Pseudorabies Virus Fast Axonal Transport Occurs by a pUS9-Independent Mechanism

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Reactivation from latency results in transmission of neurotropic herpesviruses from the nervous system to body surfaces, referred to as anterior axonal trafficking. The virus-encoded protein pUS9 promotes axonal dissemination by sorting virus particles into axons, but whether it is also an effector of fast axonal transport within axons is unknown. To determine the role of pUS9 in anterior axonal trafficking, we analyzed the axonal transport of pseudorabies virus in the presence and absence of pUS9.

All herpesviruses cycle through periods of active and latent infection, with members of the Alphaherpesvirinae subfamily typically establishing latency in neurons of the peripheral nervous system. Retrograde transport results in entry into the nervous system, transmitting particles from neuron terminals to sensory and autonomic ganglia. Reactivation from the latent state results in newly assembled viral particles traveling from the ganglia to sites of innervation at body surfaces by anterior axonal trafficking (1). The resulting infections include presentations such as herpes labialis (herpes simplex virus 1 [HSV-1]) and shingles (varicella-zoster virus [VZV]). These viruses also encompass veterinary pathogens, including the well-studied pseudorabies virus (PRV) that serves as a model for severe neuroinvasive infections (2).

Anterograde axonal trafficking consists of two steps. Cargoes, including virus proteins, are sorted into the axon from the neuronal cell body (3). Once in the axon, the cargo moves to the distal axon terminal by microtubule-dependent fast axonal transport (4). The best characterized effector of herpesvirus axonal trafficking is the type II transmembrane protein pUS9 (5–11). PRV lacking pUS9 enters the nervous system by retrograde axon transport but, following replication in neurons, is attenuated for anterior axonal trafficking (4). The best characterized effector of herpesvirus axonal trafficking is the type II transmembrane protein pUS9 (5–11). PRV lacking pUS9 enters the nervous system by retrograde axon transport but, following replication in neurons, is attenuated for anterograde trafficking both in animals and in neuronal cell culture (9, 10, 12, 13). This defect is attributed to a decrease in viral particle sorting to axons, but whether pUS9 is an effector of fast axonal transport is unknown (14, 15). The presence of viral particles in axons and transmission of infection to cells at distal terminals indicate that rare anterograde trafficking events occur, but the scarcity of these events has precluded their analysis (9, 14). To determine whether pUS9 contributes to PRV fast axonal transport, we examined the transport of wild-type (WT) and ΔUS9 PRV particles that encode red fluorescent capsids (16). The fluorescent ΔUS9 mutant used in these studies was confirmed by restriction enzyme digest, sequencing across the deletion junction, absence of pUS9 expression, lack of viral particle accumulation at axon terminals resulting from anterograde transport in culture, and inability to spread by anterograde transportation within the rat visual system following intravitreal eye injection (Fig. 1 and Table 1) (17–21).

While envelope proteins, such as pUS9, are not expected to be effectors of the retrograde axon transport that occurs upon entry into nerve endings, an analysis was performed to confirm that this initial stage of neuronal infection was unperturbed (22, 23). Experiments of avian dorsal root ganglion (DRG) sensory neurons were cultured and infected ex vivo. Individual viral particles were imaged at 10 frames/s during the first hour postinfection (h.p.i.), and the kinetics of axonal transport were measured for particles moving more than 0.5 μm using the kymograph function in the MetaMorph software package (Molecular Devices) as described previously (n > 190) (Table 1) (21, 24–26). As expected, the WT and ΔUS9 viruses had equivalent retrograde transport profiles based on run lengths and run velocities (Fig. 2). Because pUS9 supports the sorting of viral particles from soma to axons following replication (5–11), examining its subsequent role in fast axonal trans-
port of particles to axon terminals was made difficult by the low frequency of these events (Fig. 1B; note the reduced number of viral particles accumulated at axon terminals following replication). We accomplished this analysis by extensive imaging of \( /H9004\) US9 PRV infections in isolated neurons in low-density cultures to capture a minimum of 30 transport events that were unambiguously moving away from an infected neuronal soma. Although WT events were far more frequent, an equivalent number were included for the comparative analysis (Table 1). The kinetics of \( /H9004\) PRV microtubule-based anterograde axonal transport were indistinguishable from the transport kinetics of the wild type (Fig. 3). These results indicate that the well-described reduction in anterograde spread noted for \( /H9004\) PRV cannot be directly attributed to a disruption of the fundamental ability of viral particles to engage in microtubule-based transport within axons (10, 14, 27).

The finding that PRV pUS9 interacts with KIF1A, a kinesin motor involved in the fast anterograde axonal transport of presynaptic vesicles (4), suggested a direct role for pUS9 in fast axonal transport (15). However, while a mutant pUS9 (Y49-50A) that does not interact with KIF1A (15) restricts the anterograde trafficking of capsids in axons, pUS9 (Y49-50A) itself remains competent for anterograde transport (14). Based on the current findings, an explanation for this perplexing observation can be offered: the pUS9-KIF1A interaction is required for sorting capsid-containing viral particles into axons but is dispensable for subsequent fast axonal transport within axons. Consistent with this, a protein that can sort into the axon independently from pUS9 and capsids, gM, also does not require pUS9 for axonal transport (28). Therefore, in the absence of pUS9, fewer viral particles are sent down axons due to inefficient sorting from the soma, but the probability that each individual particle will reach the axon terminal, as assessed by transport dynamics, remains unchanged. Inherent in this conclusion is the prediction that \( /H9004\) viruses should traverse long distances in axons to spread across anterograde neuronal circuits in vivo, albeit at reduced frequency due to inefficient sorting.

In apparent contradiction to this prediction, pUS9 is often regarded as an essential determinant for the anterograde trafficking of PRV. The principle evidence in support of this idea comes from the complete lack of anterograde transneuronal transmission

### TABLE 1 Viruses used in this study

<table>
<thead>
<tr>
<th>Virus</th>
<th>Reporter tag</th>
<th>Mutation</th>
<th>No. of rats with visual circuit transmission/total no. of rats&lt;sup&gt;c&lt;/sup&gt;</th>
<th>No. of transport events in avian DRG sensory neurons</th>
</tr>
</thead>
<tbody>
<tr>
<td>PRV-GS4284</td>
<td>UL25/mCherry</td>
<td>None</td>
<td>5/5</td>
<td>288</td>
</tr>
<tr>
<td>PRV-GS5469</td>
<td>UL25/mCherry</td>
<td>( /H9004)US9&lt;sup&gt;b&lt;/sup&gt;</td>
<td>0/5</td>
<td>192</td>
</tr>
</tbody>
</table>

<sup>a</sup> PRV-GS4284 was described previously (16).

<sup>b</sup> A stop codon was inserted after the start ATG, and codons 2 to 72 were deleted, thus removing all internal in-frame ATG codons.

<sup>c</sup> Following intravitreal injection of WT or mutant virus into the eyes of rats, the number of rats exhibiting virus fluorescence in retinorecipient regions of the brain 48 h.p.i. (either lateral geniculate nucleus or superior colliculus) was determined (19).

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**FIG 2** Retrograde transport of WT or \( /H9004\) particles during initial infection of primary sensory neurons. E8-E10 chick DRG explants were infected with \( \sim 1 \times 10^7 \) PFU of either wild-type or \( /H9004\) PRV, and incoming particles were imaged between 15 and 60 min postinfection based on red fluorescence emissions from the UL25/mCherry capsid tags. More than 190 particles each of WT and mutant virus were tracked. (A) Gaussian (velocity) or decaying exponential (run length) curves were fit to histograms by nonlinear regression; curve-fitting produced \( R^2 \) values of \( >0.96 \). Insets show average velocities (left) and run lengths (right); error bars indicate standard deviations. (B) Representative kymographs of retrograde-moving WT or \( /H9004\) particles.
However, pUS9 in mice, although at reduced efficiency (5). More importantly, neuronal infection models do not support the conclusion that PRV lacking pUS9 does not participate in anterograde axon trafficking and transmission. Whereas an anterograde-deficient strain of PRV (PRV Bartha) does not spread in anterograde circuits within the brain (29), ΔUS9 PRV does so effectively, although not as robustly as wild-type PRV (10). Additionally, in cultures of primary neurons infected with PRV (9) or HSV-1 (7) or in cultures of neuronlike cells infected with HSV-1 (6), pUS9 is an enhancer of anterograde transmission but again is not essential for this process. Therefore, the inability of ΔUS9 PRV to traverse the rat anterograde visual circuitry may be the manifestation of a reduced ability to sort particles into axons, combined with the finite probability of a particle traveling the entire length of these long axons to reach the presynaptic terminals in the brain.

In conclusion, this study provides a kinetic characterization of PRV axonal transport in the absence of the pUS9 anterograde trafficking effector. Deletion of US9 in PRV reduced the number of capsid-containing viral particles sorted into axons but did not directly impair the subsequent step of fast axonal transport for those particles making it past the sorting barrier. The results indicate that the effector function of pUS9 in promoting anterograde axonal trafficking lies in initial sorting of particles to axons and not in fast axonal transport of particles from the proximal axon to terminals.

ACKNOWLEDGMENTS

This work was supported by NIH grants R01 AI056346 to G.A.S. and R01 NS077003 to G.E.P. and P.J.S. G.R.D was supported by the training program in the Cellular and Molecular Basis of Disease from the National Institutes of Health (grant T32 GM08601).

We thank Anne Fischer for excellent technical assistance.

REFERENCES


