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HSV, axonal transport and Alzheimer's disease: *in vitro* and *in vivo* evidence for causal relationships

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Abstract

HSV, a neurotropic virus, travels within neuronal processes by fast axonal transport. During neuronal infection HSV travels retrograde from the sensory nerve terminus to the neuronal cell body, where it replicates or enters latency. During replication HSV travels anterograde from the cell body to the nerve terminus. Postmortem studies find a high frequency of HSV DNA in the trigeminal ganglia as well as the brain. Studies correlating HSV with Alzheimer's disease (AD) have been controversial. Here we review clinical evidence supporting such a link. Furthermore, the author describes experimental data showing physical interactions between nascent HSV particles and host transport machinery implicated in AD. The author concludes that the complexity of this relationship has been insufficiently explored, although the relative ease and nontoxicity of a potential anti-HSV treatment for AD demands further study.

Keywords

A β ; Alzheimer's disease; amyloid precursor protein; APP; cerebrospinal fluid; CNS; encephalitis; fast axonal transport; HSV; senile plaques; trigeminal ganglion; trigeminal nucleus

HSV-1 and HSV-2 initially infect epithelial cells in mucous membranes and secondarily enter local sensory nerve terminals. In the case of HSV-1, initial infection typically occurs in the lip, but can also happen in the mucous membranes of the eye and, rarely, in the nose. HSV-2 typically infects the mucosa of the reproductive tract and also secondarily enters local nerve termini. Although HSV-1 and -2 have tendencies for different anatomic locations, this appears to be a tendency and not a requirement for infectivity. The two types are so similar that PCR primers and antibodies commonly used to detect them clinically do not distinguish between them, and either may cause significant acute disease. Here we will refer to types where possible. If studies did not use type-specific probes, we will refer to the two types as 'HSV'.

HSV-1 is a dsDNA virus with a genome of 150–152 kb containing 89 open reading frames encoding 87 proteins [1]. Alternative splicing and differing transcriptional start–stop sites may produce more variety in proteins synthesized from the viral genome. The infectious

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particle is an enveloped virus with four concentric compartments: the DNA core packaged in an icosahedral proteinaceous capsid surrounded by an amorphous tegument, and enveloped in a glycoprotein-rich lipid bilayer (Figure 1). Both HSV-1 and -2 display neurotropism, with HSV-1 most common in the trigeminal ganglion (TG) containing the cell bodies of the sensory nerves for the lips, eyes, face and meninges, and HSV-2 in the sacral dorsal ganglion containing the cell bodies for the sensory nerves of the perineum and reproductive tract.

Once inside the sensory process, viral particles travel retrograde towards the neuronal cell's nucleus in either the TG or sacral dorsal-root ganglia, where virus may become latent or replicate (Figure 1). Newly synthesized virus travels anterograde within the sensory process to return to the mucosal membrane, thereby causing the recurrent herpes cold sore, or 'fever blister'. Infection of mucosal epithelial cells results in lysis of the cell, termed a 'lytic' infection; whereas infection of neurons most commonly results in latency with periodic reactivations, upon which infective viral particles are produced [2,3]. Viral genomes persist in a nonlinear, episomal, nucleosomal form in infected trigeminal neurons [2].

HSV infection is incurable: the viral genome persists in a latent form as non-integrated episomal dsDNA in the nucleus of the infected neuron for the life of the neuron. Upon reactivation the virus may be shed asymptotically or may cause acute or chronic disease. Thus, infection may or may not produce disease, and reactivations may produce a range of disease severity. In many infected individuals, blisters on the lip recur at odd intervals, often with a chronicity of more than once a year. Each round of blisters arises from reactivation of the viral genome in the latently infected neuron and probably results in the infection of additional neurons in the ganglion as well. Hence many infections are 'chronic' in the sense that reactivations producing acute sores recur over the lifetime of the host. When the eye is involved, repeated herpetic lesions result in blindness [4]. Such chronic, recurrent eruptions represent the third leading cause of infectious blindness worldwide.

Sensory neurons are bipolar in that they project two processes, one to the body surface and the other into the CNS. TG neurons project their second processes into the trigeminal nucleus (TN) located in the brainstem (Figure 1). This second process provides a direct route for HSV into the CNS from infected neurons in the TG, or even during entry from the lip or other facial mucous membranes [5,6]. From the TN in the hindbrain, neuronal projections go to the thalamus and from there to the sensory cortex. Thus, HSV could move throughout the brain and could play pathological roles in various CNS disorders, including Alzheimer's disease (AD). Recent evidence links HSV-1 at a cellular level to AD. Emerging HSV-1 particles interact with amyloid precursor protein (APP), the parent protein that, when hydrolyzed, produces the A β that forms the major component of Alzheimer's plaques, and to induce maldistribution of APP in HSV-infected cells [7].

Major questions remain as to any pathological role of HSV in AD, and a cause-and-effect relationship remains controversial. Critical questions are:

- Is HSV DNA, evidence of latency, found in the brain?
- Does asymptomatic, subclinical replication of HSV occur in the CNS, or is CNS infection only acute as in encephalitis, or quiescent as in latency?
- Does subclinical viral replication in the CNS have consequences? In the case of chronic activity, are the pathological effects transient or persistent?
- How might HSV reach the brain?

- How well does the current evidence of HSV and AD support ‘Koch’s postulates’, a traditional method to determine cause–effect relationships of pathologic microorganisms and disease?

Here epidemiologic reports are reviewed, describing some current experimental *in vitro* evidence, and providing provocative ideas relevant to each of these questions.

HSV is prevalent worldwide

HSV-1 and -2 are endemic in the USA. The prevalence of HSV-1 can be determined by testing serum for antibodies to HSV-1 [8,9], by the detection of HSV-1 DNA in cerebrospinal fluid (CSF) [10], or in TG acquired at autopsy [11]. Such studies have revealed endemic infection rates of 31% in children aged 6–14, rising to 49% in adults aged 14–49, and to a high of 80–90% in the population over 65. In one study of 40 autopsied TGs, HSV-1 sequences were amplified from DNA or RNA extracted from 81% of TGs from demented subjects, and 74% of controls [11]. In another study of younger individuals (average age 38.6 years), HSV-1 or -2 genomes were found by PCR in 53% of TGs [12], and in another study, quantitative PCR detected HSV genomes in 89% of 147 American subjects, with viral load varying from a million copies per sample to almost undetectable [13].

HSV DNA is detected in the brain

HSV-1 and -2 are both found in a high percentage of brains at autopsy in England, Finland and Japan [14,15-18]. In England, 14 out of 14 autopsied brains were positive for HSV-1 tyrosine kinase DNA [17], and 23 out of 36 samples were positive in a subsequent study by the same group [14]. Another English group also found a high prevalence of HSV DNA in brain, with 77 out of 105 subjects (73.3%) found to be positive for HSV using primers based on the HSV-1 gene, *gD* [18]. Both HSV-1 and -2 may be detected by the tyrosine kinase and *gD* primers in these studies. In Finland, nested primers specific for HSV-1 were used, which detected 18 out of 114 subjects as being positive for HSV-1 [15], while in Japan specific primers detected HSV-1 in four out of six brains. Thus, HSV-1 DNA, evidence of latent infection, is found in the CNS of a large proportion of the population at death. HSV-2 DNA has also been found in brain autopsy series, but with lower incidence (13–20% positivity) [19].

Presence of HSV has been correlated with several diseases of the CNS, including AD [20], epilepsy [21] and multiple sclerosis [22]. Viral DNA was detected in the senile plaques of AD brains [20], virus has also been found to be present in epileptic seizure foci [21], and can be cultured from the plaques of multiple sclerosis [22].

Statistically significant correlations between HSV and AD were noted by one group [14,17,23]. By contrast, Hemling *et al.*, studying Finnish subjects (n = 34 with AD and 40 controls), reported lower overall rates of brain HSV infection than in England and failed to find statistically significant correlation between AD and detection of HSV [15]. A study from Japan of three familial and two sporadic post-mortem brains of subjects with AD, and six with non-Alzheimer’s supports the Itzhaki-Jamieson reports, demonstrating HSV in plaques and brains in 100% of both types of AD, but in only one out of five brains lacking A β pathology [16]. A question raised by this Japanese report is whether AD may itself induce HSV reactivation, which could result in a downward spiral.

The high prevalence of HSV latent infections in the brain and the lower incidence of AD raises the major question of what other factors may lead to AD in an HSV-infected host (see Box 1). The most likely variables influencing the effect of infection are the viral load and

the frequency of reactivations with viral production. In addition, the effect of HSV reactivations on the brain and consequent symptomatology may correlate with the anatomical location of infected neurons, their connections and function, and the individual immune response of the patient. Several susceptibility genes for HSV have recently been reported, including TLR3 for both type 1 and 2 [24]; MHC class I allotypes (*B*18*, *C*15* and the group of alleles encoding A19), the high-affinity receptor/ligand pair KIR2DL2/HLA-C1, and the CD16A-158V/F dimorphism for HSV-1 [25]; SNPs in *GRIN2b* (encoding the NMDA receptor isoform) for HSV-2 [26]; polymorphisms in Fas, the cell death receptor, for women in South Africa with HSV-2 [27]; and two SNPs within chromosome 21 open reading frame 91 for HSV-1 [28].

One of the most thoroughly studied host genotypic factors hypothesized to link HSV with AD is allelic variation of the *ApoE* gene. ApoE is a main component of the chylomicron, a serum lipoprotein that binds to specific receptors on liver and other cells, and is essential for the normal catabolism of triglyceride-rich lipoprotein constituents. Several papers report a correlation between HSV-1, *ApoE4* genotype and AD among English subjects (AD, n = 39; non-AD, n = 36), aged 54–96 years [14,23], while other studies, also of a small number of subjects (33 AD brains vs 72 non-AD), found significant amounts of HSV-1 but did not confirm a statistically significant association with ApoE4 [29]. While both of these studies demonstrated the presence of HSV DNA, representing latent virus, in the CNS of a significant number of subjects with or without AD, the correlation with *ApoE4* genotype had a high odds ratio in one study [23] and was not found to be significant by the χ^2 test in another study examining HSV-1 [29]. In female mice deleted for mouse ApoE, HSV-1 viral load was decreased 43-times below that of wildtype, ApoE^{+/+} mice [30]. However, no correlation has been found between ApoE4 and HSV encephalitis in humans [31]. While female mice are more susceptible to HSV, differences in gender infectivity in humans have been difficult to determine due to differences in male and female exposure rates. Difficulties in determining correlations between HSV-1 and ApoE include the high background of endemic viral infection, the over representation of *ApoE4* alleles among the AD subjects, and the necessity of studying outcome at a single time point: the time of death.

The presence of other pathogens in the brain may also contribute to HSV sequelae (see Box 1). Other viral occupants of the human brain in addition to HSV-1 and -2 include other herpes viruses (Varicella zoster [32-34], Human herpes virus 6 [15] and EBV [35,36]; Human herpes virus 7 and 8 [37] and CMV [19]) and other viral species [38], such as the polyoma JC virus (reviewed in [39]). Infection with prokaryotes, including treponemal spirochetes causing syphilis or Lyme disease, may also persist in the brain [40,41].

Detection of HSV DNA in the brain at autopsy by PCR and/or immunohisto chemistry probably leads to under estimation of the incidence of CNS HSV infection owing to sampling error, since only small amounts of this large organ are typically tested: the human brain weighs on average approximately 1500 g, with a volume of approximately 1500 cm³. Both the location in the brain and the amount of tissue sampled will result in false negatives, and may thus lead to a failure to find statistically significant correlations. For example, Hemling *et al.* tested only 15–61 mg of brain from the hippocampus, the frontal cortex, the temporal cortex and the anterior cingulate gyrus [15]. Only one out of 30 AD brains had detectible HSV, while ten out of 30 controls were positive for virus in this Finnish cohort.

Primer design may also lead to false negatives. Although the HSV genome is relatively stable, small nucleotide differences do occur in different strains [42,43] and can arise even in the same individual. Such base-pair changes can affect PCR efficiency or even give false-negative results, especially when nested primers are targeted against one of the more variable HSV-1 genes, *gD*. Many of the early studies cited above used different primers for

the amplification of different viral genes, which could underlie some of the inter-study variability.

Immunocytochemistry has also been used in the past in attempts to find HSV-1 and -2 in the brain. This approach greatly underestimates viral presence and load, as all viral proteins are only synthesized during viral reactivation, which could quite likely not be occurring at the time of death. Thus, viral DNA detection is a vastly superior method for ascertaining viral presence and load, as CNS HSV persists in latent form for the lifetime of the host. US FDA-approved primers now exist for PCR-based tests that are routinely applied clinically for diagnosis based on the highly conserved HSV glycoprotein G gene, which is rarely mutated and provides the most reliable results [44-46].

An autopsy series using clinically approved reliable primers and exhaustive sampling of the TN, the most likely site of viral entry into the CNS from the TG, of a large cohort of subjects would be the minimum required before a linkage between HSV and AD could be ruled out through application of these DNA detection techniques.

Asymptomatic replication of HSV occurs in both the CNS & the peripheral nervous system

Post-mortem PCR-based studies detect latent virus but are not informative about viral activity. What is the evidence that HSV is not just latent in the brain and nervous system, but also replicates?

Detection of viral DNA in the CSF demonstrates that, even during clinically silent infections, virus replicates in the CNS [10]. Although sensory innervation of the meninges emanates from the TG and superior cervical ganglia, these sensory nerve endings occur in the outer meningeal surface with little direct access to the CSF. Random testing of 3200 CSF samples revealed HSV-1 DNA in 26 and HSV-2 DNA in 36 samples from subjects without symptoms of HSV activity. Finding virus in CSF in the absence of clinical signs or symptoms is not surprising, since HSV-1 can also frequently be detected in saliva and tears of asymptomatic subjects [47]. Varicella zoster, another human alphaherpes virus and the cause of chickenpox and shingles, persists like HSV in a latent form and is shed in saliva of asymptomatic hosts [48,49]. Thus, HSV reactivates in the CNS, producing viral particles, over the lifetime of the host.

Subclinical viral replication in the CNS has consequences, & pathological effects persist

Asymptomatic replication of HSV in the CNS may produce pathologic consequences that are not yet recognized. This has become most apparent in children surviving neonatal herpes encephalitis, but may also pertain to adult CNS latency and reactivations.

In neonates, passage through the birth canal may result in perinatal infection, most commonly with HSV-2, leading to herpetic encephalitis. Neonatal mortality from HSV encephalitis has plummeted since the discovery of acyclovir (9-(2-hydroxyethoxymethyl)guanine), which inhibits viral replication through virally encoded tyrosine kinase activity [50-52], the first anti viral agent, for which Gertude Elion received the Nobel Prize in Medicine and Physiology in 1988 [53]. This success arose from earlier discoveries that purine nucleoside analogs inhibited viral replication with low host toxicity due to a unique virally encoded deoxy-thymidine kinase [54].

Treatment of acute neonatal HSV encephalitis began in the 1980s, first with vidarabine (adenine arabinoside) and more recently, acyclovir [55]. While suppressing viral replication during HSV encephalitis in the perinatal period vastly improves survival, such treatment does not eliminate latent virus, which may reactivate over the lifetime of the host. Perinatal infections with HSV-2 may not always produce acute disease, as studies in the elderly born before the advent of antivirals in the 1980s show that a significant number have HSV-2 latent DNA in the brain (13–20%) [56]. New drugs targeting other essential viral functions are urgently needed as acyclovir resistant strains emerge, and some are already in clinical trials [RUEBSAMEN-SCHAEFF H, HILL J. PERS. COMM.].

In our work, we followed an HSV-2-infected immunocompetent child from ages 8 to 12 years [57]. Her history was remarkable for a brief treatment for 2 days with acyclovir during the perinatal period for an episode of clonic seizures and fever. She presented at 8 years of age after a fall, when large asymptomatic granulomatous lesions were found incidentally in her brain on radiologic studies performed to rule out skull fracture. Biopsy of these lesions to rule out tumor revealed actively replicating HSV-2, similar in type to a known maternal infection at the time of pregnancy. Her neurological development had been unremarkable, meeting standard milestones. On presentation at 8 years of age she had no clinical, neurological or overt cognitive deficits. Yet over the 4 years of follow-up, her chronic HSV-2 encephalitis displayed a slow but relentless course, with recurrent viral reactivations and replication, producing lesions of fluctuating mass in the brain, and progressive neurological symptoms, primarily manifesting as attention deficit and learning disabilities. Pharmacologic control of the brain lesions required both antiviral therapy and immune suppression. This child was asymptomatic for >8 years, yet had large brain lesions and replicating HSV-2 in the brain. The lesions persisted despite chronic acyclovir treatment, and progressive deficits have appeared.

While this child's case may be extreme, and perhaps unique, the finding of HSV DNA in randomly sampled CSF from asymptomatic individuals suggests that chronic replication at low levels occurs frequently in many infected individuals. Subclinical symptomatology of such chronic viral replication has not yet been systematically analyzed, although we and others have proposed hypothetical roles in AD, multiple sclerosis and epilepsy as discussed above. Recent reports show that perinatal HSV infections require longer therapies with acyclovir for those surviving children to attain normal developmental milestones [58], demonstrating that subclinical viral replication produces learning and developmental deficits in children, and pathologic effects are persistent and long-lasting, causing progressive deficits. In a transgenic mouse model of familial AD carrying multiple copies of the human *APP* gene with both Indiana and Swedish mutation (*APP*^{Ind/Swe}) under control of the Tet-off promoter, suppression of the mutant APP did not result in loss of plaques that had already formed [59].

Thus, the long-term effects of chronic subclinical HSV reactivations in the CNS of children who survive neonatal HSV encephalitis or acquire virus in the perinatal period and appear asymptomatic, require urgent attention.

How HSV-1 gains entry into the CNS: transport through interaction with cellular transport machinery

HSV-1 and 2 are both known to be neurotropic and to engage in active transport within neuronal processes. Because HSV enters sensory ganglia (TG or sacral dorsal root nerves) that also project a second process to the CNS, this transport is the more likely route of entry into the brain.

The author's laboratory took the novel approach of using HSV-1 as a tool to investigate the cellular mechanisms that govern intra cellular transport [7,60-62]. A major challenge was to distinguish between incoming (infecting) viral particles and outgoing, newly synthesized (infective) particles. Indeed the long-standing controversy concerning the composition of outgoing particles may be a consequence of the experimental challenge of distinguishing incoming from out-going viral particles. While some laboratories show out-going particles lacking detectable membrane components, others report that virus travels outwards within membrane compartments, leading to the 'single' versus 'married' hypotheses for viral egress [63-71]. Data supporting both types of transport have been obtained by electronmicroscopy [72-80]. *In vitro* reconstitution of transport on purified microtubules in the presence of cytosol will allow a dissection of the key molecules involved [81].

To study retrograde transport, the author reconstituted viral particle transport in the squid giant axon by injecting biochemically treated viral particles [60,62]. Green fluorescent protein-labeled viral particles lacking membranes following detergent treatment traveled uniquely retrograde at fast axonal transport rates (2.2 $\mu\text{m}/\text{sec}$) [60]. By contrast, membranated particles traveled anterograde in the giant axon, also at fast transport rates (0.8 $\mu\text{m}/\text{sec}$). Thus, we reasoned that the membranes associated with viral particles contain the receptor that mediates interactions with axonal anterograde transport machinery. Biochemical analysis of this viral-associated membrane revealed the presence of cellular APP, which also colocalized with purified viral particles prior to injection into the axon [62].

We had a breakthrough that allowed us to analyze the composition of outgoing viral particles in the cytoplasm of a productively infected cell. To do this we developed a culture system in which viral infection is so tightly synchronized that >97% of all cytoplasmic viral particles 6 h postinfection are outgoing [3]. Our data demonstrate that >80% of outgoing viral particles inside infected cell cytoplasm are associated with cellular APP, as identified by immunofluorescence, by cotransfection/infection with fluorescent viral proteins and labeled APP, and by immunogold electronmicroscopy (Figure 2). Intracellular nascent viral particles travel more frequently when associated with APP than without, yet may also travel, albeit rarely, independent of detectable APP (Figures 2 & 3) [7]. Productive infection increased the expression of APP and changed its subcellular location. In addition, productive infection with HSV-1 induced the breakdown and reorganization of the microtubule network and reorganization of cellular organelles (Figure 3). This breakdown allows nascent viral particles to travel in either anterograde (kinesin-based) or retrograde (dynein-based) directions on individual microtubules and yet still reach the cell surface (Figure 3).

Subsequent experiments in the author's laboratory also demonstrated that a peptide derived from the cytoplasmic domain of APP was uniquely sufficient to mediate transport of inert plastic nanospheres [82]. Hence we reasoned that virus recruits APP during packaging and uses it to recruit cytoplasmic motors for travel from the perinuclear region to the distant nerve terminal. Such transport would lead to dissemination of virus throughout the arborization of neuronal projections, and thus would be a mechanism by which virus could inhabit multiple sites in the brain, depending upon the projections of each productively infected neuron.

Several virus-encoded proteins are thought to be generally involved in the anterograde transport of various alphaherpesviruses. Glycoprotein E, which forms a heterodimer with glycoprotein I, and is also an Fc receptor for immunoglobulin, travels with viral capsids moving in the anterograde direction [64,83-90]. Virus deleted or mutated for the *gE* gene fails to enter the optic nerve after retinal inoculation [86]. Another viral protein, US9, also a transmembrane envelope protein, travels with viral capsids of either pseudorabies virus or

HSV-1 [64,80,83,91-93]. Interestingly, *US9* is one of the genes altered in the H129 strain of HSV-1, which displays especially active anterograde transport [43]. Recent evidence implicates kinesin 3 (Kif1a) in US9-based HSV transport [KRAMER T *ET AL.* PERS. COMM.].

Intracellular interactions between HSV-1 and APP have been reproduced by others in neurons [94-97]. In cultured human neurons HSV-1 induces A β formation [94,95], probably in autophagic vacuoles. Transport to distant sites of A β formed abnormally in the neuronal cell body is suggested to be a mechanism of seeding the cortex with A β fibrils that act as a nidus for the deposition of normally processed A β in the cortex [98,99]. Recent work by Braak *et al.* demonstrates that similar abnormal processing of A β occurs in the hindbrain and brainstem in early AD – in the dorsal vagus area, locus ceruleus and substantia nigra; regions that project neuronal processes throughout the cortex and hippocampus, and are anatomically close to the TN [100,101]. Interactions between HSV-1 and other proteins involved in neurodegeneration have been reviewed extensively by Carter [102]. Most exciting is the recent finding that acyclovir treatment of HSV-1-infected cultured epithelial cells prevents A β accumulation even at early stages after inoculation [103].

Evidence for HSV reactivation in cognitive decline

Perhaps the most convincing and powerful evidence linking HSV with dementia comes from a prospective study of 512 cognitively normal elderly subjects in a multicenter European cohort who were followed for 14 years for anti-HSV IgG and IgM antibodies and cognitive function. This study did not distinguish between HSV-1 and -2. In the follow-up, 77 new cases of cognitive impairment consistent with AD were found. Subjects with elevated HSV IgM levels had a significantly higher risk of developing HSV (Cox hazard ratio = 2.55; $p = 0.003$), while risk in those with IgG was slightly elevated but not statistically significant (hazard ratio = 1.67; $p = 0.21$). IgM elevations indicate reactivation of HSV, whereas IgG indicates that infection has occurred but the virus is latent. A high proportion of subjects were HSV IgG positive (81%), whereas only a few (7%) had IgM elevations detected over the 14 years of the study. Both AD-like and non-AD dementias occurred at 90% frequency in the HSV-positive subjects with or without the IgM rise. Since IgM would be transitory, it is likely that failure to detect IgM in all subjects who progress to dementia could be due to the timing of blood draw relative to HSV reactivation, or failure of low-grade reactivations to produce a measureable IgM response. In a subsequent study by this group, elevated IgM levels in 1222 subjects (73.9 years, mean age) correlated inversely with plasma A β_{40-42} levels [104]. This was interpreted as indicating that HSV reactivation influences APP processing and CNS deposition, consistent with *in vitro* work described above in which nascent newly synthesized viral particles interact with APP during packaging, transport and egress [7,94].

Recent evidence implicates A β as an antimicrobial peptide that may be an adaptive (innate) immune response to protect the brain from invading organisms [105]. Antimicrobial action was demonstrated for yeast (*Candida albicans*) as well as for bacteria: both cocci (i.e., *Staphylococcus* and *Streptococcus*) and rods (i.e., *Listeria* and *Pseudomonas*, *Escherichia coli*, *Enterococcus faecalis*), with microbicidal activity higher than other well-recognized antimicrobial peptides and even in some cases antibiotics.

A β plaques are also a long-recognized hall-mark of neurosyphilis and may also occur in Lyme disease with involvement of the CNS. In 1913 Noguchi and Moore demonstrated the presence of spirochetes of *Trepanema pallidum* in the brains of cases of general paresis, a dementia with paralysis previously diagnosed as a mental disorder. Subsequently spirochetes were found to induce A β -containing plaques in the brain (reviewed in [40,106]). Other spirochetes also have this consequence, such as *Borrelia burgdorferi*, the causative agent of

Lyme disease [41]. Many other spirochete species exist, often commensal in the mouth and gut of humans, whose presence in brain has not been explored.

That A β may be protective is further supported by the results of clinical trials with drugs intended to clear A β from the brain. Two different approaches, anti-A β antibodies (i.e., bapineuzumab [107]) or inhibition of gamma secretase (i.e., semagacestat [72]) failed to improve cognition, while effectively removing plaque. These clinical drug trials have been halted. Conversely, a recent report from Iceland suggests that mutations that decrease but do not eliminate A β production have a strong protective effect against AD [108]. The A β hypothesis for AD is now under review.

Fulfillment of Koch's postulates & proof of causation

Whether infectious diseases promote or accelerate AD remains controversial, in part because Koch's postulates have not been fulfilled. Robert Koch was a 19th century physician who applied a series of logical steps to identify the causative organisms for anthrax and tuberculosis. These steps were later codified as 'postulates' and have been re-applied to identify the causative organisms in many other diseases, although in some they have had limited power (see **Table 1**).

As methodology to identify infectious disease-causing organisms progresses, revisions of Koch's postulates have been proposed and successfully applied to other organisms, first by AB Hill as presented and discussed at the British Royal Society of Medicine in 1965 [109], and then by Fredericks and Relman in 1996 [110], whose proposal replaces culture with nucleic acid techniques. For HSV, DNA has been isolated from AD brains, although HSV infection is not limited to AD in autopsy series; many non-AD harbor HSV; and many who have HSV do not have AD. The most powerful correlations comes from the European longitudinal prospective study matching serological evidence of HSV reactivations with the onset of cognitive impairment [111].

As for incurable fatal diseases, of which AD is an example, infecting a healthy person with the isolated organism is not an option. For AD there are no validated animal models other than transgenic mice carrying various human genes found in familial AD. These often display aspects of AD without the cognitive components, which are difficult to determine in mice. There are also no animal models that replicate human HSV infections in all their aspects, especially latency and the reactivations that appear to be very important in the hypothesized HSV-AD relationship. Hence, Koch's antiquated postulates cannot be ethically resolved for HSV and AD in the manner originally proposed. However, if the logic behind each postulate is extracted, it may be possible to establish a causal relationship between HSV and AD by developing a new set of 21st century postulates. For example, one such might be, in spontaneously occurring HSV encephalitis resulting in death, does the brain display A β plaques such as those pathognomic for AD?

Conclusion

Given the failure of A β -targeted therapies to improve cognition, alternative etiologies for AD pathology are currently needed. Infectious and inflammatory agents such as HSV may prove integral in our understanding of the pathogenesis and treatment of the disease.

Mounting evidence for a relationship between HSV activity and CNS injury demands further study, with clinical trials to determine if HSV suppression in the elderly protects cognition and prevents cognitive impairment. Urgent attention must be directed to survivors of perinatal HSV encephalitis, with long-term outcome studies and alternative drugs as acyclovir resistance is emerging. Satisfaction of postulates for proof of a causal relationship

will require a creative approach that respects 21st century ethical considerations and exploits 21st century methodologies not available to Koch.

Future perspective

Viral pathogens lurk in the human brain, often in a quiescent state. Even for HSV, perhaps one of the most vigorously studied, little is known about incidence, prevalence, location or viral load in the brain. While studies around the world using PCR approaches detect HSV DNA in brain, a sign of latent infection, few studies have focused on its reactivation from latency. Detection of HSV DNA in CSF and the presence of anti-HSV IgM in the serum of apparently healthy subjects demonstrates that low levels of reactivation without overt symptoms occurs. A β deposits have been also found in apparently healthy, cognitively normal adults. A β plaques have long been known to appear in tertiary syphilis and have recently been reported in Lyme disease. What is now needed is to determine if the microbiome of the brain induces A β as part of an inflammatory innate immune response. The physical interaction within the cell between nascent virus and APP, the parent protein for A β , together with the prevalence of HSV infection in the general population, demand a closer look at causal relationships between HSV and AD, where the plaques are associated with tau-based neurofibrillary tangles. In the absence of animal models and with a potentially fatal outcome, (HSV encephalitis and AD), satisfaction of Koch's postulates in the traditional manner in humans would be unethical. Determination of any causal link between HSV reactivation and Alzheimer's will thus require alternative, yet equally convincing, postulates.

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Box 1. Factors and variables affecting HSV–Alzheimer’s disease correlations

Host factors

- Age of primary infection
- Location of primary infection
- Location of infected neurons in the brain
- Host immune function and local brain inflammatory response
- Host genotype (*ApoE* allele, *TLR3* SNPs, MHC haplotype and so on)
- Environmental exposures (i.e., aspirin use or other drugs)
- Gut microbiome and its metabolites

Viral factors

- Viral load and distribution
- Frequency and timing of reactivations
- Presence of other viruses (i.e., JC virus, CMV and human herpes virus 6–8)
- Genotype of the HSV type and strains
- Role of HSV-2 compared with HSV-1
- Superinfections with more than one HSV type or strain

Confounding experimental variables

- Small number of subjects in study
- Prevalence of endemic infections (high background)
- Sampling error – small amount of total brain tested
- Reliability of PCR primers for genomically unstable viral genes
- Limited availability of normal brains for age-matched control

Executive summary

HSV is prevalent worldwide & HSV DNA is found in the brain

- Endemic infection rates for HSV in the US are immense: 31% of children aged 6–14 years, rising to 49% in adults aged 14–49 years, to a high of 89% of the population over 65 years.
- HSV-1 DNA is commonly detected by PCR in post-mortem examination of brains in England (73.3%), Finland (16%), Japan (44%) and the USA (81%).

Asymptomatic replication of HSV in both brain & the peripheral nervous system

- Chronic HSV replication occurs in the peripheral and CNS, with HSV-1 DNA found in saliva and HSV-1 and -2 found in the cerebral spinal fluid of asymptomatic subjects.
- Cognitive impairment correlates with serologic evidence of HSV reactivation in cognitively normal elderly subjects.

HSV-1 interacts with amyloid precursor protein, the substrate of the major protein, A β , *in Alzheimer's plaques*

- Nascent HSV-1 particles travel with amyloid precursor protein inside cells, resulting in abnormal amyloid precursor protein distribution.
- HSV-1 infection of neurons in culture leads to increased A β production.

Conclusion

- Evidence is accumulating that HSV plays a role in Alzheimer's disease. To satisfy a causal link between HSV and Alzheimer's disease, an approach that takes 21st century ethical considerations and methodologies into account must be developed.

Future perspective

- Monitoring cognition in HSV-positive subjects with mild cognitive impairment during long-term treatment with acyclovir, US FDA-approved for herpes virus therapy, will provide support for a causal relationship while also developing a novel treatment modality.
- If HSV replication in the brain *in vivo* results in AD-like pathology, than a causal link satisfying Koch's postulates will be obtained.

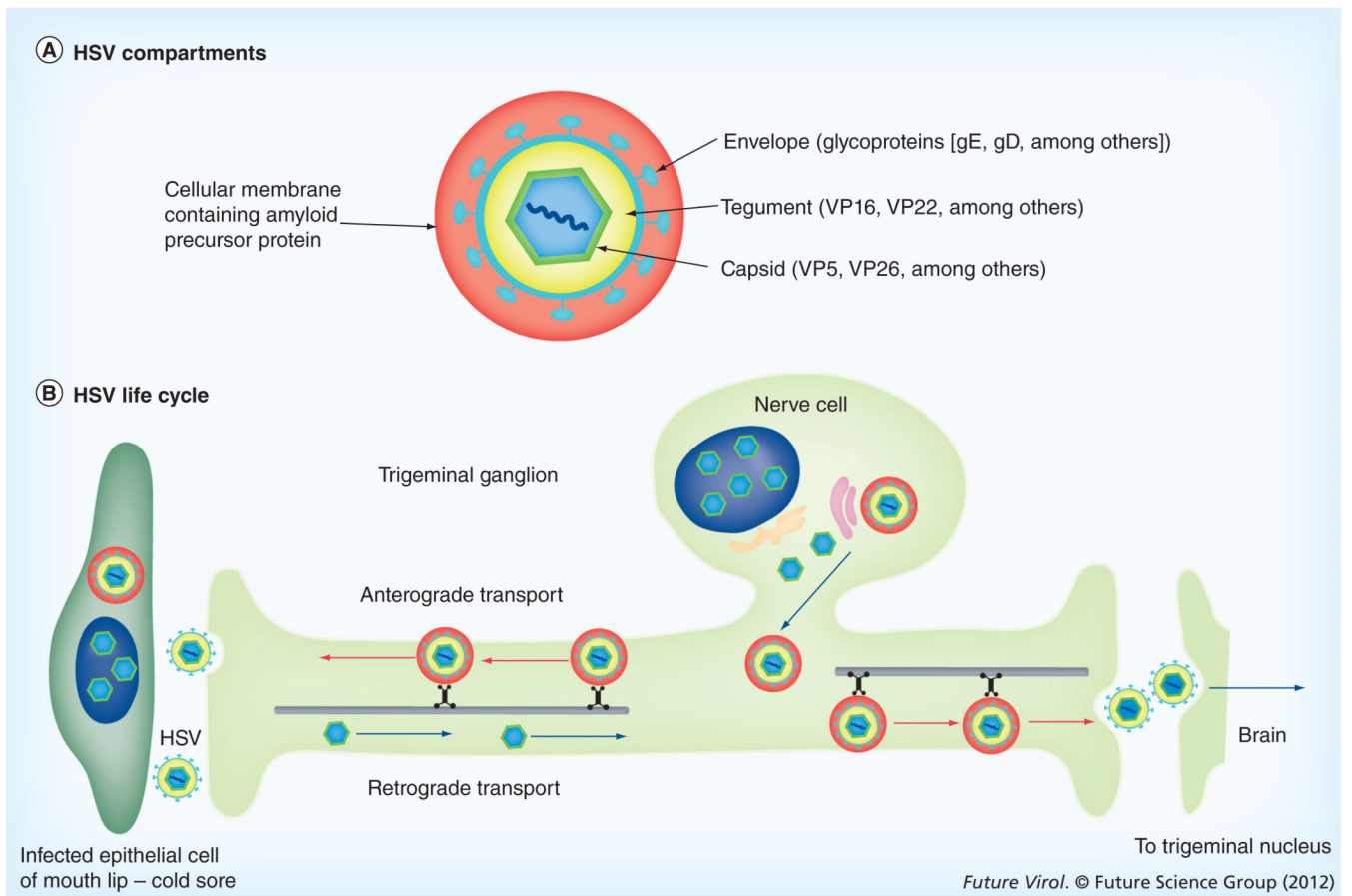


Figure 1. HSV structure and life cycle

(A) HSV compartments: this schematic depicts the four concentric compartments of the infective particle encased in a fifth compartment provided by a cellular membrane system found encircling cytoplasmic viral particles during productive infections and containing amyloid precursor protein. The capsid is approximately 135 nm in diameter and the whole assemblage may be as large as 350–400 nm. **(B) HSV life cycle:** here we outline the process the virus follows to infect the trigeminal ganglion and progress into the trigeminal nucleus in the brain. Note that the trigeminal ganglion cells project two processes: one to the lip, the other to the brain. Virus traveling in a single neuron can move within these processes to go either to the lip or into the brain. The route from the lip to the brain can be >10 cm in length.

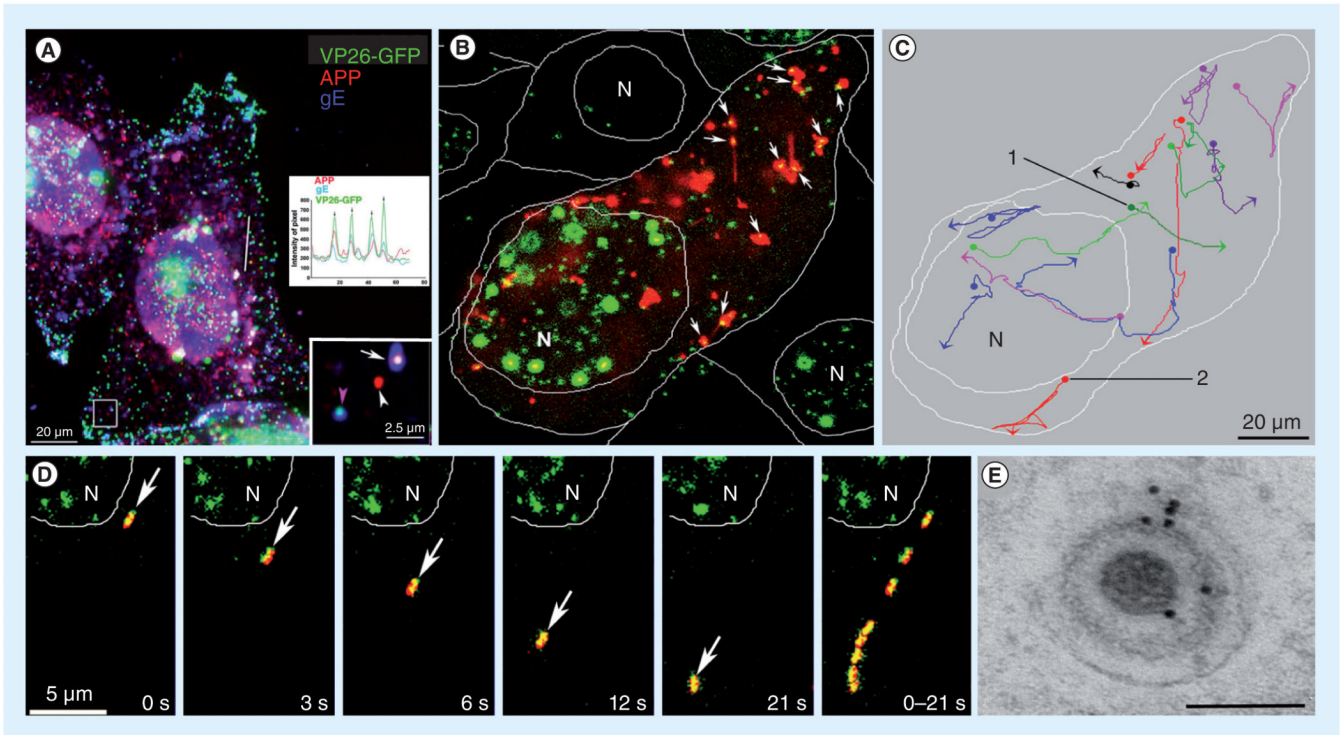


Figure 2. HSV-1 interacts with cellular amyloid precursor protein during egress

(A) VP26-GFP HSV-1 (green) particles colocalize with APP (red) during egress by immunofluorescence after synchronous infection with VP26-GFP HSV-1 (green) and fixation at 7 h postinfection. Examples of two infected cells also stained by immunofluorescence for gE (blue). Lower right inset shows a higher magnification of the boxed region. In this inset, one VP26-GFP-labeled particle displays all three labels (gE, APP and VP26), another has only APP (white arrowhead) or only gE (pink arrowhead). Such single labels demonstrate that colocalization is not due to bleed-through from other fluorescent channels. The inset graph shows intensity profiles along the line drawn across the right-hand cell. Superimposed peaks in the graph for the different colors are indicated by vertical arrows. (B) The first frame of a video sequence showing the initial positions of HSV capsids (VP26-GFP, green) and cellular APP-labeled with red fluorescent protein (APP-monomeric red fluorescent protein, red) in an infected cell at 7–9 h postinfection from a 3-s time-lapse 900-s (15 min) video. Double-labeled particles appear yellow (arrows). Many capsids (64%) colocalize with APP compartments in this frame. Capsids travel with APP vesicles, and sometimes join and separate from them. Positions of the N and of the cell boundaries are delineated by white lines. (C) The tracks of selected VP26-GFP HSV particles that move with the APP-monomeric red fluorescent protein label. Each trace has been assigned a different pseudocolor. Beginnings and ends of movements are indicated by dots and arrowheads, respectively. (D) An example of a double-labeled HSV-APP particle that moves away from the nucleus (arrow). In the last panel eight frames are superimposed to demonstrate the particle's trajectory. (E) Immunogold-labeled electron micrograph of an HSV-1 particle inside an infected cell cytoplasm probed with anti-C-APP followed by protein-A labeled with 10 nm gold. Gold particles decorate both cellular and viral membranes surrounding capsids.

Scale bar = 100 nm.

APP: Amyloid precursor protein; GFP: Green fluorescent protein; N: Nuclei.

Modified with permission from [7]. Supporting videos for these images can be found on the *PLoS One* journal website linked to the online publication.

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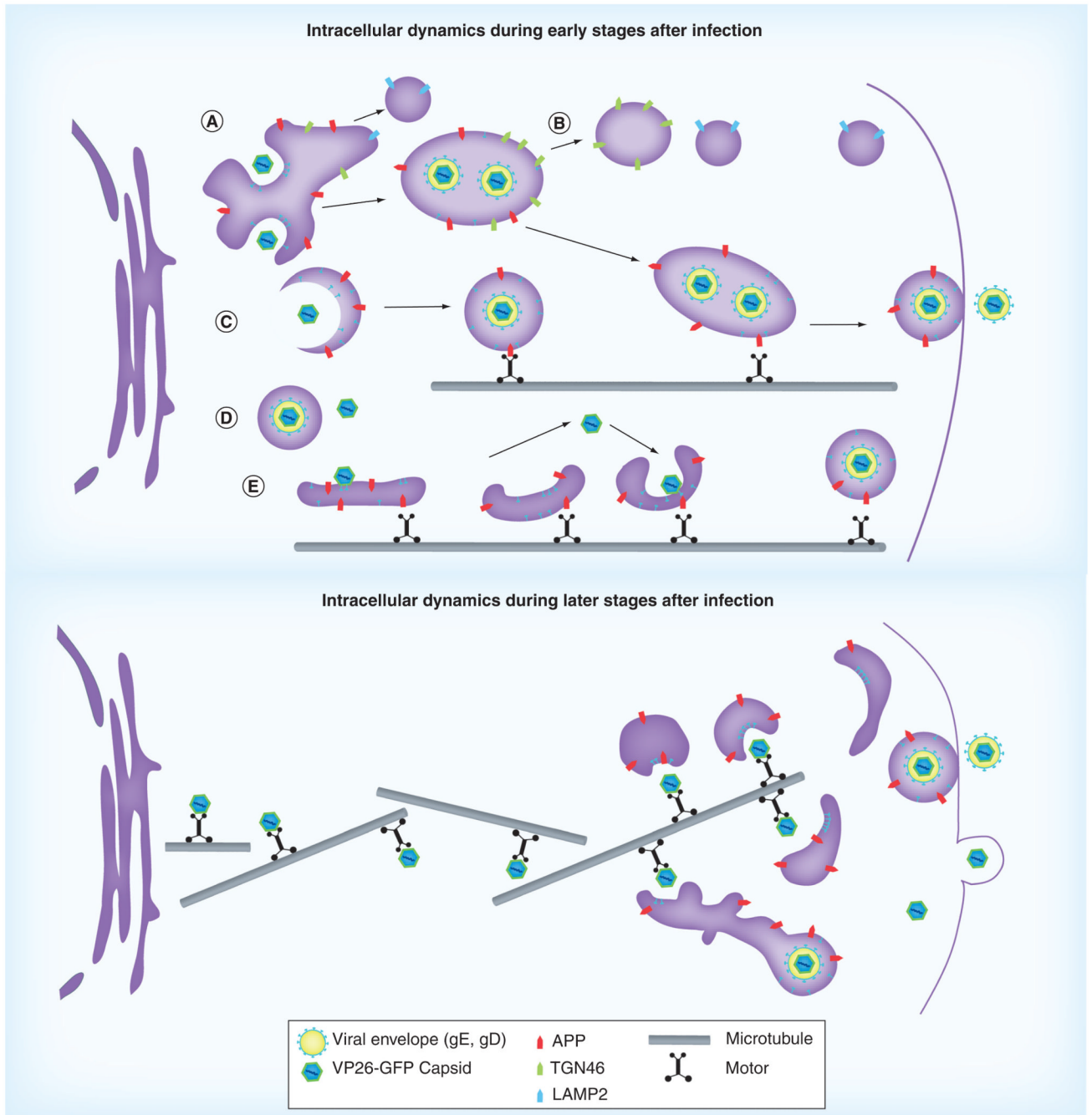


Figure 3. Schematics showing intracellular interactions between cellular amyloid precursor protein and HSV particles at an early stage (top) and a later stage (bottom) of productive infection in epithelial cells

Intracellular dynamics at early stages during viral replication (top panel): (A) in the perinuclear region, viral particles move around and within the large membranous compartments near the nucleus, and also frequently colocalize with the viral envelope glycoproteins, gE and gD, and with cellular transmembrane organelle proteins in this area of the cell, such as: LAMP2, a lysosomal membrane protein; TGN46, associated with the trans-Golgi network; and APP, whose physiologic function is unknown. In the mid-cytoplasmic area, the LAMP2 compartments are separate from this apparent Golgi network, and rarely

colocalize with any viral or Golgi components at the periphery. **(B)** In the outer cytoplasm neither LAMP2 nor TGN46-positive organelles colocalize with viral particles, while APP particles remain associated both with viral capsids and with viral envelope glycoproteins, gE and gD, probably on their way to the cell surface. **(C)** Capsids entering smaller post-Golgi compartments that stain uniquely for APP undergo fast transport towards the cell surface. APP: Amyloid precursor protein; GFP: Green fluorescent protein.

Adapted from [7], with permission as modified from [61].

(D) After leaving the nucleus, viral particles are also found without detectible APP or the other cellular membrane proteins studied here. These particles could be inside some other unlabeled membrane system, or be free in the cytoplasm. These solo particles will be transported only rarely. **(E)** Capsids, possibly with tegument, may ride on the cytoplasmic surface of APP-labeled membrane systems, attach and detach from these membranes, or bud into them. Any particular capsid may employ all of these mechanisms during transit in the cytoplasm. In each case, we hypothesize that microtubule motors, such as one or more of the kinesin family, are recruited to the capsid-containing membrane-bound particles, possibly via APP or another cellular motor receptor. Dimensions are not to scale. Intracellular dynamics at later stages (bottom panel): as the microtubule network breaks down and reorganizes during productive infection, capsids must travel further to reach the cellular membrane synthesis and packaging centers in the Golgi apparatus, which drifts out during productive infections. In this case capsids travel directly on microtubules, possibly recruiting retrograde machinery as for infective entry, and yet reach the cell surface due to the disorganization of the microtubule network. Upon arrival at the cortex, where the Golgi has drifted, viral envelopment takes place. Dimensions are not to scale.

APP: Amyloid precursor protein; GFP: Green fluorescent protein.

Adapted from [7], with permission as modified from [61].

Table 1

Koch's postulates revisited.

Postulates	Caveats
The agent must be found in abundance in all individuals suffering from the disease but not in healthy individuals (Koch later abandoned the universal requirement for this postulate)	Healthy individuals may harbor cholera and typhoid and not be affected but become carriers; polio only causes paralysis in a small subset of cases; asymptomatic or subclinical infections occur for many viruses, such as HIV and hepatitis C as well as HSV For HSV, many variables influence disease (see Box 1)
The microorganism must be isolated from a diseased individual and grown in culture	Prions have never been grown in culture, and many other difficult- or impossible-to-culture infectious disease-causing organisms exist that are now identified by nucleic acid assays and confirmed by consensus as disease-causing For HSV, this postulate has been fulfilled for mucosal blisters and acute encephalitis. DNA has been found in the CNS in many studies
The cultured microorganism should cause disease when introduced into a healthy organism	Infection may depend on host factors as well, as in the resistance to malaria of individuals carrying the sickle-cell gene. Koch himself used the word 'should' instead of 'must' to indicate that this is not universal. This postulate cannot be ethically satisfied for fatal diseases with no animal model, such as sporadic Alzheimer's disease For HSV, this postulate has been fulfilled for cold sores, encephalitis and teratology
The microorganism must be re-isolated from the inoculated diseased individual	Re-isolation requires that the microorganism can be inoculated For HSV, this postulate has been fulfilled for cold sores, encephalitis and teratology