



Functional analysis of the baculovirus per os infectivity factors 3 and 9 by imaging the interaction between fluorescently labelled virions and isolated midgut cells

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1                   **Functional analysis of the baculovirus**  
2                   ***per os* infectivity factors 3 and 9 by imaging the**  
3                   **interaction between fluorescently labelled virions**  
4                   **and isolated midgut cells**

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13  
14                   **Abstract**

15                   Baculovirus occlusion derived viruses (ODVs) contain ten known *per os* infectivity factors  
16                   (PIFs). These PIFs are crucial for midgut infection of insect larvae and form, with the exception  
17                   of PIF5, an ODV entry complex. Previously, R18 dequenching assays have shown that PIF3 is  
18                   dispensable for binding and fusion with midgut epithelial cells. Oral infection nevertheless fails  
19                   in absence of PIF3. PIF9 has not been analysed in much depth yet. Here, the biological role of  
20                   these two PIFs in midgut infection was examined by monitoring the fate of fluorescently  
21                   labelled ODVs when incubated with isolated midgut cells from *Spodoptera exigua* larvae.  
22                   Confocal microscopy showed that in absence of either PIF3 or PIF9, the ODVs bound to the  
23                   brush borders, but the nucleocapsids failed to enter the cells. Finally, we discuss how the results  
24                   obtained for PIF3 with dequenching assays and confocal microscopy can be explained by a  
25                   two-phase fusion process.

27 **Full text**

28 The *Baculoviridae* are a family of arthropod-specific viruses with large circular DNA genomes  
29 (1). Insect larvae become orally infected when food is consumed that is contaminated with viral  
30 occlusion bodies (OBs). Infectious occlusion derived viruses (ODVs) are embedded in the  
31 protein matrix of these OBs and are released in the alkaline environment of the insect's midgut  
32 lumen. The first step in the infection process is binding and fusion of the ODV-envelope with  
33 the plasma membrane of the microvilli of midgut epithelial cells (2, 3). After primary infection  
34 of the midgut epithelium, the virus spreads systemically to other tissues by means of a different  
35 virus phenotype, the budded viruses (BVs).

36 Infection of the midgut epithelium cells by ODVs requires at least ten different ODV-envelope  
37 proteins: the *per os* infectivity factors or PIFs (4-16). Deletion of any of the *pif*-genes abolishes  
38 the oral infectivity of the ODVs without affecting their formation or occlusion into OBs. The  
39 production and infectivity of BVs remain unaffected by *pif*-gene deletions. All PIFs, except  
40 PIF5, participate in the formation of the ODV entry complex (14, 16-19). The initial steps of  
41 midgut infection have previously been studied by fluorescence dequenching assays with R18-  
42 loaded ODVs and brush border membrane vesicles (7, 20, 21). These studies showed that PIF0,  
43 1 and 2 were important for the binding of ODVs to the vesicles, while PIF3 was dispensable  
44 for ODV binding and fusion (7, 20, 21). These findings would indirectly imply that the binding  
45 and fusion of the ODV envelope with the cellular plasma membrane do not require the presence  
46 of the entry complex, as the entry complex fails to form in absence of PIF3 (18), a concept that  
47 is not easy to understand. To gain more insight into the biological function of PIF3 we  
48 visualized the interaction of ODVs with the brush border of isolated midgut cells of *S. exigua*  
49 larvae using confocal microscopy imaging, as previously described (22). We also examined the  
50 role of the recently discovered PIF9 in this analysis. Like PIF3, PIF9 is also needed for entry  
51 complex formation (15).

52 *Generation of fluorescent ODVs with an inactivated pif3 or pif9 gene*

53 For this study, we produced fluorescently labelled ODVs of *Autographa californica* multiple  
54 nucleopolyhedrovirus (AcMNPV) lacking either PIF3 or PIF9. The *pif3* gene of the bacmid  
55 bMON14272 derived from the E2 strain (23, 24) was deleted by insertion of the  
56 chloramphenicol acetyl transferase (*cat*) resistance gene via  $\lambda$ -red mediated homologous  
57 recombination (25). To this aim, a *cat*-amplicon, with arms homologous to sequences flanking  
58 the *pif3* region to be deleted, was produced by PCR with Phusion polymerase (Thermo Fischer).  
59 We used the pCR-TOPO-loxLE-*cat*-loxRE plasmid (kindly provided by L. Galibert, at the time

60 at Généthon, France) as template for the PCR in combination with the previously described  
61 primers (26). *E. coli* MW003 cells (MW001 (27) with the *bla* gene removed), carrying the  
62 AcMNPV bacmid were transformed with the *cat*-amplicon. The expression of the  $\lambda$  RED  
63 recombinase (25) was activated in these bacterial cells by heat induction at 42°C for 10 min.  
64 The transformed bacteria were incubated on LB-agar plates in the presence of chloramphenicol  
65 (34  $\mu$ g/ml) and kanamycin (50  $\mu$ g/ml) for up to two days at 32°C. The insertion of the *cat*-gene  
66 in the *pif3*-ORF was confirmed by PCR (**Supplementary Figure 1**) with primers that annealed  
67 outside the recombined region (5'-CAGTTGTAAACGCGTCTGTAC-3' and  
68 5'-CATGTTGAACTTTTGGCAAATACTTATTC-3'). The *pif3* deletion bacmid was  
69 electroporated into DH10 $\beta$  *E. coli* cells that contained the transposase helper plasmid  
70 pMON7124 (24). These resulting cells were used for site-directed transposition using the  
71 pFastBac-*polh-egfp:vp39* plasmid (22) with the aim to restore the polyhedrin (*polh*) gene to  
72 allow OB formation and to introduce a *gfp:vp39* fusion construct under control of the *p10*  
73 promoter to label the nucleocapsids (**Fig. 1**, del-*pif3*). The *pif9* (*ac108*) mutant bacmid  
74 previously described (15) was transposed in a similar way to obtain mut-*pif9* (**Fig. 1**). BV-  
75 stocks were generated by transfection of Sf21 cells with the bacmids, using Expres2TR  
76 transfection reagent (Expres2ion Biotechnologies). High titre BV stocks were obtained by an  
77 additional infection round in Sf21 cells. OBs were produced by infection of TnH5 cells with  
78 BVs at an MOI of 2 and isolated as described before (15). The fluorescent “wild type” (Wt)  
79 virus used as control in this study has been described before (22). Western blot analysis  
80 confirmed the absence of PIF3 in the envelope of *pif3*-deletion ODVs, while all the tested  
81 viruses were shown to incorporate the eGFP:VP39 fusion protein into their virions (**Fig. 2**).

## 82 *Infectivity assays in S. exigua larvae*

83 First the infectivity of the BVs of the fluorescent *pif3* and *pif9* mutant viruses was determined  
84 by injection of 1  $\mu$ l BV-stock ( $3,5 \times 10^7$  TCID<sub>50</sub> / ml) into the hemocoel of L4 *Spodoptera*  
85 *exigua* larvae, using an Humapen Luxura insulin pen. When the larvae were injected with Wt  
86 or one of the mutant BVs stocks, fluorescence was observed in at least 11 out of 12 larvae for  
87 each virus at 3 days post infection (dpi) (**Fig. 3a**), and almost all larvae finally died of the  
88 infection (**Table 1**), indicating that the Wt and *pif* mutant viruses replicated efficiently in the  
89 larvae when bypassing the midgut. The oral infectivity of the del-*pif3* and mut-*pif9* OBs was  
90 determined in separate bioassays, in which the del-*pif3* was tested in duplicate (as others already  
91 tested *pif3* deletion mutants extensively) and the *pif9* mutant in triplicate. When L3 instars were  
92 orally inoculated with OB stocks ( $3 \times 10^8$  ml<sup>-1</sup>) using droplet feeding, only the larvae that were  
93 fed with the Wt OBs showed fluorescence (**Fig. 3b**) and significant mortality (**Table 1**). Hence,

94 the fluorescently labelled del-*pif3* and mut-*pif9* viruses have the same phenotype as previously  
95 described for the non-fluorescent variants (15, 26).

#### 96 *Interaction between ODVs and the brush borders of midgut epithelial cells*

97 To visualize the interaction between (mutant) ODVs and the epithelial cell brush border,  
98 midguts of twenty *S. exigua* L5 instar larvae were dissected by cutting of the larva's head and  
99 posterior abdominal segment, after which the midgut was pulled out and the peritrophic  
100 membrane was removed. The midguts were collected in Grace's medium, supplemented with  
101 50 µg/ml gentamycin, rinsed twice with this medium and further incubated in Graces medium,  
102 containing 2 mg/ml collagenase from *Clostridium histolyticum* (Sigma, blend type H), for 90  
103 min. at 27°C under vigorous shaking at 850 rpm. The cells were separated from the remaining  
104 midgut tissues by filtering through a 200 µm meshwork of nylon, fixed at the bottom of a  
105 microtube. The primary midgut cells were collected and pelleted by centrifugation at 200 g for  
106 10 min, and subsequently washed three times with Grace's medium. In parallel, the ODVs were  
107 isolated in a slightly modified way from what has been described before (19), in which ODVs  
108 were released from larva-derived OBs by EDTA-containing DAS-buffer and were further  
109 purified by ultracentrifugation in a sucrose gradient. For our experiments, the ODVs were  
110 released from  $\sim 3.0 \times 10^8$  cell culture-derived OBs by a 10 min incubation in EDTA-free DAS-  
111 buffer (0.1 M Na<sub>2</sub>CO<sub>3</sub>, 166 mM NaCl; pH 10.5) and subsequent disruption of the OB membrane  
112 by two sonication bursts.

113 This alkaline ODV suspension was further used to suspend the isolated midgut cells after the  
114 last washing step and the resulting cell-virus suspension was incubated on ice for 15 min to  
115 allow the ODVs to bind to the brush borders prior to imaging. The cell-virus suspension (50-  
116 100 µl) was transferred to a circular compartment on the object holder, which was created by  
117 placing a silicone spacer between the object- and cover glass. The midgut cells were examined  
118 for (bound) ODVs in the brush border area by detection of GFP fluorescence by confocal  
119 microscopy (Zeiss LSM-510), using a 63x oil objective lens and excitation with an Argon laser.  
120 The ODV-midgut cell interaction was then analysed by construction of a z-stack consisting of  
121 15 – 20 optical slices of 600-700 nm each in a time lapse experiment with a duration of 20 min.  
122 It turned out to be crucial to add limited amounts of ODVs to be able to clearly discriminate  
123 individual ODVs over time. The images were analysed by using Zen 2.3 lite (blue edition)  
124 software.

125 First, more than twenty Wt ODVs were followed over time in their interaction with eight  
126 separate midgut cells to validate the experimental conditions (**Table 2**). These ODVs were

127 detected in the brush border for 2 to 6 minutes before they disappeared from the cell surface  
128 (**Fig. 4a**). About half of the ODVs disappeared within 2-4 minutes after the start of the time  
129 lapse experiment. As the ODVs under study could not be found in the surrounding focal planes,  
130 it was concluded that the nucleocapsids had entered the midgut cells. Apparently, the  
131 fluorescent signal disappeared by decomposition of the grouped nucleocapsids of the multiple  
132 capsid ODVs upon cell entry to allow transport of individual nucleocapsids through the  
133 microvilli on their way towards the nucleus, as was observed before in electron microscopy  
134 studies (2, 3, 28). It should be noted that it took longer for the Wt ODVs to enter the cells under  
135 the present experimental conditions than previously described (22). This might be explained by  
136 the usage of ODVs that were released from cell culture-derived OBs (C-ODVs) instead of OBs  
137 from infected larvae (L-ODVs), as used in the previous study. Infection with C-ODVs might  
138 initially be hampered by remnant polyhedrin protein that sticks to the virus envelope, but will  
139 be degraded by co-occluded host proteases in L-ODV preparations (29-31). However, as we  
140 now know that the remaining PIF-proteins in *pif* deletion mutants are highly sensitive to  
141 degradation by such co-occluded proteases during OB-dissolution (19), we decided to use C-  
142 ODVs for our experiments. The remnant polyhedrin was probably degraded by digestive  
143 enzymes of the primary midgut cell (32) when the virus and cell suspensions were mixed.

144 The experiment was repeated with mutant ODVs, for which seven ODVs of the *pif3* deletion  
145 mutant and fourteen of the *pif9* mutant were analysed, each on four distinct midgut cells  
146 (**Table 2**). The ODVs of both mutants were detected at the cell surface until the end of the  
147 measurements, even after continuous excitation with the laser for 20 minutes of the eGFP:VP39  
148 fusion protein (**Fig. 4b**). Hence, the disappearance of wild type ODVs could not be explained  
149 by fading of the fluorescent signal by continuous excitation of the eGFP:VP39 fusion protein  
150 with the laser. These observations show that in absence of PIF3 or PIF9 the ODVs are still able  
151 to bind to the brush borders of midgut epithelial cells (**Fig. 4b**), but the nucleocapsids fail to  
152 enter the cell.

153 *PIF3 and 9 might be required to overcome an intermediate hemifusion state during midgut*  
154 *infection.*

155 Two different phenotypes regarding binding and fusion have been observed before with R18  
156 dequenching assays using brush border vesicles with various mutant ODVs. It had been found  
157 that PIF0, 1 and 2 are important for ODV binding, while PIF3 is dispensable for binding and  
158 fusion (7, 20, 21). The ODVs of the *pif3* deletion mutant nevertheless fail to establish an midgut  
159 infection. Our observations show that in absence of PIF3, the ODVs bind to the brush border

160 (Fig. 4b), just as indicated before by the R18 dequenching assays (7). Whether the ODV-  
161 envelope of this mutant also fused with the brush border plasma membrane could not be  
162 confirmed with the currently used assay. Our observations nevertheless show that the  
163 nucleocapsids stayed together and failed to enter the cell. This indicates that PIF3 is indeed  
164 required for the translocation of the nucleocapsids into the host cell as proposed by others (7).  
165 However, it has been observed with mutants of other viruses that R18-dequenching also takes  
166 place after hemifusion of the lipid membranes, in which only the outer leaflets of the bilayers  
167 fuse (33, 34). It could therefore be reasoned that in absence of PIF3, only the outer leaflets fuse  
168 and that this PIF is required to overcome this hemifusion intermediate state (Fig. 5). This might  
169 explain why the R18-dequenching assays with the *pif3* deletion mutant suggested that  
170 membrane fusion took place, while the infection of the midgut epithelium failed.

171 ODVs that lacked PIF9 also bound to the epithelial cell brush borders. Whether the binding of  
172 these mutant ODVs is indeed not affected quantitatively is not known, as the live imaging assay  
173 is biased for ODVs that bind to the brush borders; i.e. non-interacting ODVs were not analysed  
174 in the time lapse experiments. However, as we regularly encountered bound ODVs, it is  
175 reasoned that binding is not significantly affected by the absence of PIF9. Based on our assay  
176 we cannot say for sure whether PIF9 is involved in fusion, but our observations nevertheless  
177 show that midgut cell entry of the nucleocapsids failed in absence of PIF9, as shown previously  
178 for PIF1 and 2 (22) and now also for PIF3.

179 Despite the fact that we know that both mutants are unable to form the ODV entry complex (15,  
180 18), the mutant ODVs were nevertheless found to be able to bind to the brush border of the  
181 isolated midgut cells. Formation of the entry complex is apparently not important for the ODV  
182 binding properties as long as PIF0, 1 and 2 are present in the ODV envelope. It was also  
183 observed for both *pif 3* and *pif 9* mutants that the entry of the nucleocapsids was abolished. It  
184 could be concluded from these observations that formation of the entry complex is crucial for  
185 early stages of midgut infection to complete the membrane fusion process at the epithelial cell  
186 brush border and/or for the translocation of the nucleocapsids into the cytosol.

187

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295

296

297 **Table 1:** Mortality of *S. exigua* larvae after injection with BVs or oral inoculation with OBs

<b>Injection with BVs</b>	<b>3.5 x 10<sup>7</sup> TCID<sub>50</sub> / ml</b>		
mock	0/12		
<i>del-pif3-egfp:vp39</i>	11/12		
<i>mut-pif9-egfp:vp39</i>	11/12		
<i>Wt-egfp:vp39</i>	12/12		
<b>Oral infectivity 3.0 x 10<sup>8</sup> OBs / ml</b>	<b>Exp. 1</b>	<b>Exp. 2</b>	
mock	0/14	0/12	
<i>del-pif3-egfp:vp39</i>	0/13	0/11	
<i>Wt-egfp:vp39</i>	12/13	11/11	
<b>Oral infectivity 3.0 x 10<sup>8</sup> OBs / ml</b>	<b>Exp. 1</b>	<b>Exp.2</b>	<b>Exp. 3</b>
mock	0/12	0/14	0/12
<i>mut-pif9-egfp:vp39</i>	0/15	0/12	0/12
<i>Wt-egfp:vp39</i>	12/15	12/13	10/12

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301 **Table 2:** Overview of the observations of fluorescent ODVs interacting with the brush border of primary

302 midgut cells.

<b>Virus</b>	<b>Nr. of observations</b>	<b>Nr. of observed cells</b>	<b>Average observation time before disappearance (min.)</b>
Wild type	21	8	4.5 ± 2
<i>del-pif3</i>	7	4	-
<i>mut-pif9</i>	14	4	-

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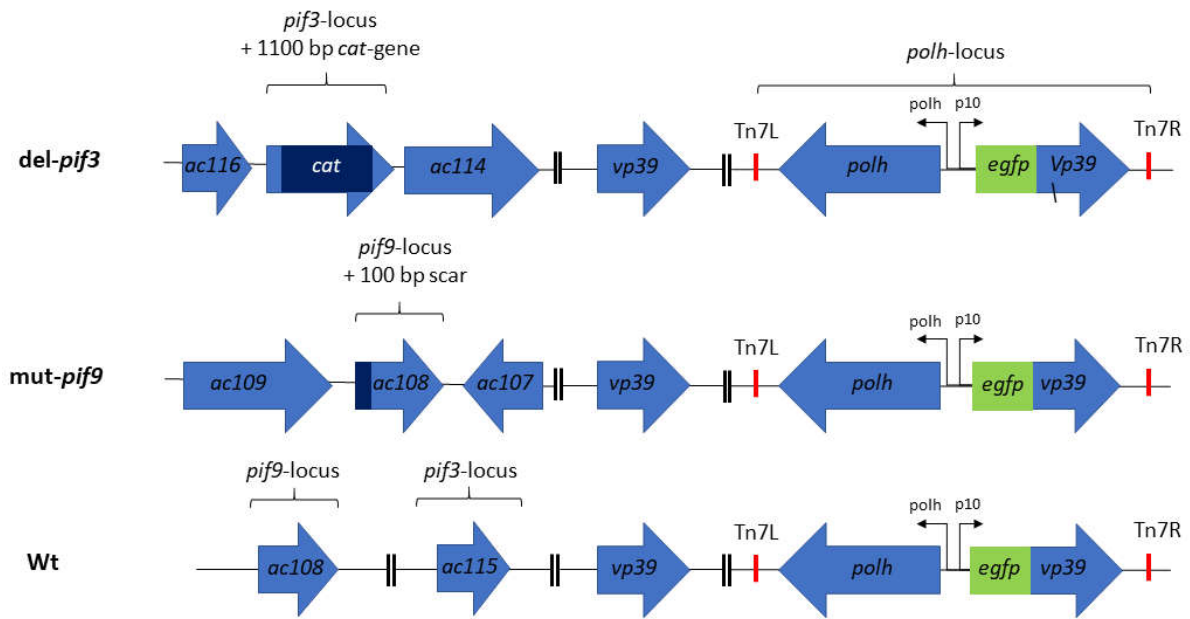
305

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309 **Figures with legends**



310

311 **Fig. 1.** Overview of the constructed bacmids. The *pif3* deletion and *pif9* mutant bacmids (the

312 latter is described in Boogaard *et al.*, 2019) were transposed with a pFastbac-*polh-egfp:vp39*

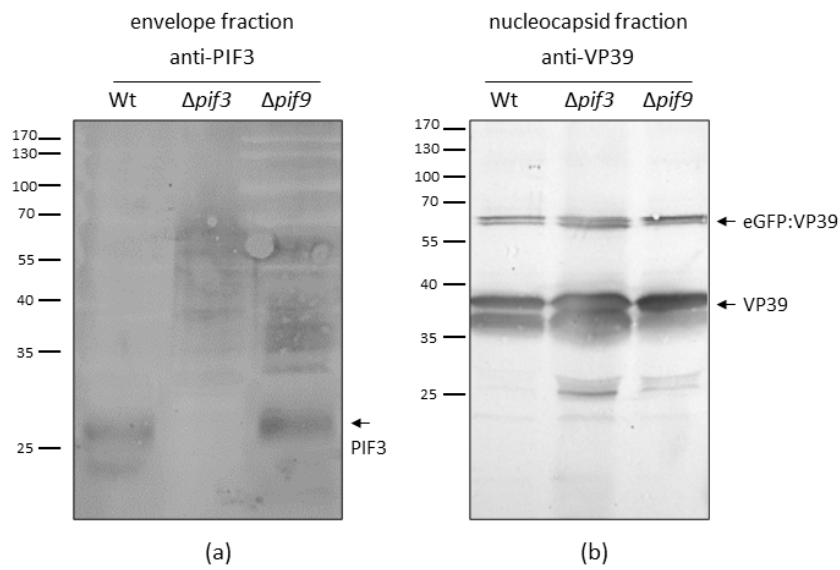
313 vector to enable the mutant viruses to produce OBs and fluorescently labelled nucleocapsids.

314 The fluorescent wild type virus was previously described in (22) and was used in this study as

315 wild type control.

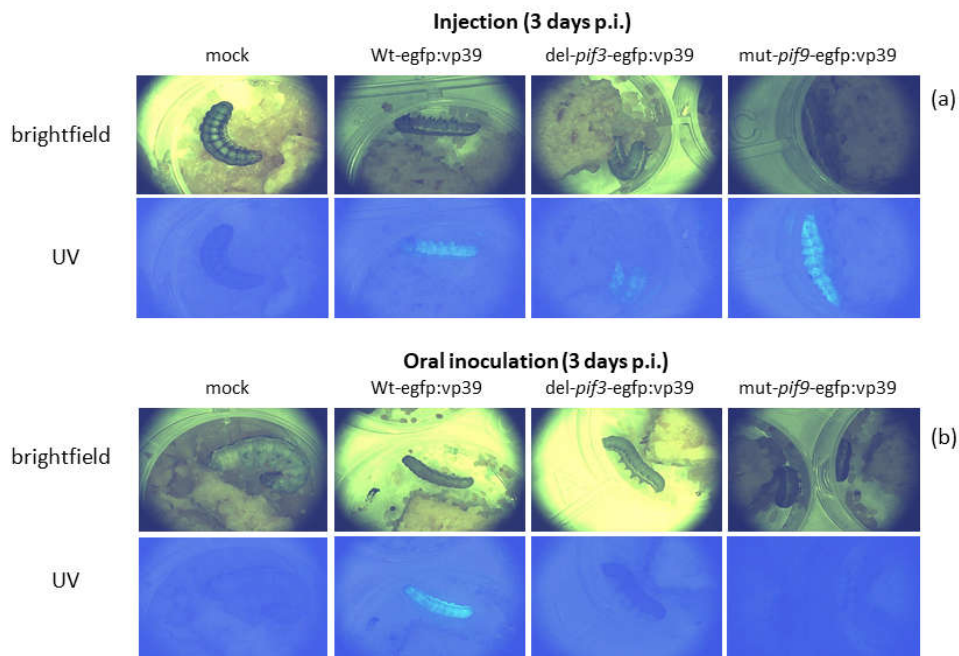
316

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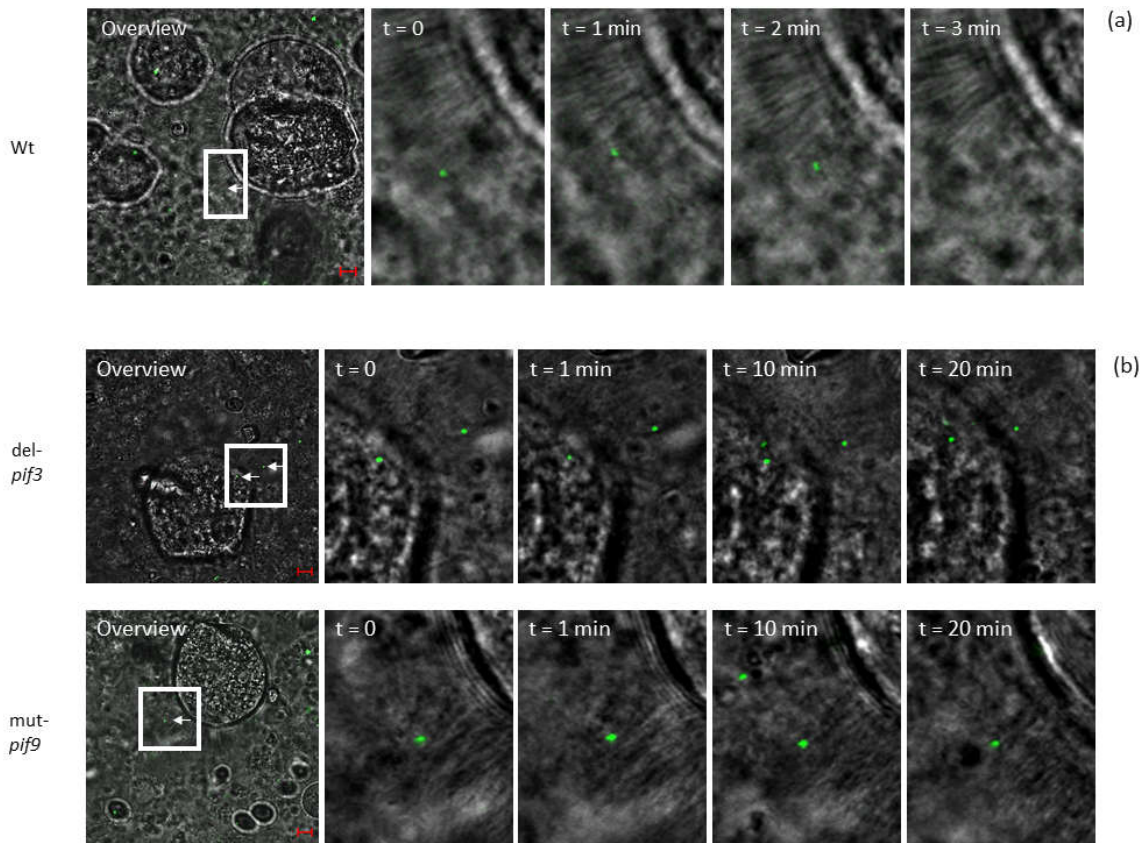
318

319 **Fig. 2.** Western blot analysis of the ODV-envelope and nucleocapsid fractions of wild type  
 320 (Wt), *pif3* deletion mutant and *pif9* mutant viruses. The envelope fractions were analysed with  
 321 antiserum against PIF3 (a) and the nucleocapsid fractions with VP39 antiserum (b).



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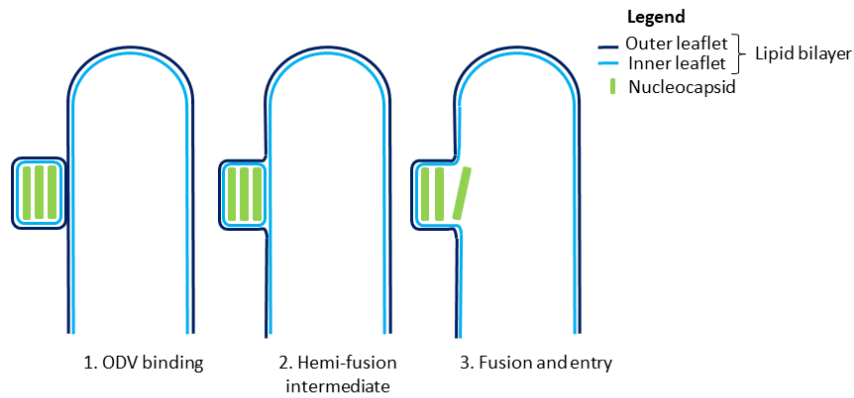
323 **Fig. 3.** Detection of viral infection. The larvae were examined under UV-light to observe  
 324 green fluorescence after injection with BVs (a) or oral inoculation with OBs (b). Green  
 325 fluorescence is indicative of GFP:VP39 production and hence for viral replication.



326

327 **Fig. 4.** Time lapse analysis of single fluorescently labelled ODVs on the brush border of a  
 328 primary midgut cell. Cells were incubated with ODVs from the wilt type (Wt) virus (a), the  
 329 *pif3* deletion mutant (b) or the *pif9* mutant (c) and these ODVs were followed by confocal  
 330 microscopy in their interaction with the brush border of a primary midgut cell. The red scale  
 331 bars correspond with 5  $\mu$ m. The Wt ODV shown in (a) disappeared within 3 minutes after the  
 332 start of the observation.

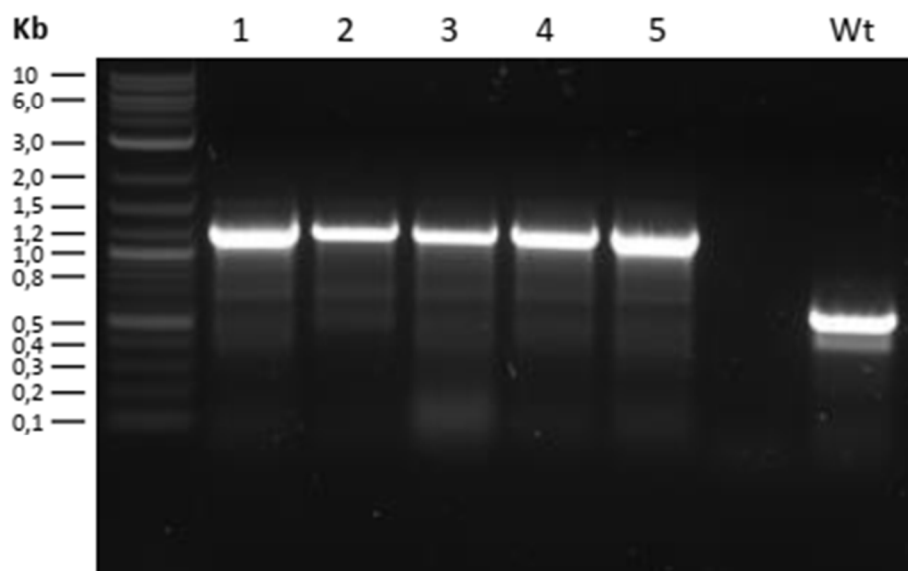
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335 **Fig. 5.** Model for the membrane fusion event during primary infection of the midgut  
 336 epithelium. In the first step (1) the ODV binds to the microvilli membrane. The fusion of the  
 337 outer and inner leaflets of the lipid bilayers are considered as separate events. First the outer  
 338 leaflets fuse leading to the formation of a hemifusion intermediate (step 2), followed by the  
 339 fusion of the inner leaflets and the release of the nucleocapsids into the cytosol (step 3). It is  
 340 hypothesized that in absence of PIF3, membrane fusion is stuck in the hemifusion state, and  
 341 hence, entry of the nucleocapsids into the host cell will be prevented.

PCR validation of the insertion of a  
chloramphenicol resistance (*cat*) gene  
into the *pif3*-ORF



**Fig. S1:** Confirmation of the insertion of the chloramphenicol acetyl transferase (*cat*) resistance gene into the *pif3*-locus by PCR. Insertion of the *cat*-gene resulted in the formation of an amplicon of 1500 bp (clone 1-5), while without insertion a 500 bp amplicon was produced (Wt). Clone 1 was used for further studies.

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