsive element (GRE) responds to  
86 dexamethasone in the presence of glucocorticoid receptors and Tax [25]. Additionally,  
87 there are multiple binding sites for several transcription factors, including two AP-4 sites,  
88 a glucocorticoid response element (GRE), and a PU.1/Spi-B–binding site. In the U5  
89 region, a binding site for the interferon-responsive factor is present. These binding sites  
90 regulate BLV transcription, dependent on or independently of BLV-Tax expression [21].

91 Moreover, BLV transcriptional activity is influenced by the acetylation and methylation  
92 of these binding sites [26].

93 Using lentiviral vectors, we generated a stable reporter cell line to measure BLV  
94 infectivity, in which the EGFP reporter gene was expressed under the control of the full-  
95 length BLV-LTR. We have demonstrated that the Tax protein of BLV and histone  
96 deacetylases inhibitors (HDACi) such as, VPA and TSA, can transactivate the BLV LTR.  
97 When cultured with BLV-infected cells, the reporter cell line forms fluorescent syncytia.  
98 By integrating automated scanning and image acquisition using a confocal microscope,  
99 and developing an analysis software, we can detect single GFP cells and fluorescent  
100 multinucleated cells with two or more nuclei. Our results show that this reporter cell line  
101 is sensitive and, when combined with automated analysis, provides a useful tool to study  
102 both cell-to-cell and cell-free infection.

## 103 MATERIAL AND METHODS

### 104 Cell culture

105 HEK293T (ATCC CRL-3216) and CC81 (feline cell line transformed by mouse sarcoma  
106 virus ECACC 90031403) cells were grown in Dulbecco's modified Eagle's medium  
107 (DMEM)- high glucose (GlutaMAX, Gibco) supplemented with 1%  
108 penicillin/streptomycin and 10% fetal bovine serum (Gibco, USA). Persistently infected  
109 FLKBLV (Fetal lamb kidney, DSMZ-ACC 153) and BL3.1 (bovine B-lymphosarcoma,  
110 ATCC CRL-2306) cells were grown in Roswell Park Memorial Institute (RPMI) 1640  
111 Medium-high glucose (GlutaMAX, Gibco) supplemented with 10% inactivated fetal  
112 bovine serum, 1% penicillin/streptomycin, 1% Sodium pyruvate. All cell lines were  
113 incubated at 37°C in a 5%CO<sub>2</sub> humidified atmosphere.

### 114 Plasmids

115 DNA sequences encoding the full BLV Tax protein (GenBank, access number:  
116 EF600696) was synthesized and cloned in pcDNA3.1(+)-C-DYK by Genescrypt. The  
117 resulting plasmid, named pcDNA3.1BLVTAX-FLAG, encodes FLAG-tagged BLV Tax.  
118 pTrip lentiviral vector (pTripCMVGFP) [27], in which GFP are under the control of CMV  
119 promoter was used as a reporter or transfer vector. The CMV promoter was substituted  
120 with the 5' LTR of BLV (531 pb, GenBank, access number: EF600696) to generate the  
121 pTripLTRBLVGFP vector. 5' LTR of BLV was synthesized and cloned in  
122 (pTripCMVGFP) by Genescrypt. The commercial plasmids Lenti DR8.75 (Addgene  
123 #22036) and pMD2.G (Addgene #12259) were employed as packaging and envelope  
124 vectors, respectively.

### 125 Transfection of HEK293T cells to obtain lentiviral particles (LVs)

126 HEK293T cells were co-transfected with the following plasmids: pTripLTRBLVGFP or  
127 pTripCMVGFP, DR8.74, and pMD2.G, to obtain lentiviral particles (LVs) with the VSV  
128 G protein on the surface and coding for GFP under the LTR of BLV (LVs LTRBLV-  
129 GFP) or under the CMV promoter as a control (LVs CMV-GFP). The day before  
130 transfection, 2.5x10<sup>6</sup> HEK293T cells were seeded in 10 mL of complete medium  
131 (DMEM-GlutaMAX, 10% FBS and 1% PenStrep) into 100 mm petri dishes. PEI at 1  
132 mg/mL (Polysciences) was used as the transfection agent. DNA:PEI complexes in a 1:10  
133 ratio were prepared using 10 µg of pTripLTRBLBGFP or pTripCMVGFP, 6.5 µg of  
134 DR8.74, and 3.5 µg of pMD2.G, and added to the cells. After 12-18 h post-transfection,  
135 the medium was substituted with fresh medium. Forty-eight hours post-transfection,

136 supernatant containing LVs was harvested and filtered by 0.45  $\mu$ m and stored in aliquots  
137 at -80 °C.

138 Transduction for the establishment of the stable reporter cell line.

139 The day before transduction,  $0.5 \times 10^5$  CC81, or FLKBLV cells per well were seeded in a  
140 24-well plate to achieve a monolayer with 70% confluence. For transduction, the culture  
141 medium was removed, and 250  $\mu$ L of LVs (LVs LTRBLVGF or LVs CMVGF) or cell  
142 culture medium (in negative control cells) together with 10  $\mu$ L of polybrene (250  $\mu$ g/mL)  
143 were added to each well. An additional 250  $\mu$ L of cell culture medium was added to all  
144 wells, and the plate was incubated at 37°C and 5% CO<sub>2</sub> for 60 h. GFP expression was  
145 evaluated by means of an epifluorescence microscope and images were inspected using  
146 ImageJ. A frozen stock of transduced cells was generated, and some cells were expanded  
147 for further experiments. The stable cell lines generated were designated  
148 CC81LTRBLVGF or FLKLTRBLVGF.

149 Transient transfection of CC81LTRBLVGF cells with plasmids encoding BLV Tax  
150 protein

151  $1 \times 10^5$  CC81LTRBLVGF cells per well were seeded in a 12-well plate for 24 hours, and  
152 the cell culture medium was changed 3 hours before transfection. The PEI transfection  
153 reagent at 1 mg/mL was prepared in OptiMEM medium and added to a tube containing  
154 pcDNA3.1BLVTAX-FLAG or pcDNA3.1 (empty vector, as a control) (final  
155 concentration of 0.5  $\mu$ g/ $\mu$ l) in OptiMEM and incubated for 30 minutes. The DNA:PEI  
156 complexes were added drop by drop to CC81LTRBLVGF cells. Twenty-four hour post-  
157 transfection, fluorescence was observed using fluorescence microscopy and the  
158 percentage of GFP positive cells was determined by flow cytometry as described below.

159 Transactivation of CC81LTRBLVGF cells by valproic acid (VPA), trichostatin A (TSA)  
160 or INF $\alpha$

161 CC81LTRBLVGF cells were incubated for 24 hours to form a confluent monolayer.  
162 Subsequently, the cells were treated with 10mM VPA or 500nM TSA for 24 hours. To  
163 evaluate the effect of INF- $\alpha$  on BLV LTR promoter, CC81LTRBLVGF cells were  
164 treated without or with 75, 125, 250, 500 or 1000 U/mL of INF- $\alpha$  for 48 hours. The  
165 percentage of GFP positive cells was determined by flow cytometry as described below.

166 GFP protein expression analysis by flow cytometry

167 CC81LTRBLVGF cells transiently transfected with pcDNABLVTAX-FLAG or  
168 pcDNA3.1 (empty) or treated with VPA or TSA were detached through trypsinization  
169 and transferred to a V-bottom plate. Subsequently, the cells were washed twice with PBS-  
170 BSA 3% and fixed with 4% PFA. Cell acquisition was performed using the Attune NxT  
171 (Thermo Fischer Scientific) cytometer, and the data was analyzed with the FlowJo  
172 software (Tree Star Inc.). Non-transfected or transfected with empty vector, and untreated  
173 cells were used as a control of the fluorescence background.

174 Evaluation of the reporter cell line by infection with BLV

175 For setting up co-culture experiments, CC81LTRBLVGF cells (as target cells) were co-  
176 incubated with FLKBLV producing cells (donors) at ratios of 1:1, 1:2, and 1:4  
177 (donor:target) for 24 or 48 h at 37°C. Subsequently, the cells were washed with PBS,  
178 fixed with 4% PFA, and stained with DAPI (1/1000) for nuclei visualization. GFP  
179 expression was evaluated by means of an epifluorescence microscope and images were

180 inspected using ImageJ. Once the experiment was set up, the co-culture was carried out  
181 by co-incubating  $1,6 \times 10^5$  CC81LTRBLVGFP cells with  $0,4 \times 10^5$  FLKBLV in a 12-well  
182 plate for 48 h at 37°C. The co-culture experiments using the BL3.1 cell line were  
183 performed as follows: CC81LTRBLVGFP cells were seeded at a density of  $5 \times 10^4$  cells  
184 per well in a 12-well plate and cultured at 37°C for 24 hours. Subsequently, they were co-  
185 cultured with BL3.1 cells at a density of  $1 \times 10^5$  cells per well for 72 h. For all experiments,  
186 co-culture cells were washed with PBS, fixed with 4% PFA, and stained with DAPI  
187 (1/1000) for nuclei visualization. GFP expression was evaluated by means of an  
188 epifluorescence microscope and images were inspected using ImageJ.

189 *Transwell infection assay*

190  $1,6 \times 10^4$  FLKBLV cells, serving as viral particle-producing cells, were cultured on  
191 transwell inserts with a 0.4  $\mu$ m pore size for 24 h to establish a monolayer. Following cell  
192 culture, the transwell inserts were incubated with CC81LTRBLVGFP reporter cells per  
193 well, which were previously plated in a 24-well. As a control, a similar number of  
194 FLKBLV and CC81 reporter cells were directly co-cultured in 24-well plates without an  
195 insert. The system was then incubated at 37°C for 48 h to facilitate the interaction between  
196 viral particles and CC81LTRBLVGFP cells. Cells were then fixed with 4% PFA, washed  
197 three times with PBS and subsequently blocked for 15min with PBS-3%BSA-0.1% Triton  
198 X-100. Cells were further incubated 1 h at RT with monoclonal antibodies anti-p24  
199 (BLV3, 1/200, VMRD, USA). After three washes with PBS, cells were incubated for 1 h  
200 at RT with Alexa Fluor-594 conjugated goat anti-mouse IgG (Invitrogen) diluted 1/1,000  
201 in PBS added with 3% (w/v) BSA and DAPI 1/1000, washed three times with PBS and  
202 mounted in 70% (v/v) glycerol pH 8.8. Cell infection was assessed using Zeiss LSM800  
203 confocal microscopy.

204 *Fluorescence microscopy imaging*

205 Mosaics of  $5 \times 5$  tiles adding to twenty-five images per well were automatically acquired  
206 on Zeiss LSM800 confocal microscopy with a 10X objective. Two-source excitation was  
207 performed using 488 (power 0,92%) and 405 (0,86%) lasers, detection wavelength 510-  
208 700nm and 400-510nm respectively. Pinhole set at 25  $\mu$ m. Images were taken at 1,03  $\mu$ s  
209 pixel dwell time, a resolution of 4650x4650 pixel.

210 *Automated segmentation of fluorescently infected cells*

211 A custom set of scripts was written in MATLAB to perform automated segmentation of  
212 nuclei and GFP+ cells and automated counting of nuclei per cell and classification into  
213 syncytia. The two emission channels (DAPI and GFP) were processed independently to  
214 segment objects based on the knowledge of the expected area of nuclei. In both cases, a  
215 pre-processing step was performed, involving brightness normalization to the [0.1- 99.9]  
216 % quantiles of the total brightness and gaussian filtering; for the GFP channel using a  
217 standard deviation of  $\frac{1}{4}$  of the expected radius of nuclei and for the DAPI images a  
218 standard deviation of 0.1 of the expected radius of nuclei. Nuclei were segmented using  
219 a watershed-based algorithm with a size stopping rule at 4 times the expected area of a  
220 nuclei. GFP+ cells were segmented using simple Otsu thresholding allowing for a user-  
221 inputted factor to fine tune the segmentation. A post-filtering step was required to remove  
222 objects that were either smaller than twice the expected area of a nuclei or 200 time larger.  
223 Software is available upon request and will be uploaded to a public repository.

224

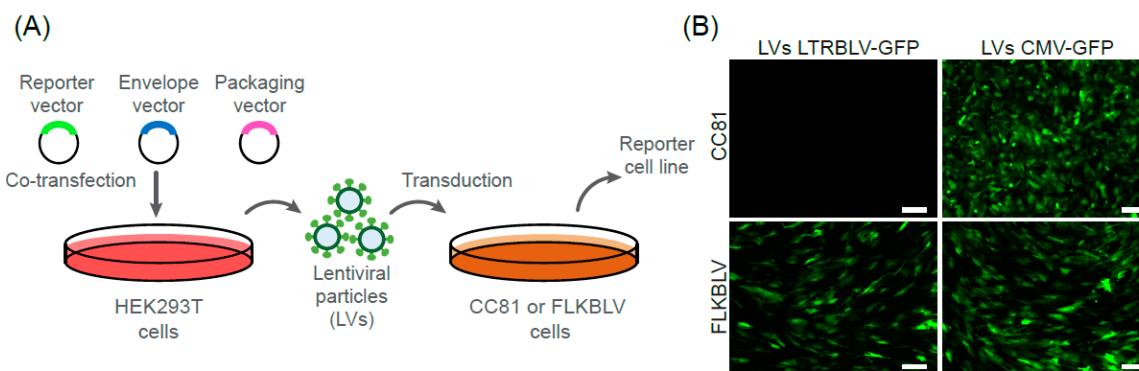
225 *Automated counting and feature measurement*

226 Once segmentation step was completed in both channels, for each individual segmented  
227 GFP+ cell, the number of segmented nuclei within the same spatial region were counted,  
228 distinguishing between partially included and completely included accounting for  
229 segmentation errors. The nuclei number per GFP+ cell was established as the average  
230 between the total distinct nuclei inside and the total nuclei that were completely inside.  
231 The criteria to classify into syncytia was those GFP+ segmented cells with more than 3  
232 detected nuclei. For completion a clustering step was performed using the distances  
233 between the centers of the detected nuclei, using a density-based algorithm (dbSCAN). The  
234 set of scripts allowed to automatically export a list of detected GFP+ cells (in the order  
235 of 300 per field of view), with their measured number of nuclei, relative brightness and  
236 measured area, and a list of detected nuclei (in the order of 10k per field of view).  
237 Software is available upon request and will be uploaded to a public repository.

## 238 RESULTS

### 239 *Generation of a reporter cell line for BLV infection*

240 A second-generation lentiviral system was utilized to develop the stable reporter cell line  
241 **Figure 1 A.** We replaced the CMV promoter in the pTripCMVGFP vector with the full-  
242 length 5'LTR of BLV to generate the reporter lentiviral vector. As shown in **Figure 1 B**,  
243 no GFP expression was detected for CC81 cells transduced with LVs LTRBLVGFP. This  
244 outcome is expected since the BLV LTR promoter has deficient basal activity and can  
245 only be activated and induce GFP expression in the presence of Tax or upon BLV  
246 infection. On the other hand, cell line expressing the BLV Tax transactivator protein  
247 (FLKBLV), transduced with LVs LTRBLV-GFP or CMV-GFP showed similar levels of  
248 GFP expression, indicating that the BLV promoter would have similar activity to the  
249 CMV promoter. Therefore, observing GFP in this cell line allows us to conclude that both  
250 the constructions and LVs production were successful. Finally, in both cell lines (CC81  
251 or FLKBLV) transduced with CMVGFP LVs (control), GFP expression was observed.  
252 This result was anticipated, as these LVs containing the constitutive CMV promoter can  
253 induce high levels of GFP transgene expression.



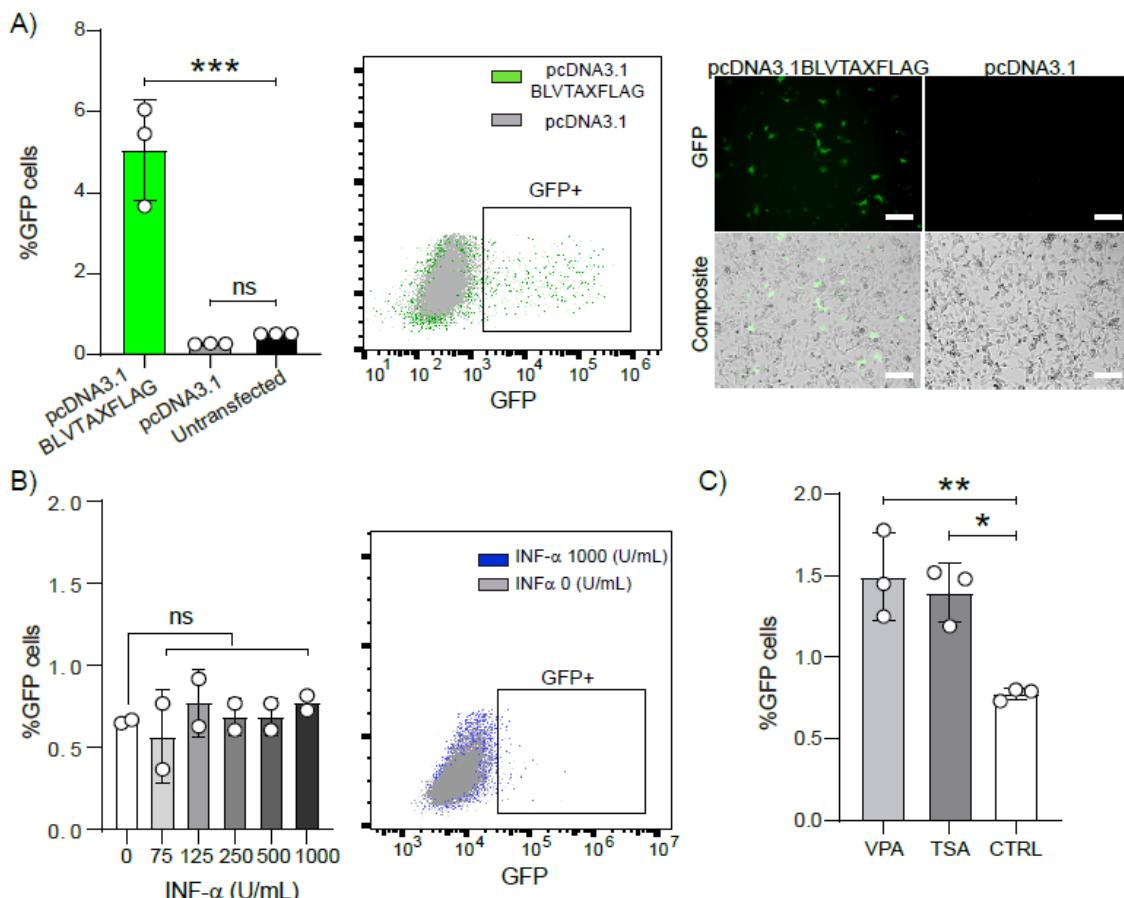
255 **Figure 1. Generation of the reporter cell line for BLV infection.** **(A)** Representative  
256 diagram of the process for obtaining the reporter cell lines CC81LTRBLVGFP and  
257 FLKBLVLTRGFP. HEK293T cells were co-transfected with packaging (DR8.74),  
258 envelope (VSV-G), and reporter (pTripLTRBLVGFP or pCMVGFP) plasmids. Forty-  
259 eight hours post-transfection, the supernatant containing lentiviral particles (LVs, LV-  
260 LTRBLVGFP or LV-CMVGFP) was collected. CC81 or FLKBLV cells were transduced  
261 with 250 µL of LV-LTRBLVGFP or LV-CMVGFP diluted in culture medium and 10 µL  
262 of polybrene 250 µg/mL and incubated for 60 h. **(B)** Representative images of transduced

263 cell lines. The FLKBLV cell line served as a positive control. GFP expression was  
264 evaluated by fluorescent microscopy and the obtained images were analyzed using Image  
265 J. Scale bar, 100  $\mu$ m.

266

267 *Functional testing of the reporter construct LTRGFP*

268 To assess the effect of BLV Tax protein expression due to transactivation of the BLV  
269 LTR, CC81LTRBLVGFP reporter cells were transiently transfected with an expression  
270 vector pcDNABLVTAX-FLAG or the empty vector as a negative control. Tax expression  
271 in transfected cells was confirmed by immunofluorescence and western blot 24 h after  
272 transfection (**Supplementary Figure 1 A, B**). We show that the induction of BLV Tax  
273 expression is enough to activate the BLV LTR in the reporter cell line, as evidenced by a  
274 significative increase of GFP+ cells. No GFP expression was detected in cells transfected  
275 with an empty vector or in untransfected cells (**Figure 2 A**). The U5 region present in the  
276 LTR contains regulatory interferon factor binding sites, which have previously been  
277 reported to drive tax-independent replication of BLV [28]. We tested the effect of IFN- $\alpha$   
278 treatment (0-1000 U/mL) on BLV LTR-driven GFP expression in the  
279 CC81LTRBLVGFP cell line. After 48 h of IFN treatment, we found no GFP expression  
280 in the cells at any of the concentrations tested (**Figure 2 B**). Previous reports have shown  
281 that deacetylase inhibitors can induce Tax-independent transactivation of the LTR. For  
282 example, TSA or VPA has been shown to be an efficient activator of BLV expression  
283 [19, 26, 29]. We demonstrate that a small percentage of CC81LTRBLVGFP cells show  
284 GFP expression after treatment with TSA and VPA (**Figure 1 C**). While there is a slight  
285 effect, it should be considered when designing experiments using our reporter cell line.

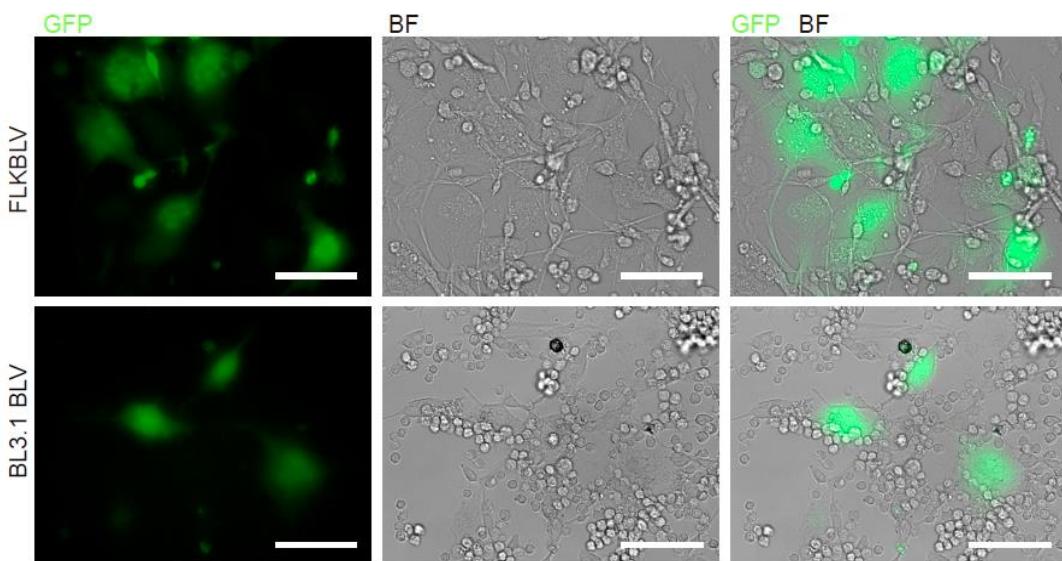


286

287 **Figure 2. Functional testing of the reporter constructs LTRBLVGFP.** (A) Transient  
288 transfection of the reporter cell line with Tax expressing vector. CC81LTRBLVGFP cells  
289 were transfected with the pcDNABLVTAX-FLAG or pcDNA3.1 empty vector (control).  
290 The %GFP+ cells were determined by flow cytometry. A representative dot plot of three  
291 replicates is shown. Bars represent the mean + SD of 3 replicates. \*\*\*p=0.0005, ns p=0.88  
292 One-way ANOVA test. GFP expression was evaluated by fluorescent microscopy in  
293 CC81LTRGFP cells 48 h after transfection with Tax or the empty vector. Untransfected  
294 cells are shown as control. Scale bar, 100  $\mu$ m. (B) The effect of INF- $\alpha$  on BLV LTR  
295 promoter activity was evaluated in CC81LTRBLVGFP reporter cells. Cells were  
296 cultivated for 48 h without or with 75, 125, 250, 500 or 1000 U/mL of INF- $\alpha$ , and the  
297 percentage of GFP+ cells was evaluated by flow cytometry. Bars represent the mean +  
298 SD of 2 replicates. One-way ANOVA test. (C) The effect of deacetylase inhibitors (VPA  
299 and TSA) on BLV LTR promoter activity was evaluated in CC81LTRBLVGFP reporter  
300 cells. Cells were cultivated for 24 h without (CTRL) or with 10 mM VPA or 500 nM TSA  
301 and the percentage of GFP+ cells were evaluated by flow cytometry. Bars represent the  
302 mean + SD of 3 replicates. \*\*p=0.007 \*p=0.01 One-way ANOVA test. For statistical  
303 analysis, GraphPad Prism 9.1.0 was used.

304 Validation of the reporter cell line CC81LTRBLVGFP upon BLV infection

305 The next step was to evaluate the effect of BLV infection on LTR GFP expression by co-  
306 culturing the reporter cell line CC81LTRBLVGFP with persistently infected FLKBLV  
307 or BL3.1 cell lines. We first optimized the co-culture between FLKBLV cells and the  
308 reporter cell line, assessing different ratios (1:1, 1:2, 1:4 FLKBLV:CC81LTRBLVGFP),  
309 and fluorescent syncytia (cell-cell fusion) formation was evaluated 24 and 48 h later by  
310 fluorescent microscopy. We determined that cell-cell fusion (the number of fluorescent  
311 syncytia) increases with the proportion of reporter cells to BLV-infected cells,  
312 particularly at the 4:1 ratio. Fluorescent syncytium can be detected 24 h after co-culture,  
313 but the number of syncytia and nuclei per syncytium increases at 48 h (see  
314 **Supplementary Figure 2**). Thus, we conclude that the optimal conditions for co-culture  
315 between the FLKBLV and CC81LTRBLVGFP cell lines involve a 1:4 ratio, respectively,  
316 maintained for 48 hours (**Figure 3, upper panel**). Regarding the BL3.1 cell line co-  
317 culture, we first evaluated the same conditions used with FLKBLV (1:4 and 48 h).  
318 However, under these conditions, we did not observe the formation of fluorescent  
319 syncytia. Since BL3.1 cells grow in suspension, experimental conditions, such as the  
320 number of cells and the co-culture time were modified. Thus, we used  $5 \times 10^4$   
321 CC81LTRBLVGFP cells per well (plated 24 h before co-culture),  $1 \times 10^5$  BL3.1 cells per  
322 well and 72 h co-cultured period. With this experimental set up, stronger fluorescence  
323 and larger syncytium were observed (**Figure 3, lower panel**). In summary, our reporter  
324 cell line was able to express GFP and form syncytia when subjected to co-culture assays  
325 using cell lines persistently infected with BLV.



326

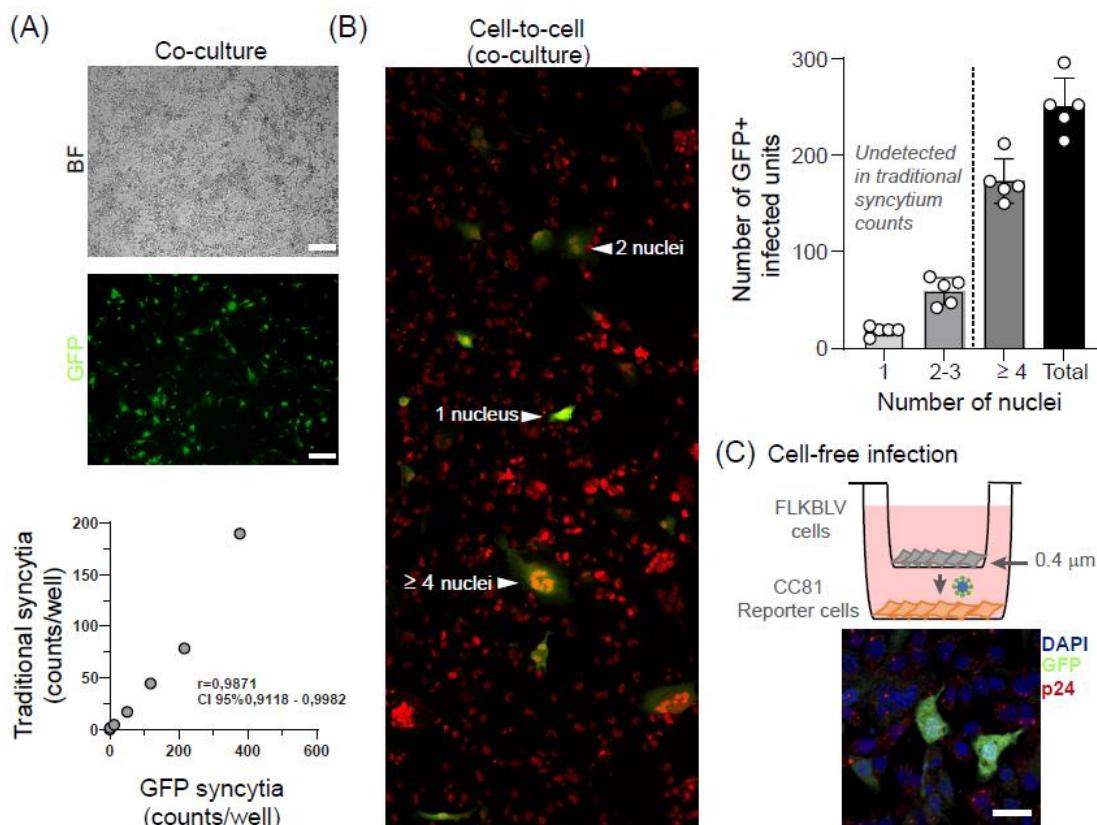
327 **Figure 3. Co-culture of C81LTRBLVGFP reporter cell with the BLV infected cell**  
328 **lines FLKBLV or BL3.1.** CC81LTRBLV cells were co-cultured in a 4:1 ratio with  
329 FLKBLV cells (above) or BL3.1 BLV cells (below) for 48 or 72 h, respectively.  
330 Syncytium expressed GFP was evaluated by fluorescent microscopy, and images were  
331 analyzed using Image J software. Representative images from three independent  
332 experiments are shown. BF: Bright field. Scale bar: 100  $\mu$ m.

333 *CC81LTRBLV reporter cell system is useful to measure cell-to-cell (via cell-fusion) and*  
334 *cell-free BLV infection*

335 To compare our BLV infection reporter system with traditional syncytia formation assay,  
336 CC81LTRBLVGFP reporter cells were co-cultured with FLKBLV cells in a 4:1 ratio for  
337 48 h. BLV infection was measured concurrently by counting the number of syncytia using  
338 bright-field microscopy and the number of GFP+ syncytia using fluorescent microscopy  
339 in the same field. We found a significant correlation between the GFP syncytia number  
340 using the reporter cell line and the syncytia number assessed by the traditional syncytia  
341 formation assay (Pearson correlation coefficient,  $r = p < 0.0001$ ) (**Figure 4 A, Supplementary Table 1**). However, with our reporter cell line, we were able to  
342 distinguish GFP+ syncytia that were not detected with the traditional method (>30%). In  
343 addition, since traditional syncytia formation assays consider multinucleated cell with 4  
344 or more nuclei, we detected individual infected cells and multinucleated cells with fewer  
345 than 4 nuclei (herein termed “GFP+ infected units”). Taking these results into  
346 consideration, we utilized an automated image acquisition system (25 images per well)  
347 using confocal microscopy and developed a software that allows for detection and  
348 counting of GFP+ infected units. This automated method allows detection of individual  
349 infected and multinucleated cells with less than 4 nuclei (30% of total counts) indetectable  
350 by traditional method (**Figure 4 B, Supplementary Figure 3, Supplementary Table 2**).  
351 To test the ability of CC81LTRBLVGFP cells to detect BLV cell-free infection, we  
352 initially used FLKBLV supernatant containing BLV viral particles to infect the cell line;  
353 however, no GFP+ cells were observed. To further investigate the ability of our reporter  
354 cell line to detect cell-free BLV infection, we performed a transwell experiment between  
355 the CC81LTRBLVGFP reporter cells and an insert with FLKBLV cells producing BLV  
356 particles and immunofluorescence. Here, we detected CC81 reporter cells infected with  
357 BLV by detecting the capsid protein and individual cells with a strong GFP signal using  
358

359 fluorescent microscopy (**Figure 4C**). The number of GFP+ cells is lower than when cell  
360 contact is allowed.

361



362

363 **Figure 4. CC81LTRBLV reporter cell system is useful to measure cell-to-cell (via**  
364 **cell-fusion) and cell-free BLV infection.** (A) The correlation between the stained  
365 syncytia counts in traditional assay and the fluorescence syncytia counts.  $5 \times 10^4$   
366 CC81LTRBLVGFP cells were co-cultured with 0, 90, 180, 750, 1500, 3000 and 6000  
367 FLKBLV cells in a 24-well plate at  $37^\circ\text{C}$  for 48 h after that, the cells were fixed with 4%  
368 PFA. The number of syncytia per well, defined as multinucleated cells with 4 or more  
369 nuclei, was counted (in the same field) by two independent experimenters in bright field  
370 (BF) and fluorescence images. Syncytia count measured by the traditional syncytia assay  
371 was plotted against syncytia number determined by GFP expression and the correlation  
372 was evaluated with the Pearson correlation test,  $r=0.9871$ . Scale bar 200  $\mu\text{m}$ . (B) A  
373 representative section of an image acquired automatically obtained from the co-culture  
374 between CC81LTRBLVGFP cells and FLKBLV cells is shown. GFP+ cells with 1, 2 or  
375 more than 4 nuclei are indicated (white triangles). DAPI staining is shown in red. The  
376 number of GFP+ cells per well was measured with 1, 2-3, 4 nuclei, and total. Bars  
377 represent the mean + 3SD of five independent experiments. (C) A BLV cell-free infection  
378 experiment was performed using the transwell system. After 48 hours of culture, the cells  
379 were fixed with 4% PFA and subjected to immunofluorescence with monoclonal anti-p24  
380 (to detect BLV capsid protein, in red). Cell nuclei were stained with DAPI (blue). Cell  
381 infection was assessed using Zeiss LSM800 confocal microscopy. Scale bar, 25  $\mu\text{m}$ .

382

383

384 **DISCUSSION**

385 It is well established that purified BLV virions exhibit inefficient infection of  
386 susceptible cells and transmission in cell culture is greatly improved when cell-to-cell  
387 contact is allowed. When BLV-infected cells are co-cultured with susceptible cells, they  
388 undergo cell-to-cell fusion events (syncytia) in cell culture [18], [16]. This syncytium  
389 formation assay is a primary tool in studying BLV infectivity [15, 17, 30]. Although this  
390 method is highly robust, the complexity of visually counting syncytia poses a significant  
391 technical challenge, especially in high-throughput screening assays.

392 An alternative to this challenge is the development of reporter cell lines capable  
393 of emitting a fluorescence or luminescence signal in response to viral infection. In this  
394 sense, an improved reporter cell system utilizing the BLV full-length LTR [22] or  
395 LTR/U3 promoter [19–21] was described. Recently, Sato *et al.*, [21] established a reporter  
396 cell line, namely CC81-GREM, harboring the EGFP reporter gene under the control of  
397 the GRE-mutated LTR-U3 promoter. This assay was used to evaluate the BLV-infected  
398 white blood cells, the capacity of neutralizing antibodies, and the infectious potential of  
399 cells in milk from BLV-infected dams, demonstrating greater sensitivity in quantifying  
400 BLV infectivity than traditional cell lines used in conventional cell fusion assays.

401 In this study, we constructed a reporter cell line to measure BLV infectivity using  
402 a lentiviral vector system due to the ease of establishing stable cell lines [31]. This cell  
403 line has stably integrated into its genome a construct containing the viral BLV full-length  
404 5'LTR associated with the fluorescent protein GFP coding region. The LTR promoter is  
405 activated in the presence of the viral protein Tax (transcriptional activator), which is only  
406 present when viral infection of the reporter line occurs.

407 We first characterize the functionality of our reporter cell line through Tax-  
408 dependent and Tax-independent transactivation. As previously described, the 5'LTR  
409 region present in the BLV genome contains protein binding regulatory sites and is  
410 responsible for virus integration and replication. This mechanism is significant because  
411 the BLV LTR-U3 regulates BLV replication induced by Tax protein [32, 33]. In this  
412 regard, through transient transfection of the CC81LTRBLVGF reporter cell line with a  
413 Tax expression plasmid, we demonstrated that the Tax protein efficiently activated the  
414 CC81LTBLVGF cell line, observing cells expressing GFP. This result is consistent with  
415 previous studies [22], [20]. On the other hand, it has been demonstrated that using HDACi  
416 can induce the expression of both viral and cellular genes. In this context, previous studies  
417 using reporter-based assays have shown that BLV expression is upregulated in response  
418 to TSA and VPA stimulation in a Tax-independent manner [26]. In this sense, as  
419 expected, low levels of GFP expression were observed when stimulating our reporter cell  
420 line with VPA and TSA. A binding site for interferon (IFN) regulatory factors 1 and 2  
421 (IRF-1 and IRF-2) has been identified in a transcriptional enhancer in the LTR-U5 region,  
422 and the possibility of BLV LTR basal activation by IFN in a Tax-independent manner has  
423 been suggested [28]. One of the questions we asked ourselves was if stimulation of the  
424 reporter cell line with IFN was sufficient to activate the promoter. We showed that high  
425 concentrations of IFN do not induce GFP expression, suggesting that our reporter cell line  
426 (containing the complete LTR) could be used to study the effect of IFN at different stages  
427 of BLV infection.

428 We validated the CC81LTRBLVGF reporter cell line following BLV infection  
429 by assessing its ability to form syncytia through cocultivation with FLKBLV or BL3.1

430 cell lines persistently infected with BLV. The quantification of fluorescent syncytia using  
431 CC81LTRBLVGFP strongly correlated with traditional syncytium counts ( $r=0.9871$ ).  
432 However, when the reporter cell line was cocultured with FLKBLV, we observed both  
433 single GFP+ cells and fluorescent multinucleated cells with 2 or 3 nuclei. These  
434 observations suggest that CC81LTRBLVGFP can detect infection events that are not  
435 distinguishable in traditional syncytium formation assays improving the robustness of our  
436 method.

437 In this regard, we integrate automated scanning and image acquisition using a  
438 confocal microscopy, with the development of an analysis software that allows us to  
439 detect and count single GFP-positive cells or fluorescent multinucleated cells with 2 or  
440 more nuclei. This approach significantly accelerates sample analysis compared to  
441 traditional visual counting methods. Through coculture assays between  
442 CC81LTRBLVGFP and FLKBLV, we observed that BLV predominantly spreads  
443 through cocultured cells via cell fusion transmission. Additionally, we evaluated the  
444 usefulness of the reporter cell line in measuring cell-free infection through a transwell  
445 assay. We successfully detected GFP cells and viral particles within the reporter cell line  
446 by labeling a BLV capsid. However, we cannot exclude the possibility that Tax is secreted  
447 and can enter CC81 LTRBLVGFP to transactivate the LTRGFP. Furthermore, Tax has  
448 been identified in exosomes released from cells infected with HTLV [34]

449 It has been previously documented that retrovirus spread in cultured cells and  
450 tissues via two routes: through cell-to-cell contact and cell-free mode through the  
451 extracellular environment and infecting new cells [13, 35–37]. Cell-to-cell transmission  
452 typically involves tight cell-cell contact, such as virological synapses, biofilms, or the  
453 formation of nanotubes. On the other hand, cell-cell fusion has been identified as an  
454 alternative pathway for cell-to-cell transmission [12, 38]. Cell-cell fusion has been  
455 described in HIV and HTLV. However, the relevance of this property for virus spread *in*  
456 *vivo* is unclear, and there is no clear evidence that syncytia formation can enhance HIV-  
457 1 dissemination *in vivo* [13]. BLV is naturally and mainly transmitted via cell-to-cell  
458 rather than cell-free mechanisms, as the extent of cell-free infectivity of virions in the  
459 blood of BLV-infected cattle is significantly lower [39]. Currently, very little is known  
460 about the mode of transmission between cells or the efficiency of cell-free infection for  
461 BLV. Our results support that cell-to-cell transmission is the primary mechanism for BLV  
462 spread in cell co-culture. In addition, we observed that more than 90% of infection events  
463 (GFP+ infected units with >1 nuclei) involved cell-cell fusion. In this regard, our reporter  
464 cell line, combined with the development of analysis software, is a useful tool for  
465 understanding the role of cell fusion in BLV transmission.

466 Although the co-culture of infected and target cells is a frequently used and  
467 efficient method for studying BLV infection, some concerns remain. For example, it is  
468 challenging to distinguish between donor and acceptor cells. Cell-free BLV infection  
469 models allow for more controlled experimental conditions. Further exploration of the  
470 mechanisms involved in cell-to-cell or cell-free transmission of BLV is necessary to  
471 better understand its pathogenesis and biology. Systems that can measure both  
472 mechanisms of infection, like our reporter cell line, could be handy for this.

473 Finally, our reporting system could be implemented to evaluate the infective  
474 potential of BLV present in different biological samples of importance in the veterinary  
475 management of cattle, such as colostrum, semen, mucous fluids, and more. Additionally,  
476 it could be used to assess the effect of different disinfectant measures on infectivity, such  
477 as heat treatment, specific disinfectants, and other methods of viral inactivation. Lastly,  
478 it could also be employed in the evaluation of molecules with antiviral activity (e.g., BLV

479 protease inhibitor compounds) or molecules capable of neutralizing infection (e.g.,  
480 specific monoclonal antibodies and their smaller derivatives).

## 481 **AUTHOR CONTRIBUTIONS**

482 NO-D and OP: conceptualization and designed the project; NO-D and FR designed the  
483 experiments; NO-D and FR performed the experiments; NO-D, FR, MF, FC, CO and  
484 FDN analyzed the data, AV: design image processing software. All authors contributed  
485 to the preparation of the manuscript.

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## 498 **CONFLICT OF INTEREST**

499 The authors declare that have no conflict of interest.

## 500 **ETHICAL APPROVAL**

501 This article does not contain any studies with animals performed by any authors.

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