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6   **Neuronal hyperexcitability is a DLK-dependent trigger of HSV-1 reactivation that**

7   **can be induced by IL-1**

9   Sean Cuddy[1,2#], Austin R. Schinlever[1#], Sara Dochnal[1], Jon Suzich[1], Parijat Kundu[1],

10   Taylor K. Downs[3], Mina Farah[1], Bimal Desai[3], Chris Boutell[4] and Anna R. Cliffe[1]*

13   1. Department of Microbiology, Immunology and Cancer Biology, University of

14   Virginia, Charlottesville, VA, 22908.

15   2. Neuroscience Graduate Program, University of Virginia, Charlottesville, VA,

16   22908

17   3. Department of Pharmacology, University of Virginia, Charlottesville, VA, 22908.

18   4. MRC-University of Glasgow Centre for Virus Research (CVR), Garscube

19   Campus, Glasgow, Scotland, United Kingdom

21   # Denotes equal contribution

23   * Correspondence to Anna R. Cliffe, cliffe@virginia.edu

Abstract

Herpes Simplex Virus (HSV) establishes a latent infection in neurons and periodically reactivates to cause recurrent disease. The stimuli that act on neurons to trigger HSV reactivation have not been fully elucidated. Here we demonstrate that HSV reactivation is triggered by stimuli that induce neuronal hyperexcitability. Neuronal stimulation-induced reactivation was dependent on voltage-gated ion and hyperpolarization-activated cyclic nucleotide-gated (HCN) channels, demonstrating that neuronal activity is required for HSV reactivation. Hyperexcitability-induced reactivation was dependent on the neuronal specific pathway of DLK/JNK activation and progressed via an initial wave of viral gene expression that was independent of histone demethylase activity and linked to increased levels of histone phosphorylation. IL-1β induces neuronal hyperexcitability and is released under conditions of psychological stress and fever; both known triggers of clinical HSV reactivation. IL-1β treatment of sympathetic neurons induced histone phosphorylation, and importantly HSV reactivation, which was dependent on both DLK and neuronal excitability. Thus, HSV co-opts an innate immune pathway resulting from IL-1 stimulation of sympathetic neurons to induce reactivation from latency.

 Introduction

Herpes simplex virus-1 (HSV-1) is a ubiquitous human pathogen that is present

in approximately 40-90% of the population worldwide[1]. HSV-1 persists for life in the

form of a latent infection in neurons, with intermittent episodes of reactivation.

Reactivation from a latent infection and subsequent replication of the virus can cause

substantial disease including oral and genital ulcers, herpes keratitis and encephalitis.

In addition, multiple studies have linked persistent HSV-1 infection to the progression of

Alzheimer's disease[2]. Stimuli in humans that are linked with clinical HSV-1 reactivation

include exposure to UV light, psychological stress, fever and changes in hormone

levels[3]. How these triggers result in reactivation of latent HSV-1 infection is not fully

understood.

During a latent infection of neurons, there is evidence that the viral genome is

assembled into a nucleosomal structure by associating with cellular histone proteins[4].

The viral lytic promoters have modifications that are characteristic of silent

heterochromatin (histone H3 di- and tri-methyl lysine 9; H3K9me2/3, and H3K27me3)[5-8],

which is thought to maintain long-term silencing of the viral lytic transcripts. Hence, for

reactivation to occur, viral lytic gene expression is induced from promoters that are

assembled into heterochromatin and in the absence of viral proteins, such as VP16,

which are important for lytic gene expression upon *de novo* infection. Reactivation is

therefore dependent on the host proteins and the activation of cellular signaling

pathways[3]. However, the full nature of the stimuli that can act on neurons to trigger

reactivation and the mechanisms by which expression of the lytic genes occurs have

not been elucidated.

One of the best characterized stimuli of HSV reactivation at the cellular level is

nerve-growth factor (NGF) deprivation resulting from loss of PI3K/AKT activity[9-11].

Previously, we found that activation of activation of the c-Jun N-terminal kinase (JNK)

cell stress response via activation of dual leucine zipper kinase (DLK) was required for

reactivation in response to loss of NGF signaling. In addition, recent work has identified

a role for JNK in HSV reactivation following perturbation of the DNA damage/repair

pathways, which also trigger reactivation via inhibition of AKT activity[12]. DLK is a master

regulator of the neuronal stress response, and its activation can result in cell death,

axon pruning, axon regeneration or axon degeneration depending on the nature of

activating trigger[13,14]. Therefore, it appears that HSV has coopted this neuronal stress

pathway of JNK activation by DLK to induce reactivation. One key mechanism by which

JNK functions to promote lytic gene expression is via a histone phosphorylation on S10

of histone H3[15]. JNK-dependent histone phosphorylation occurs on histone H3 that

maintains K9 methylation and is therefore known as a histone methyl/phospho switch,

which permits transcription without the requirement for recruitment of histone

demethylases[16,17]. This initial wave of viral lytic gene expression is known as Phase I,

and also occurs independently of the lytic transactivator VP16. In addition, late gene

expression in Phase I occurs independent of viral genome replication[18,19]. A sub-

population of neurons then progress to full reactivation (also known as Phase II), which

occurs 48-72h post-stimulus and requires both VP16 and histone demethylase activity

[15,20-23]. However, not all models of reactivation appear to go through this bi-phasic

progression to reactivation as axotomy results in more rapid viral gene expression and

dependence on histone demethylase activity for immediate viral gene expression.

The aim of this study was to determine if we could identify novel triggers of HSV

reactivation and determine if they involved a bi-phasic mode of reactivation. We decided

to focus on stimuli that cause heightened neuronal activity because hyperstimulation of

cortical neurons following forskolin treatment or potassium chloride mediated

depolarization has previously been found to result in a global histone methyl/phospho

switch[24]. Whether this same occurs in different types of neurons, including sympathetic

neurons, is not known. Although forskolin has previously been found to induce HSV

reactivation,[25-28], whether this is a result of inducing hyperexcitability or as a

consequence of activation of alterative cAMP-responsive proteins including PKA and

CREB is not known. Therefore, we investigated if forskolin-mediated reactivation was a

result of increased neuronal activity. Hyperexcitability of neurons is correlated with

changes in cellular gene expression, increased DNA damage[29,30] and epigenetic

changes including H3 phosphorylation[24]. However, DLK-mediated activation of JNK has

not been linked to changes in cellular gene expression in response to hyperexcitability.

Therefore, we were also interested in determining whether DLK and JNK are required

for induction of HSV gene expression in response to hyperexcitability.

IL-1β is released under conditions of psychological stress and fever[31-34]; both

known triggers of clinical HSV reactivation[35-37]. IL-1β has previously been found to

induce heightened neuronal activity[38-40]. However, an intriguing feature of IL-1β

signaling is its ability to have differential effects on different cell types. For example, IL-

1β is involved extrinsic immune response to infection via activation of neutrophils and

lymphocytes. In addition, it can act on non-immune cells including fibroblasts to initiate

an antiviral response[41,42], as has previously been described for lytic infection with HSV-

1[41]. Given these differential downstream responses to IL-1β signaling, we were

particularly interested in the effects of IL-1β treatment of latently infected neurons.

Interestingly, we found that IL-1β was capable of inducing reactivation of HSV from

mature sympathetic neurons. Inhibition of voltage-gated sodium and hyperpolarization

activated cyclic nucleotide gated (HCN) channels prevented reactivation mediated by

both forskolin and IL-1β. Activity of the cell stress protein DLK was also essential for IL-

1β-mediated reactivation. We therefore identify IL-1β as a novel trigger from HSV

reactivation that acts via neuronal hyperexcitability and highlight the central role of JNK

activation by DLK in HSV reactivation.

<u>Results</u>

<u>Adenylate Cyclase Activation Triggers DLK/JNK-Dependent Reactivation of HSV from</u>

<u>Latent Infection in Sympathetic Neurons</u>

130   Both forskolin and cAMP mimetics are well known to induce neuronal

hyperexcitation and have previously also been found to trigger HSV reactivation[25-28].

Using a model of HSV latency in mouse sympathetic neurons isolated from the super-

cervical ganglia (SCG)[15] we investigated whether forskolin treatment induced

reactivation in this system and the potential mechanism resulting in the initial induction

of viral lytic gene expression. Sympathetic SCG neurons were infected with a Us11-

GFP tagged HSV-1[43] at a multiplicity of infection (MOI) of 7.5 PFU/cell in the presence

of acyclovir (ACV). After 6 days the ACV was washed out and the neuronal cultures

monitored to ensure that no GFP-positive neurons were present. Two days later,

reactivation was triggered by addition of forskolin (Figure 1A). As represented in figure

1B, forskolin can act either extra-cellularly on ion channels or intracellularly to activate

adenylate cyclase[44-46]. Dideoxy-forskolin is a cell impermeable forskolin analog that can

act directly on voltage gated ion channels but does not activate cAMP[44,47]. We found

addition of forskolin but not dideoxy-forskolin triggered HSV reactivation (Figure 1C). In

addition, treatment of latently infected primary neurons with a cAMP mimetic (8-bromo-

cAMP) was sufficient to trigger reactivation (Figure 1D) and inhibition of adenylate

cyclase activity using SQ22, 536[48] significantly diminished HSV reactivation (Figure 1E).

Therefore, activation of adenylate cyclase activation and subsequent increased

intracellular levels of cAMP are required for forskolin-mediated reactivation.

DLK/JNK Activity is Required for the Early Phase of Viral Gene Expression in Response

to Forskolin Treatment

We previously found that DLK-mediated JNK activation was essential for Phase I

reactivation following interruption of nerve growth factor signaling. To determine

whether DLK and JNK activation were crucial for reactivation in response to

hyperexcitability, neurons were reactivated with forskolin in the presence of the JNK-

inhibitor SP600125 (Fig. 2A) or the DLK inhibitor GNE-3511[49] (Fig. 2B), which

prevented reactivation based on the number of GFP-positive neurons at 3-days post-

Figure 1. HSV-1 Reactivation Induced by Adenylate Cyclase Activation is DLK/JNK-Dependent. (A) Schematic of the primary superior sympathetic ganglia-derived model of HSV latency. Reactivation was quantified based on Us11-GFP positive neurons in presence of WAY-150168, which prevents cell-to-cell spread. (B) Schematic of the cellular pathways activated by forskolin treatment. Forskolin can act both intracellularly to activate adenylate cyclase increasing the levels of cAMP or extracellularly on ion channels. (C) HSV reactivation was induced by treatment with forskolin (60μM) but not the cell-impermeable dideoxy-forskolin (60μM). (D) Reactivation could be triggered by the cAMP mimetic 8-Bromo-cAMP (125μM). (E) Forskolin-induced reactivation was blocked by the adenylate cyclase inhibitor SQ22,536 (50μM). Each point represents a single replicate. Statistical comparisons were made using Welch's t-test.

reactivation Inhibition of either JNK or DLK blocked HSV reactivation in response to

forskolin. These data therefore indicate hyperexcitability-induced reactivation is

dependent on the neuronal stress pathway mediated by DLK activation of JNK.

Because we previously found that JNK-activation results in a unique wave of viral

gene expression in response to inhibition of nerve-growth factor signaling, we especially

intrigued to determine whether hyperexcitability triggers a similar wave of JNK

dependent viral gene expression. The previously described bi-phasic progression to

viral reactivation is characterized by viral DNA replication and production of infectious

virus, occurring around 48-72 post-stimulus[18], but an earlier wave of lytic gene

expression occurring around 20h post-stimulus. To determine whether forskolin-

mediated reactivation results in a similar kinetics of reactivation, we investigated the

timing of Us11-GFP synthesis, viral DNA replication, production of infectious virus and

lytic gene induction following forskolin treatment. In response to forskolin treatment,

GFP synthesis in neurons started to appear around 48h post-reactivation, with more

robust reactivation observed at 72h (Figure 2C). In contrast to forskolin-mediated

reactivation, the number of GFP-positive neurons following superinfection with a

replication competent wild-type virus resulted in a rapid induction of GFP-positive

neurons by 24h post-superinfection (Figure 2C). Therefore, forskolin triggered

reactivation results in slower synthesis of Us11-GFP than superinfection. In addition,

these data highlight the ability of forskolin to trigger reactivation from only a

subpopulation of latently infected neurons (approximately 1 in every 3.4 neurons

compared to superinfection).

Figure 2

Figure 2. Reactivation Triggered by Forskolin Involves a DLK/JNK Dependent Phase I of Viral Gene Expression.

(A) Reactivation was induced by forskolin in the presence of JNK inhibitor SP600125 (20μM). (B) Reactivation was induced by forskolin in the presence of the DLK inhibitor GNE-3511 (4μM). (C) Reactivation was induced by forskolin or superinfection with a wild-type (F strain) HSV-1 (MOI of 10 PFU/cell) and qualified based on Us11-GFP positive neurons (n=3). (D-F) RT-qPCR for viral mRNA transcripts following forskolin treatment of latently infected SCGs. (G-I) RT-qPCR for viral lytic transcripts at 20h post-forskolin treatment and then in presence of the JNK inhibitor SP600125 (20μM) and the DLK inhibitor GNE-3511 (4μM). In D-L each experimental replication is represented. Statistical comparisons were made using two-tailed paired student's t-test.

Figure S1

<u>Figure S1(A) Reactivation triggered by forskolin triggers a wave of lytic gene expression that precedes DNA replication and infectious virus production (supplement to Figure 2)</u>
(A) Titers of infection virus detected from reactivating neurons induced with forskolin (n=4).
(B) Quantification of the relative viral genome copy number following forskolin-mediated reactivation based on gC copy number normalized to cellular GAPDH and expressed relative to the 0h time-point (n=7). (C-E) RT-qPCR for viral mRNA transcripts following forskolin treatment of latently infected SCGs. In C-E each biological replicate is represented. Statistical comparisons were made using two-tailed paired student's t-test (C-E).

The production of infectious virus also mirrored the data for the detection of

Us11-GFP positive neurons, with a robust increase in viral titers between 24 and 60h

post-stimulus (Figure S1A). An increase in viral genome copy was also not detected

until 48h post-stimulus, which continued between 48h and 72h (Figure S1B). The

quantification of viral genome copy number was also carried out in presence of WAY-

150138[50], which prevents packaging of the viral genome[51], therefore indicating that

DNA replication occurs in reactivating neurons and not as a consequence of cell-to-cell

spread.

Given the observed 48h delay in viral DNA replication and production of

infectious virus, we were interested to determine if there was a Phase I wave of lytic

gene expression that occurred prior to viral DNA replication. We therefore carried out

RT-qPCR to detect representative immediate-early (*ICP27* and *ICP4*), early (*ICP8* and

UL30) and late (*UL48* and *gC*) transcripts between 5- and 20-hours post addition of

forskolin (Figures 2D-F and S1C-E). For all six transcripts, a significant up-regulation of

mRNA occurred at 20h post-treatment, including the true late gene *gC*, whose

expression would usually only be stimulated following viral genome replication in the

context of *de novo* lytic replication. Therefore, this indicates that lytic gene expression is

induced prior to viral DNA replication and that neuronal hyperexcitability does trigger a

Phase I wave of lytic gene expression. Notably, we did detect small but reproducible

induction of *ICP27* mRNA at 5h post-stimulus, followed by a second induction at 20h

(Figure 2D), indicating that there is likely differential regulation of some viral lytic

transcripts during Phase I reactivation induced by hyperexcitability that is distinct from

both NGF-deprivation and *de novo* lytic infection.

To determine whether JNK and DLK were required Phase I gene expression in

response to hyperexcitability, we investigated viral mRNA levels following forskolin-

mediated reactivation in the presence of the JNK inhibitor SP600125. We found a

significant reduction in *ICP27* (2.2-fold), *UL30* (3.3-fold) and *gC* (5.5-fold) mRNA levels

at 20h post-stimulus in the presence of SP600125 (Figure 2J-L). For all genes tested,

there was no significant increase in mRNAs in the JNK-inhibitor treated neurons

compared to mock. We observed comparable results following treatment with the DLK

inhibitor GNE-3511, with a 2.3-, 3-, 8.8-fold decrease in *ICP27*, *UL30* and *gC* mRNAs

respectively compared to forskolin treatment alone, and no significant increase in

mRNA levels compared to the reactivated samples (Figure 2J-L).

It is possible that in addition to JNK, other signal transduction proteins are

important in forskolin-mediated reactivation. Previous data has found that DLK can be

activated by PKA, which is well known to be activated by cAMP. However, using well

characterized inhibitors of PKA, along with the PKA-activated transcription factors

CREB, in addition to two other cAMP responsive proteins Rapgef2 and EPAC, we did

not find that these cAMP activated proteins were required for Phase I reactivation

(Figure S2). Inhibition of PKA or CREB did reduce Phase II reactivation (Figure S2A

and C) but had no effect on Phase I (Figure S2B and D). Inhibition of Rapgef2 or PEAC

had no effect on HSV reactivation (Figure SE and F). Taken together, these data

Figure S2

Figure S2. Effect of PKA, CREB, Rapgef2 and EPAC Inhibition on HSV-1 Reactivation. (A) Latently infected cultures were reactivated with forskolin (60 μM) in the presence of the PKA inhibitor KT 5720 (3 μM) and the number of Us11-GFP positive neurons quantified at 3 days post-reactivation. (B) RT-qPCR for the viral lytic transcript ICP27 at 20h post-forskolin treatment and in the presence of KT 5720. (C) Latently infected cultures were reactivated with forskolin in the presence of the CREB inhibitor 666-15 (2 μM). (D) RT-qPCR for ICP27 at 20h post-forskolin treatment and in the presence of 666-15. (E) Latently infected cultures were reactivated with forskolin (60μM) in the presence of the EPAC inhibitor ESI09 (10 μM). (F) Latently infected cultures were reactivated with 8-Bromo-cAMP (125μM) in the presence of the Rapgef2 inhibitor SQ22,536 (50 μM). Individual experimental replicates are represented. Statistical comparisons were made using Welch's t-test (A) or two-tailed unpaired t-test (B).

232   indicate that it is hyperexcitability induced by forskolin that induces a Phase I wave of

233   gene expression via activation of DLK and JNK.

235   <u>Forskolin Triggers a Phase-I Wave of Viral Gene Expression that is Independent of</u>

236   <u>Histone Demethylase Activity.</u>

237     Hyperexcitability results in the propensity of neurons to fire repeated action

potentials, and is associated with specific changes in histone posttranslational

modifications. The first is physiological DNA damage[29,30], measured by the intensity of

γH2AX staining in neuronal nuclei. Forskolin treatment was associated with an increase

in the levels of γH2AX at 5h post-treatment, which resolved by 15h post-treatment

(Figure S2A and C), and is therefore indicative of physiological DNA damage and

repair, which occurs upon neuronal hyperexcitability. A second reason for probing the

DNA damage/repair pathway in response to forskolin treatment is that previously

reactivation of HSV from latency has been associated with perturbation of the DNA

damage/repair response[12]. Both inhibition of repair and exogenous DNA damage

resulted in loss of AKT phosphorylation by PHLPP1, which was required for HSV

reactivation. Although we did observe increased levels of γH2AX following forskolin

treatment, this was not accompanied by a loss of pAKT measured at 15h post-treatment

(Figure S2D). This indicates that HSV reactivation in response to forskolin treatment

does not involve dephosphorylation of AKT. Therefore, hyperexcitability triggers

reactivation via an alternative mechanism that does not feed into the AKT.

Figure 3

Figure 3. The Initial Wave of Viral Lytic Gene Expression During Forskolin-mediated Reactivation is Independent on Histone Demethylase Activity. (A) Quantification of the percentage of genome foci stained using click-chemistry that co-localize with H3K9me3/S10p. At least 15 fields of view were blindly scored from two independent experiments. Whiskers represent the 2.5-97.5 percentile range. (B) Representative images of click-chemistry based staining of HSV-EdC genomes and H3K9me3/S10p staining at 5h post-forskolin treatment. (C and D). Effect of the LSD1 inhibitors OG-L002 and S 2101 on forskolin-mediated Phase I of reactivation determined by RT-qPCR for ICP27 (C) and gC (D) viral lytic transcripts at 20h post-forskolin treatment and in the presence of 15μM OG-L002 and 20μM S 2102. (E) Effect of the JMJD3 and UTX inhibitor GSK-J4 (2μM) on forskolin-mediated Phase I measured by RT-qPCR for viral lytic transcripts ICP27 (E) and gC (F) at 20h post-forskolin treatment and in the presence of GSK-J4. For C-F each experimental replicate is represented. Statistical comparisons were made using two-tailed unpaired t-test (A) or Welch's t-test (C-F).

Figure S3

<u>Figure S3. Forskolin Treatment Induces Hyperexcitability-associated Histone Post-translational Modification in Sympathetic Neurons</u>. SCG neurons were treated with forskolin and immunofluorescence staining was carried out for H3K9me3/S10p, the DNA damage marker γH2AX and the neuronal marker beta III-tubulin. (B) Quantification of neuronal nuclear staining intensity for H3K9me3 (>150 cells/condition). (C) Quantification of neuronal nuclear staining for γH2AX. In B and C, whiskers represent the 2.5-97.5 percentile range. (D). Western blotting for pS475-AKT, total AKT, pS73-c-Jun and tubulin at 15h post-treatment with the PI3-kinase inhibitor LY294002 (20μM) or forskolin (60 μM). (E) Effect of the LSD1 inhibitors OG-L002 (15μM) and S 2101 (20μM) on forskolin-mediated reactivation measured by Us11-GFP positive neurons. (F) Effect of the JMJD3

and UTX inhibitor GSK-J4 (2μM) on forskolin-mediated reactivation measured by Us11-GFP positive neurons. Statistical comparisons were made using Welch's t-test (E, F).

Previously, we found that Phase I reactivation is accompanied with a JNK-

dependent histone methyl/phospho (marked by H3K9me3/pS10) switch on lytic

promoters[15]. In cortical neurons, one study has found that hyperexcitability results in

increased H3K9me3/pS10[24]. Therefore, we were particularly interested to determine

whether forskolin treatment of sympathetic neurons triggered a histone S10

phosphorylation on H3K9me3. Forskolin triggered a transient increase in

H3K9me3/S10p at 5h post-treatment that had returned to baseline by 10h (Figure S3A

and B). This indicates that, in keeping with cortical neurons, forskolin induces a histone

H3K9me3/pS10 methyl/phospho switch on regions on cellular chromatin.

We next sought to determine whether the phospho/methyl switch that arises as a

result of hyperexcitability plays a role in Phase I of HSV reactivation. We therefore

investigated whether viral genomes were co-localized with H3K9me3/S10p following

forskolin treatment. To visualize HSV genomes, viral stocks were grown in the presence

of EdC as described previously[52,53]. Click-chemistry was performed on latently infected

and neurons following forskolin treatment. As shown in Figure 3A and B, viral genomes

co-localized with H3K9me3/pS10 following robust H3K9me3/S10p staining at 5h. The

percentage of viral genomes that co-localized with H3K9me3/S10p was significantly

increased compared to the unreactivated samples at 5h and 20h post-forskolin

treatment (Figure 3A).

Serine phosphorylation adjacent to a repressive lysine modification is thought to

permit transcription without the removal of the methyl group[17,24]. Therefore, we

investigated whether histone demethylase activity was required for the initial induction in

lytic gene expression following forskolin treatment. Previously, the H3K9me2 histone

demethylase LSD1 has been found to be required for full HSV reactivation[20,23], and in

our in vitro model this was determined by the production of infectious viral particles or

late gene synthesis at 48-72h post-reactivation[15]. Addition of two independent LSD1

inhibitors (OG-L002 and S 2102) inhibited Us11-GFP synthesis at 72h post-reactivation

(Figure 3C). Hence, LSD1 activity, and presumably removal of H3K9-methylation is

required for forskolin-mediated reactivation. However, LSD1 inhibition did not prevent

the initial induction of *ICP27* and *g*C mRNA expression at 20h post-forskolin treatment

(Figure 3D and E). Therefore, this initial wave of viral lytic gene expression following

forskolin-mediated reactivation is independent of histone H3K9 demethylase activity.

We previously found that H3K27me demethylase activity is required for full

reactivation but not the initial wave of gene expression[15]. However, because of the lack

of an antibody that specifically recognizes H3K27me3S28p and not also

H3K9me3S10p, we are unable at this point to investigate genome co-localization with

this combination of modifications. However, we could investigate the role of the

H3K27me demethylases in forskolin-mediated reactivation. Treatment of neurons with

the UTX/JMJD3 inhibitor GSK-J4[54] prevented the synthesis of Us11-GFP at 72h post-

reactivation, indicating that removal of K27 methylation is required full reactivation

(Figure 3F). However, the initial burst of gene expression (assessed by *ICP27* and *g*C

mRNA levels) was robustly induced at 20h post-forskolin treatment in the presence of

GSK-J4 (Figure 3G and H). Taken together, our data indicate that the initial phase of

gene expression following forskolin treatment is independent of histone demethylase

activity and therefore consistent with a role for a histone methyl/phospho switch in

permitting lytic gene expression.

<u>Forskolin-Mediated Reactivation Requires Neuronal Excitability</u>

Given that the HSV genome co-localized with regions of hyperexcitability-induced

changes in histone phosphorylation, we investigated whether reactivation was linked to

neuronal excitability. To inhibit action potential firing, we treated neurons with

tetrodotoxin (TTX), which inhibits the majority of the voltage gated sodium channels and

therefore depolarization. Addition of TTX significantly inhibited HSV reactivation

triggered by forskolin, as measured by Us11-GFP positive neurons at 72 hours post-

stimulus (Figure 4A). To further confirm a role for repeated action potential firing in

forskolin-mediated reactivation, we investigated the role of voltage gated potassium

channels, which are required for membrane repolarization. Addition of TEA, which

inhibits voltage-gated potassium channel activity, also blocked HSV reactivation

measure by Us11-GFP positive neurons at 3 days post-forskolin treatment (Figure S4B.

Taken together, these data indicate that action potential firing is required for forskolin-

mediated reactivation.

Increased levels of cAMP can act on nucleotide gated ion channels, including the

hyperpolarization activated cyclic nucleotide gated (HCN) channels. HCN channels are

Figure 4

<u>Figure 4. HSV Reactivation Mediated by Forskolin Requires Neuronal Activity.</u>
(A) Latently infected cultures were reactivated with forskolin in the presence of the voltage-gated sodium channel blocker tetrodotoxin (TTX; 1μM) and the number of Us11-GFP positive neurons quantified at 3 days post-reactivation. (B) Latently infected cultures were reactivated with forskolin in the presence of the voltage-gated potassium channel blocker Tetraethylammonium (TEA; 10 mM) and the number of Us11-GFP positive neurons quantified at 3 days post-reactivation. (C) Forskolin-mediated reactivation in the presence of the HCN channel blockers ZD 7288 (10μM) quantified as the numbers of Us11-GFP positive neurons at 3 days post-reactivation. (D) The effect of ZD 7288 on the HSV lytic gene transcript ICP27 during Phase I reactivation measured at 20h post-forskolin treatment by RT-qPCR. Individual experimental

replicates are represented. (E and F) Quantification of the relative nuclear staining for H3K9me3/S10p and γH2AX in SCG neurons at 5h post-forskolin treatment and in the presence of ZD 7288 from two independent experiments. Statistical comparisons were made using Welch's t-test (A-C), two-tailed unpaired t-test (D-F).

K^+ and Na^+ channels that are activated by membrane hyperpolarization[55,56]. In the

presence of high levels of cAMP, the gating potential of HCN channels is shifted in the

positive direction, such that HCN channels can open at resting membrane potential,

resulting in an increased propensity of neurons to undergo repeated firing[56-58]. HCN

channel activity can be blocked by ZD 7288, Ivabradine or cesium chloride. Addition of

ZD 7288 (Figure 4A), Ivabradine (Figure S4A) or CsCl (Figure S4B) all significantly

reduced HSV reactivation triggered by forskolin, as measured by Us-11 GFP positive

neurons at 3 days post-stimulus. To determine whether HCN channel activity was

required for the initial induction of HSV lytic mRNA expression, we assessed viral

mRNA expression during Phase I in the presence and absence of ZD 7288. Expression

of representative lytic mRNAs *ICP27* (Figure 4D), *UL30* and *gC* (Figure S4C and D)

were significantly decreased in the presence of ZD 7288 compared to the forskolin

treated neurons alone, and were not significantly increased compared to the mock

treated samples. Therefore, HCN channel activity is required for the initial induction of

lytic gene expression during Phase I reactivation mediated by forskolin.

We also confirmed that inhibition of HCN-channel activity affected the levels of

hyperexcitability-associated changes in histone post-translational modifications.

Addition of ZD 7288 resulted in significantly decreased staining intensities of both

H3K9me3/S10p and γH2AX at 5h post-forskolin treatment (Figure 4F and G), which was

the peak time-point for which we observed these changes upon forskolin treatment

alone (Figure S3 A and B). Therefore, activity of the HCN channels in response to

Figure S4

Figure S4. HSV Reactivation Mediated by Forskolin Requires Neuronal Activity. (A and B) Latently infected cultures were reactivated with forskolin in the presence of the HCN channel inhibitors ivabradine (20µM; A) and CsCl (3mM; B). Latently infected cultures were reactivated with forskolin in the presence of the HCN inhibitor ZD 7288 (10 µM) and viral lytic transcripts measured at 20h post-reactivation. Individual experimental replicates are represented. Statistical comparisons were made using Welch's t-test (A, B) or two-tailed unpaired t-test (C, D).

increased levels of cAMP results in hyperexcitability-associated changes in

histone modifications and reactivation of HSV from latent infection.

HSV Reactivation can be Induced by Stimuli that Directly Increase Neuronal Excitability

The role of ion-channel activity in forskolin-mediated reactivation prompted us to

investigate whether additional stimuli that induce hyperexcitability in neurons also

trigger HSV reactivation. We were also interested in whether reactivation required

chronic versus short term hyperexcitability. Increasing the extracellular concentration of

KCl is well-known to induce action potential firing. Therefore, we investigate the timing

of both KCl and forskolin-mediated hyperexcitability in HSV reactivation. Both of these

treatments triggered HSV reactivation more robustly if applied for 8h or more (Figure

5A). This indicates that chronic neuronal hyperactivity is important in inducing

reactivation of HSV.

To further clarify that hyperexcitability can directly trigger HSV reactivation, we

investigated the effects of removal from a TTX block on latently infected neurons.

Addition of TTX to neurons results in synaptic scaling, so that when the TTX is removed

the neurons enter a hyperexcitable state[59-62]. TTX was added to the neurons for 2 days

and then washed out. This resulted in a robust HSV reactivation as determined by

Us11-GFP synthesis (Figure 5B). We also investigated whether the JNK-cell stress

pathway was important in HSV reactivation in response to TTX-release. Addition of the

JNK inhibitor SP600125 or the DLK inhibitor, GNE-3511, blocked HSV reactivation

Figure 5

Figure 5. HSV Reactivation Triggered by Prolonged Neuronal Hyperexcitability is DLK/JNK Dependent. (A) Latently infected SCG cultures were treated with forskolin or KCl (55mM) for the indicated times followed by wash-out. Reactivation was quantified by number of Us11-GFP positive neurons at 3 days after the initial stimulus was added. (B) Latently infected neurons were placed in tetrodotoxin (TTX; 1μM) for 2 days and the TTX was then washed out. At the time of wash-out the JNK inhibitor SP600125 (20μM) or DLK inhibitor GNE-3511 (4μM) was added. Reactivation was quantified at 3-days post-wash-out. Individual experimental replicates are represented. Statistical comparisons were made using Welch's t-test.

following TTX-release. Therefore, directly inducing neuronal hyperexcitability triggers

HSV reactivation in a DLK/JNK-dependent manner.

IL-1β Triggers HSV Reactivation in Mature Neurons in a DLK and HCN Channel

Dependent Manner

Our data thus far point to a reactivation of HSV following increasing episodes of

neuronal hyperexcitability in a way that is requires activation of the JNK-cell stress

pathway. However, we wished to link this response to a physiological trigger that may

stimulate HSV reactivation *in vivo*. Increased HCN-channel activity has been associated

to inflammatory pain resulting from the activity of pyrogenic cytokines on neurons[63]. In

addition, IL-1β is known to act on certain neurons to induce neuronal excitation[38-40]. IL-

1β is released in the body during times of chronic, psychological stress. In addition, IL-

1β contributes to the fever response[31-34]. In sympathetic neurons, we found that

exposure of mature neurons to IL-1β induced an accumulation of the hyperexcitability-

associated histone post-translational modifications γH2AX and H3K9me3/S10p (Figure

6A-C). We did not observe the same changes for post-natal neurons. The reasons for

this maturation-dependent phenotype are unknown at this point but we hypothesize it

could be due to changes in the expression of cellular factors required to respond to IL-

1β. Therefore, these experiments were carried out on neurons that we more than 28

days old. The kinetics of induction of these histone modifications was different from

what we had previously observed for forskolin treatment, as both γH2AX and

H3K9me3/S10p steadily accumulated to 20h post-treatment. This likely reflects the

activation of upstream signaling pathways in response to IL-1β prior to inducing

Figure 6

Figure 6. IL-1β-Induced HSV Reactivation is Linked to Heightened Neuronal Activity and DLK Activation. (A) Adult P28 SCG neurons were treated with IL-1β (30ng/mL) for 15 hrs and stained for H3K9me3/S10p, γH2AX and beta II-tubulin to mark neurons. (B-C) Quantification of the intensity of H3K9me3/S10p and γH2AX in neuronal nuclei following forskolin treatment from two independent experiments. (D). Addition of IL-1β to latently infected cultures of mature SCG neurons triggers HSV reactivation. (E). Quantification of IL-1β induced reactivation in the presence of the voltage gated sodium channel blocker TTX (1μM), the

HCN channel blocker ZD 7288 (10µM) and the DLK inhibitor GNE-3511 (4µM). In D and E individual experimental replicates are represented. Statistical comparisons were made using or two-tailed unpaired t-test (B, C) or Welch's t-test (D, E)

neuronal excitation as IL-1β increases the expression of voltage-gated sodium

channels[40]. Importantly, IL-1β was able to trigger HSV reactivation in mature neurons

(Figure 5D). Reactivation was reduced in the presence of the HCN-channel inhibitor ZD

7288 and the voltage-gated sodium channel inhibitor TTX (Figure 6F), indicating that IL-

1β triggered reactivation via increasing neuronal activity. Importantly, addition of the

DLK inhibitor GNE-3511 blocked reactivation in response to IL-1β (Figure 6F).

Therefore, IL-1β can induce HSV reactivation that is both dependent on neuronal

activity and induction of the JNK neuronal cell stress response.

<u>Discussion</u>

As herpesviruses hide in the form of a latent infections of specific cell types, they

sense changes to the infected cell, resulting in the expression of viral lytic genes and

ultimately reactivation. HSV establishes latency in neurons and has previously been

found to respond to activation of a neuronal stress signaling pathway[15]. As an excitable

cell type, the function of neurons is to rapidly transmit stimuli via the firing of action

potentials, and under conditions of hyperexcitability, neurons increase their propensity

to fire repeated action potentials. Here we show that this state of hyperexcitability

induces HSV to undergo reactivation in a DLK/JNK dependent manner, indicating that

the virus responds to both activation of cell stress signaling and prolonged

hyperexcitability via a common pathway to result in reactivation. This common pathway

also permits viral lytic gene expression from silenced promoters without the requirement

of histone demethylase activity via a histone phospho/methyl switch. Conditions that

result in hyperexcitability include prolonged periods of stress and inflammation, which

are both linked to the release of IL-1β[31-34]. Consistent with this, here we show that IL-1β

induces DNA damage, and histone H3 phosphorylation in sympathetic neurons, which

are both markers of neuronal excitability. Importantly, IL-1β triggered HSV reactivation

that was dependent on neuronal activity and activation of DLK. Therefore, this study

identifies a physiological stimulus that induces HSV reactivation via increasing neuronal

excitability and places DLK/JNK signaling and a histone phospho/methyl switch as

central to HSV reactivation.

Neuronal hyperexcitability results in DNA damage followed by repair, which

together are thought to mediate the expression of cellular immediate early genes[29,30].

Here we show that forskolin treatment and IL-1β also induce DNA damage in

sympathetic neurons. Previously, HSV reactivation has been found to occur following

inhibition of DNA damage, inhibition of repair and exogenous DNA damage[12]. In the

context of repair inhibition or exogenous DNA damage, reactivation was dependent on

dephosphorylation of AKT by the PHLPP1 phosphatase and activation of JNK, and

therefore feeds into the same pathway as PI3K-inhibition. However, we did not observe

decreased AKT phosphorylation in response to forskolin treatment, indicating that the

mechanism of reactivation is distinct following physiological levels of DNA damage

resulting from neuronal hyperexcitability versus perturbation of the damage/repair

pathways.

Conditions that result in hyperexcitability include prolonged periods of stress and

inflammation, which are both linked to the release of IL-1β[31-34]. Consistent with these

findings, we show that IL-1β treatment induces two markers of neuronal excitability,

DNA damage and histone H3 phosphorylation, in primary sympathetic neurons. The IL-

1 family of cytokines act via the IL-1 receptor to activate downstream signaling

pathways [64]. IL-1β is released systemically during prolonged periods of psychological

stress and upon infection via activation of the inflammasome[31-34]. IL-1α, which also

signals via the IL-1R, is released locally as an alarmin. Interesting IL-1α and IL-1β are

found at high levels in keratinocytes and are released upon HSV-1 infection[41], where

they can mediate antiviral responses in underlying stromal fibroblasts and endothelial

cells. Antiviral responses mediated by IL-1 signaling have been found to involve NF-κB,

IRF3 and/or IRF1[42]. The downstream signaling elicited by IL-1 in neurons has not been

clearly defined and likely varies between different subtypes of neurons. NF-κB has been

reported to be absent in certain subtypes of neurons but constitutively active in

others[65,66], and a recent study suggests that NF-κB levels increase with neuronal

maturation[67], which may be why we only observed IL-1-mediated reactivation in mature

neurons. Additional studies have found a role for p38MAPK signaling and AKT/mTOR

signaling in neuronal IL-1-mediated responses[68,69]. A common feature of IL-1 signaling

in neurons is increased excitability, which has been associated with neurotransmitter

release, and mediates a variety of physiological responses including behavior

modulation and an intersection with the hosts' immune response[38-40]. IL-1 is also

associated with pathological conditions, including neurodegenerative disease such as

Alzheimer's disease[70]. There are multiple studies linking HSV-1 infection to the

progression of Alzheimer's[2]; therefore, the combination of both HSV infection and

increased IL-1 could have a feed forward effect on the progression of Alzheimer's

disease by promoting increased reactivation of HSV from latency.

Experiments using primary neuronal *in vitro* model systems and inducing

reactivation by PI3-kinase inhibition have shown that reactivation progresses over two

phases. Phase 1 involves the synchronous up-regulation of lytic gene expression that

occurs independently of the viral transactivator VP16 and the activity of cellular histone

demethylases[15,18]. A population of neurons progress to full reactivation (Phase II), which

is dependent on both VP16 and HDM activity[15,18]. We previously found that lytic gene

expression in Phase I is DLK/JNK dependent and is corelated with a JNK-dependent

histone methyl/phospho switch on lytic gene promoters[15]. Here we demonstrate that a

Phase 1 wave of viral gene expression that is dependent on activation of JNK but not

histone demethylases also occurs in response to neuronal hyperexcitability. The co-

localization of viral genomes with H3K9me3/pS10 indicates that a histone

methyl/phospho switch also permits lytic gene expression to occur following forskolin

treatment in a manner that is independent of HDM activity. This indicates that

reactivation proceeds via a Phase 1-wave of gene expression in response to multiple

different stimuli. However we note that there may be differences in the mechanism and

kinetics of reactivation with different stimuli and/or strains of HSV-1 as reactivation

triggered by axotomy may bypass Phase I[19,20] and reactivation induced in vivo by heat

shock with a more pathogenic strain of HSV triggered more rapid reactivation [71]. It will

be especially interesting to determine in the future whether there are differences in the

progression to reactivation with different strains of HSV. Ultimately, these reactivation

kinetics may relate differences in the epigenetic structures of viral genomes that vary

based on virus strains or differential manipulation of host-cell signaling pathways.

The Wilcox lab demonstrated in 1992 that reactivation can be induced by

forskolin, and it has since been used as a trigger in multiple studies[25-28]. However, the

mechanism by which increasing levels of cAMP induces lytic gene expression was not

known. Here we link cAMP-induced reactivation to the excitation state of the neuron and

show that the initial induction of viral gene expression is dependent on DLK and JNK

activity but independent of CREB and PKA. The activity of PKA may be required for full

reactivation, which is also consistent with a role for PKA in overcoming repression of the

related Pseudorabies Virus during *de novo* axonal infection[72]. Our data also suggest

that CREB may be involved in the progression to full reactivation. However, the

mechanism of action of the inhibitor used here, 666-15, is not entirely clear. It has been

reported as preventing CREB-mediated gene expression, but may act to prevent

recruitment of histone acetyltransferases[73]. Therefore, inhibition of Phase II reactivation

by 666-15 would be consistent with more large-scale chromatin remodeling on the viral

genome at this stage. In addition, previous work has identified a role for inducible cAMP

early repressor (ICER) in HSV reactivation[26]. ICER is a repressor of gene expression

via heterodimerization with members of the CREB/ATF family of transcription factors.

CREB expression is also known to be down-regulated by loss of NGF-signaling[74].

Therefore, it is conceivable that inhibition, rather than activation, of CREB is important

for reactivation of HSV from latency.

Previously, we found that JNK activation by DLK is required for reactivation

following interruption of the NGF-signaling pathway. Here we find that forskolin and IL-

1β-mediated reactivation also required both DLK activity, further reinforcing the central

role of DLK and JNK in reactivation of HSV from latency. DLK is known as a master

regulator of neuronal response to stress stimuli and mediates whole cell death, axon

pruning, regeneration or generation depending on the nature of the stimuli. However, it

has not before been linked to neuronal hyperexcitability or the response to IL-1β

signaling. The known mechanisms of DLK activation include loss of AKT activation and

phosphorylation by PKA, neither of which could be linked to HSV reactivation mediated

by forskolin in this study. Following activation by DLK, one mechanism by which JNK is

thought to permit lytic gene expression is via recruitment to viral promoters and histone

phosphorylation. However, it is likely that there are additional, JNK-dependent effects

including activation of pioneer or transcription factors that also mediate viral gene

expression. Further insight into how HSV has hijacked this cellular pathway to induce

lytic gene expression may lead to novel therapeutics that prevent reactivation, in

addition to providing information on how viral gene expression initiates from promoters

assembled into heterochromatin.

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Materials and Methods

Reagents

Compounds used in the study are as follows: Acycloguanosine, FUDR, Uridine,

SP600125, GNE-3511, GSK-J4, L-Glutamic Acid, and Ivabradine (Millipore Sigma);

Forskolin, LY 294002, 666-15, SQ 22536, KT 5720, Tetraethylammonium chloride,

Cesium chloride, OG-L002, S2101, Tetrotdotoxin, and ESI-09 (Tocris); 1,9-dideoxy

Forskolin, ZD 7288 and 8-bromo-cyclic AMP (Cayman Chemicals); Nerve Growth

Factor 2.5S (Alomone Labs); Primocin (Invivogen); Aphidicolin (AG Scientific); IL-1β

(Shenandoah Biotechnology); WAY-150138 was kindly provided by Pfizer, Dr. Jay

Brown and Dr. Dan Engel at the University of Virginia, and Dr. Lynn Enquist at

Princeton University. Compound information and concentrations used can be found

below in Table S1.

Preparation of HSV-1 Virus Stocks

HSV-1 stocks of eGFP-Us11 Patton were grown and titrated on Vero cells obtained

from the American Type Culture Collection (Manassas, VA). Cells were maintained in

Dulbecco's Modified Eagle's Medium (Gibco) supplemented with 10% FetalPlex

(Gemini Bio-Products) and 2 mM L-Glutamine. eGFP-Us11 Patton (HSV-1 Patton strain

with eGFP reporter protein fused to true late protein Us11[43]) was kindly provided by Dr.

Ian Mohr at New York University.

Primary Neuronal Cultures

Sympathetic neurons from the Superior Cervical Ganglia (SCG) of post-natal day 0-2

(P0-P2) or adult (P21-P24) CD1 Mice (Charles River Laboratories) were dissected as

previously described[15]. Rodent handling and husbandry were carried out under animal

protocols approved by the Animal Care and Use Committee of the University of Virginia

(UVA). Ganglia were briefly kept in Leibovitz's L-15 media with 2.05 mM L-Glutamine

before dissociation in Collagenase Type IV (1 mg/mL) followed by Trypsin (2.5 mg/mL)

for 20 minutes each at 37 °C. Dissociated ganglia were triturated, and approximately

10,000 neurons per well were plated onto rat tail collagen in a 24-well plate.

Sympathetic neurons were maintained in CM1 (Neurobasal® Medium supplemented

with PRIME-XV IS21 Neuronal Supplement (Irvine Scientific), 50 ng/mL Mouse NGF

2.5S, 2 mM L-Glutamine, and Primocin). Aphidicolin (3.3 µg/mL), Fluorodeoxyuridine

(20 µM) and Uridine (20 µM) were added to the CM1 for the first five days post-

dissection to select against proliferating cells.

Establishment and Reactivation of Latent HSV-1 Infection in Primary Neurons

Latent HSV-1 infection was established in P6-8 sympathetic neurons from SCGs.

Neurons were cultured for at least 24 hours without antimitotic agents prior to infection.

The cultures were infected with eGFP-Us11 (Patton recombinant strain of HSV-1

expressing an eGFP reporter fused to true late protein Us11). Neurons were infected at

a Multiplicity of Infection (MOI) of 7.5 PFU/cell (assuming 1.0×10^4 neurons/well/24-well

plate) in DPBS +$CaCl_2$ +$MgCl_2$ supplemented with 1% Fetal Bovine Serum, 4.5 g/L

glucose, and 10 µM Acyclovir (ACV) for three hours at 37 ˚C. Post-infection, inoculum

was replaced with CM1 containing 50 µM ACV for 5-6 days, followed by CM1 without

ACV. Reactivation was carried out in DMEM/F12 (Gibco) supplemented with 10% Fetal

Bovine Serum, Mouse NGF 2.5S (50 ng/mL) and Primocin. Inhibitors were added either

one hour prior to or concurrently with the reactivation stimulus. WAY-150138 (2-10

µg/mL) was added to reactivation cocktail to limit cell-to-cell spread. Reactivation was

quantified by counting number of GFP-positive neurons or performing Reverse

Transcription Quantitative PCR (RT-qPCR) of HSV-1 lytic mRNAs isolated from the

cells in culture.

Analysis of mRNA expression by reverse-transcription quantitative PCR (RT-

qPCR)

To assess relative expression of HSV-1 lytic mRNA, total RNA was extracted from

approximately 1.0×10^4 neurons using the Quick-RNA™ Miniprep Kit (Zymo Research)

with an on-column DNase I digestion. mRNA was converted to cDNA using the

SuperScript IV First-Strand Synthesis system (Invitrogen) using equal amounts of RNA

(20-30 ng/reaction). To assess viral DNA load, total DNA was extracted from

approximately 1.0×10^4 neurons using the Quick-DNA™ Miniprep Plus Kit (Zymo

Research). qPCR was carried out using *Power* SYBR™ Green PCR Master Mix

(Applied Biosystems). The relative mRNA or DNA copy number was determined using

the Comparative C_T ($\Delta\Delta C_T$) method normalized to mRNA or DNA levels in latently

infected samples. Viral RNAs were normalized to mouse reference gene GAPDH. All

samples were run in duplicate on an Applied Biosystems™ QuantStudio™ 6 Flex Real-

Time PCR System. Primers used are described in Table S2.

Western Blot Analysis

Neurons were lysed in RIPA Buffer with cOmplete, Mini, EDTA-Free Protease Inhibitor

Cocktail (Roche) and PhosSTOP Phosphatase Inhibitor Cocktail (Roche) on ice for one

hour with regular vortexing to aid lysis. Insoluble proteins were removed via

centrifugation, and lysate protein concentration was determined using the Pierce

Bicinchoninic Acid Protein Assay Kit (Invitrogen) using a standard curve created with

BSA standards of known concentration. Equal quantities of protein (generally 20-50 µg)

were resolved on 4-20% gradient SDS-Polyacrylamide gels (Bio-Rad) and then

transferred onto Polyvinylidene difluoride membranes (Millipore Sigma). Membranes

were blocked in PVDF Blocking Reagent for Can Get Signal (Toyobo) for one hour.

Primary antibodies were diluted in Can Get Signal Immunoreaction Enhancer Solution 1

(Toyobo) and membranes were incubated overnight at 4°C. Antibodies and

concentrations are described in Table S3 below. HRP-labeled secondary antibodies

were diluted in Can Get Signal Immunoreaction Enhancer Solution 2 (Toyobo) and

membranes were incubated for one hour at room temperature. Blots were developed

using Western Lightning Plus-ECL Enhanced Chemiluminescence Substrate

(PerkinElmer) and ProSignal ECL Blotting Film (Prometheus Protein Biology Products)

according to manufacturer's instructions. Blots were stripped for reblotting using

NewBlot PVDF Stripping Buffer (Licor).

Immunofluorescence

Neurons were fixed for 15 minutes in 4% Formaldehyde and blocked in 5% Bovine

Serum Albumin and 0.3% Triton X-100 and incubated overnight in primary antibody.

Antibodies and concentrations are described in Table S4 below. Following primary

antibody treatment, neurons were incubated for one hour in Alexa Fluor® 488-, 555-,

and 647-conjugated secondary antibodies for multi-color imaging (Invitrogen). Nuclei

were stained with Hoechst 33258 (Life Technologies). Images were acquired using an

sCMOS charge-coupled device camera (pco.edge) mounted on a Nikon Eclipse Ti

Inverted Epifluorescent microscope using NIS-Elements software (Nikon). Images were

analyzed and intensity quantified using ImageJ.

Click Chemistry

Click chemistry was carried out a described previously[52] with some modifications.

Neurons were washed with CSK buffer (10 mM HEPES, 100 mM NaCl, 300 mM

Sucrose, 3 mM $MgCl_2$, 5 mM EGTA) and simultaneously fixed and permeabilized for 10

minutes in 1.8% methonal-free formaldehyde (0.5% Triton X-100, 1%

phenylmethylsulfonyl fluoride (PMSF)) in CSK buffer, then washed twice with PBS

before continuing to the click chemistry reaction and immunostaining. Samples were

blocked with 3% BSA for 30 minutes, followed by click chemistry using EdC-labelled

HSV-1 DNA and the Click-iT EdU Alexa Flour 555 Imaging Kit (ThermoFisher Scientific,

C10638) according to the manufacturer's instructions. For immunostaining, samples

were incubated overnight with primary antibodies in 3% BSA. Following primary

antibody treatment, neurons were incubated for one hour in Alexa Fluor® 488-, 555-,

and 647-conjugated secondary antibodies for multi-color imaging (Invitrogen). Nuclei

were stained with Hoechst 33258 (Life Technologies). Images were acquired at 60x

using an sCMOS charge-coupled device camera (pco.edge) mounted on a Nikon

Eclipse Ti Inverted Epifluorescent microscope using NIS-Elements software (Nikon).

Images were analyzed and intensity quantified using ImageJ.

Statistical Analysis

Power analysis was used to determine the appropriate sample sizes for statistical

analysis. All statistical analysis was performed using Prism V8.4. Welch's t-test was

used for GFP fluorescence experiments and two-tailed unpaired t-test was used for RT-

qPCR and IF experiments; specific analyses are included in the figure legends. EdC

virus and H3K9me3S10/p co-localization was quantified using ImageJ after sample

blinding. Mean fluorescence intensity of γH2AX was quantified using ImageJ.

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865 Supplemental Tables of Reagents

866 Table S1: Compounds Used and Concentrations

Compound	Supplier	Identifier	Concentration
Acycloguanosine	Millipore Sigma	A4669	10 µM, 50 µM
FUDR	Millipore Sigma	F-0503	20 µM
Uridine	Millipore Sigma	U-3003	20 µM
SP600125	Millipore Sigma	S5567	20 µM
GNE-3511	Millipore Sigma	533168	4 µM
GSK-J4	Millipore Sigma	SML0701	2 µM
L-Glutamic Acid	Millipore Sigma	G5638	3.7 µg/mL
Forskolin	Tocris	1099	60 µM
LY 294002	Tocris	1130	20 µM
666-15	Tocris	5661	2 µM
SQ 22,536	Tocris	1435	50 µM
KT 5720	Tocris	1288	3 µM
TEA	Tocris	3068	10 mM
CsCl	Tocris	4739	3 mM
OG-L002	Tocris	6244	15 µM, 30 µM
S2101	Tocris	5714	10 µM, 20 µM
Tetrodotoxin	Tocris	1069	1 µM
ESI-09	Tocris	4773	10 µM
ZD 7288	Cayman	15228	20 µM
8-bromo-cyclic AMP	Cayman	14431	125 µM

NGF 2.5S	Alomone Labs	N-100	50 ng/mL
Primocin	Invivogen	ant-pm-1	100 µg/mL
Aphidicolin	AG Scientific	A-1026	3.3 µg/mL
IL-1β	Shenandoah Bio.	100-167	30ng/mL
WAY-150138	Pfizer	NA	10 µg/mL

869 Table S2: Primers Used for RT-qPCR

Primer	Sequence 5' to 3'
mGAP 1SF	CAT GGC CTT CCG TGT GTT CCT A
mGAP 1SR	GCG GCA CGT CAG ATC CA
ICP27 F	GCA TCC TTC GTG TTT GTC ATT CTG
ICP27 R	GCA TCT TCT CTC GAC CCC CG
ICP8 1SF	GGA GGT GCA CCG CAT ACC
ICP8 1SR	GGC TAA AAT CCG GCA TGA AC
ICP4 F	TGC TGC TGC TGT CCA CGC
ICP4 R	CGG TGT TGA CCA CGA TGA GCC
UL30 F	CGC GCT TGG CGG GTA TTA ACA T
UL30 R	TGG GTG TCC GGC AGA ATA AAG C

UL48 F	TGC TCG CGA ATG TGG TTT AG
UL48 R	CTG TTC CAG CCC TTC ATG TT
gC #1 F	GAG TTT GTC TGG TTC GAG GAC
gC #1R	ACG GTA GAG ACT GTG GTG AA

873 Table S3: Antibodies Used for Western Blotting and Concentrations

Antibody	Supplier	Identifier	Concentration
Rb Phospho-Akt (S473)	CST	4060	1:500
Rb Akt (pan)	CST	C67E7	1:1000
Rb Phospho-c-Jun (S73)	CST	3270	1:500
Ms Monoclonal α-Tubulin	Millipore Sigma	T9026	1:2500
HRP Goat Anti-Rabbit IgG Antibody (Peroxidase)	Vector	PI-1000	1:10000
HRP Horse Anti-Mouse IgG Antibody (Peroxidase)	Vector	PI-2000	1:10000

877   Table S4: Antibodies Used for Immunofluorescence and Concentrations

Antibody	Supplier	Identifier	Concentration
Rb H3K9me3S10P	Abcam	ab5819	1:250
Ch Beta-III Tubulin	Millipore sigma	AB9354	1:1000
Ms γH2A.X	CST	80312S	1:100
Ms c-Fos	Novus	NB110-75039	1:125
F(ab')2 Goat anti Mouse IgG (H+L) Alexa Fluor® 647	Thermo Fisher	A21237	1:1000
F(ab') Goat anti Rabbit IgG (H+L) Alexa Fluor® 555	Thermo Fisher	A21425	1:1000
Goat anti Chicken IgY (H+L) Alexa Fluor® 647	abcam	ab150175	1:1000
Goat Anti-Chicken IgY H&L (Alexa Fluor® 488) preabsorbed	abcam	ab150173	1:1000
F(ab')2 Goat anti-Rabbit IgG (H+L) Alexa Fluor® 488)	Thermo Fisher	B40922	1:1000

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