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Helminth infection reactivates latent γ -herpesvirus via cytokine competition at a viral promoter — Source link \square

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Published on: 01 Aug 2014 - Science (American Association for the Advancement of Science)

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gene. Also, mutations in IZKF3, which also had HIV integrations detected in two of our three participants, was recently associated with a form of acute lymphoblastic leukemia (35). Fourth, as somatic mutations that "drive" cancers are estimated to only convey a 0.4% growth advantage (36), HIV integration into genes with subtle enhancement of cell proliferation may be difficult to detect as clonal due to our limited sampling.

HIV-infected cells that express viral proteins are likely to be eliminated by immune surveillance, or virus replication may lead to cell lysis. Whether the proliferating and persisting HIV-infected cells that we describe harbor replication-competent virus is critical to defining their role in perpetuating the infectious virus reservoir. Undoubtedly, some clonal populations persist due to defects in expression of the proviral genome (9, 10, 37). Although we did not evaluate viral sequences for replication competency, lethally hypermutated viral genomes were linked to three integration sites (Fig. 2). However, cells producing viremias with identical env sequences have been shown to harbor replication-competent virus (38). Also, approximately 12% of proviruses refractory to in vitro induction were found to have intact genomes and may be infectious (27). Although transcriptional interference was not detected in the aforementioned noninduced viral transcripts (26, 27), others have observed that the site of integration may cause transcriptional interference (34, 39-43).

In conclusion, HIV integration into genes associated with cancer or cell cycle regulation appears to confer a survival advantage that allows these cells to persist during suppressive ART, with cell proliferation appearing to serve as an important mechanism of HIV persistence. To be defined are the mechanisms contributing to cell proliferation, the role of proliferating cells in perpetuating the infectious virus reservoir, and whether therapies that target HIV-infected proliferating cells, specific genes, or their products may contribute to a curative strategy for HIV infection.

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COINFECTION

ACKNOWLEDGMENTS

We wish to express gratitude to the study participants who donated their time and provided specimens, and to N. H. Tobin, A. J. Melvin, and K. M. Mohan for their contributions to this study. Additionally, we thank Y. C. Ho, R. and J. Siliciano, H. Imamichi and C. Lane, and N. Malani and F. Bushman for providing their HIV integration site data. HIV env sequences generated from specific integration sites were deposited in GenBank under accession numbers (KM025062 to KM025124 and KM201287 to KM201295). Additional HIV env sequences were deposited in GenBank under accession numbers AY075701 to AY077450 (44), AY483287 to AY484389 (2), and KC314011 to KC315287 (8). Sequences of HIV-3'-LTR and adjacent chromosome integration sites are available in table S3. Other data are tabulated in the main paper and in the supplementary materials. Integration site sequences and HIV sequence alignments are available for download at http:// mullinslab.microbiol.washington.edu/publications/supplemental html. These studies were funded by R01AI091550 (L.M.F.). K23AI077357 (T.A.W.), R01 AI111806 (J.I.M.), the Molecular Profiling and Computational Biology Core of the University of Washington's Center for AIDS Research (P30 AI027757), and the Canadian Institutes of Health (HIV/AIDS Research Initiative award 201311CVI-322424-244686) (C.Y.K.C.). We wish to thank A. Embry and D. Lawrence at the National Institutes of Health for their support. The authors report no conflicts of interest related to this research.

SUPPLEMENTARY MATERIALS

www.sciencemag.org/content/345/6196/570/suppl/DC1 Materials and Methods Figs. S1 to S3 Tables S1 to S3 References (46-59)

20 May 2014; accepted 27 June 2014 Published online 10 July 2014; 10.1126/science.1256304

Helminth infection reactivates latent γ -herpesvirus via cytokine

competition at a viral promoter

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Mammals are coinfected by multiple pathogens that interact through unknown mechanisms. We found that helminth infection, characterized by the induction of the cytokine interleukin-4 (IL-4) and the activation of the transcription factor Stat6, reactivated murine γ -herpesvirus infection in vivo. IL-4 promoted viral replication and blocked the antiviral effects of interferon- γ (IFN γ) by inducing Stat6 binding to the promoter for an important viral transcriptional transactivator. IL-4 also reactivated human Kaposi's sarcoma-associated herpesvirus from latency in cultured cells. Exogenous IL-4 plus blockade of IFN γ reactivated latent murine γ -herpesvirus infection in vivo, suggesting a "two-signal" model for viral reactivation. Thus, chronic herpesvirus infection, a component of the mammalian virome, is regulated by the counterpoised actions of multiple cytokines on viral promoters that have evolved to sense host immune status.

ammals are populated by many chronic viruses, termed the virome, which can regulate host physiology and disease susceptibility (1). For example, more than 90% of humans are latently infected with herpesviruses that, after clearance of acute infec-

tion, produce little infectious virus and often cause no overt disease. Like the human γ -herpesviruses Epstein-Barr virus (EBV) and Kaposi's sarcomaassociated herpesvirus (KSHV), murine γ -erpesvirus-68 (MHV68) establishes lifelong latency. Studies in this model system showed that the cytokine ¹Department of Pathology and Immunology, Washington University School of Medicine, St. Louis, MO 63110, USA.
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Fig. 1. Challenge with *H. polygyrus* and *S. mansoni* eggs reactivates MHV68. (A) C57BL6/

J mice were infected intraperitoneally (i.p.) with MHV68-M3-FL and challenged with H. polygyrus 42 days later. Mice were imaged before H. polygyrus infection (day 0, d0) and 5, 7, and 9 days after. Three representative mice imaged on days 0 and 7 are shown. (B) Total flux (photons per second, p/s) was quantitated for mice in two independent experiments for the timecourse after infection with H. polygyrus. Data from four independent experiments at day 7 after H. polygyrus (H.p.) infection are also shown. (C) C57BL/6J mice were infected intranasally (i.n.) with MHV68-M3-FL. The diagram indicates the timecourse of the experiment and challenge with Sm eggs or phosphate-buffered saline (PBS) as a control. Mice were injected with D-Luciferin and imaged before intravenous (i.v.) challenge with Sm eggs (day 0). They were subsequently imaged 5, 8, and 11 days after challenge with Sm eggs. Three representative mice imaged on days 0 and 8 are shown. (D) Total flux was quantitated from mice in two independent experiments after Sm egg challenge. Symbols represent individual mice, and the mean and standard error are indicated. *P < 0.05, **P < 0.01, ***P < 0.001 by two-way repeated measures analysis of variance (ANOVA) with Tukey's and Bonferroni's post-test.

Fig. 2. IL-4 and IFN_Y signatures identified in different macrophage populations during MHV68 infection. (A) GSEA of virus-positive and -negative cells sorted from the peritoneum of MHV68infected mice compared to BMDMs stimulated with IL-4 or IFN_Y/LPS. (B) C57BL/6J mice or YARG/ R26-stop-RFP mice were either mock infected or infected with MHV68-cre. CD11b+F4/80+ cells were gated and examined for RFP and YFP expression. Representative plots from two independent experiments, with three to five mice per experiment, are shown. (C) Quantitation of fluorescenceactivated cell sorting (FACS) analysis in (B), with each symbol representing a single mouse. PEC, peritoneal exudate cells. interferon- γ (IFN γ) inhibits MHV68 replication and reactivation from macrophages, a major cellular site for latency (2–5); controls persistent replication in vivo (6); and is present at low amounts during latency (7).

Many people around the world are coinfected with herpesviruses and intestinal helminths. Although herpesviruses can modulate immunity to harm or benefit the host (7–11), the effects of helminth coinfection on chronic herpesvirus infection are unexplored. Intestinal helminths generate strong T helper 2 ($T_{\rm H}$ 2)-driven cytokine responses, which counter the biological effects of IFN γ and drive the activation of macrophages with an M2 (immunoregulatory) rather than M1 (proinflammatory) phenotype (*12*). Parasitic worms may influence control of pathogens—including *Mycobacterium tuberculosis*, HIV, and *Plasmodium* species in humans—but there are few studies elucidating the mechanisms behind this immunomodulation (*13*). Thus, we considered the hypothesis that parasite infection would induce MHV68 reactivation in vivo.

We examined the effects of acute infection with *Heligmosomoides polygyrus* or *Schistosomiasis* mansoni (Sm) egg administration on MHV68



reactivation from latency using a MHV68 virus expressing luciferase under the control of a lytic viral promoter upon reactivation from latency in vivo (MHV68-M3-FL) (14). Both acute *H. polygyrus* infection and *Sm* egg challenge reactivated MHV68 infection (Fig. 1, A to D). Mice latently infected for more than 100 days also showed increased luciferase expression after *Sm* egg challenge (fig. S1). By contrast, infection with the systemic bacteria *Listeria monocytogenes* did not stimulate viral reactivation (fig. S2). Thus, responses to either a nematode parasite or trematode eggs

induced herpesvirus reactivation, suggesting a role for $\rm T_{H}2$ cytokines in viral reactivation.

To determine whether $T_{\rm H}2$ cytokines affect latently infected macrophages, we compared host gene expression patterns in virally infected and uninfected macrophages during chronic infection. We engineered MHV68 to express cre-recombinase (MHV68-cre) from a locus permitting heterologous gene expression without altering viral replication or reactivation (*15*) (fig. S3). Reporter mice in which fluorescent protein expression is induced by cre recombination [Rosa26-floxed stop-eYFP] or tandem dimer (td)RFP (eYFP, enhanced yellow fluorescent protein; RFP, red fluorescent protein) (*16*)] were infected with MHV68-cre. Virus-positive and -negative cells sorted from latently infected mice (fig. S3G) were subjected to RNAseq analysis. Transcription in these cells was compared to that in bone marrow–derived macrophages (BMDMs) stimulated with interleukin-4 (IL-4) (M2) or IFN γ plus lipopolysaccharide (LPS) (M1) compared to untreated BMDMs (M0). Gene set enrichment analysis (GSEA) revealed that genes up-regulated in M1 BMDMs were enriched in





tors expressing luciferase under control of four different gene 50 promoters (23, 24). Cells were then treated with or without IL-4 or IL-13 for 24 hours, lysed, and assayed for luciferase activity. (**F**) Cells were transfected with the N4/N5 promoter luciferase construct and treated with IL-4, IFN₇, or both. (**G**) N4/N5 luciferase mutants were transfected into RAW264.7 cells and assayed for sensitivity to IL-4. (**H**) RAW264.7 cells were infected with MHV68 at multiplicity of infection = 5 and treated with IL-4 for 8 hours. After ChIP with Stat6 antibody, quantitative PCR (qPCR) was performed for the N4/N5 promoter region or vascular endothelial growth factor (VEGF). Percentage of input after normalizing to immunoglobulin G control was calculated for both N4/N5 and VEGF. One experiment representative of four independent experiments is shown. Error bars indicate SEM. n.d. not detected; n.s. not significant. **P* < 0.05, ***P* < 0.01, *****P* < 0.001 by *t* test or one-way ANOVA with Sidak's multiple comparisons test.

virus-positive macrophages, whereas genes upregulated in M2 BMDMs were enriched in virusnegative cells (Fig. 2A and table S1). This was consistent with the role of IFNy, which drives M1 macrophage polarization, in inhibiting MHV68 replication and reactivation (3, 4). We therefore tested whether latent MHV68 infection was restricted to M1-type macrophages by infecting tdRFP mice carrying the Arginase-1 (Arg1)-YFP reporter (YARG, a marker for macrophages stimulated with $T_{\rm H}2$ cytokines) (17) with MHV68-cre. Surprisingly, virus-positive macrophages were either positive or negative for Arg1 (Fig. 2, B and C), suggesting that despite the role for IFN γ in controlling chronic MHV68 infection, at least some virus-infected cells were exposed to cytokines that drive Arg-1 expression in vivo.

Along with the observation that $T_{\rm H}2$ cytokineinducing parasites promote reactivation from latency, the presence of an IL-4 signature in some virus-infected macrophages suggests a role for

IL-4 in viral infection. We tested this by determining the effect of IL-4 on MHV68 replication in BMDMs. Treatment with IL-4 increased Arg1 expression (fig. S4A), consistent with M2 polarization (18). As expected, few infected BMDMs expressed lytic viral antigens upon MHV68 infection (19). However, IL-4 pretreatment increased the number of BMDMs expressing viral proteins and enhanced viral replication (Fig. 3A and fig. S4, B to D); it also increased infection of transformed RAW264.7 macrophages (fig. S4E). Treatment with IL-4 after MHV68 infection increased viral replication (fig. S4F), indicating that IL-4 acts on replication rather than by increasing the number of infected cells. Enhancement of replication was dependent on the T_H2-associated transcription factor Stat6 (Fig. 3A) and occurred with IL-13 stimulation, another T_H2-associated cvtokine that uses the IL-4 receptor $\boldsymbol{\alpha}$ chain and signals via Stat6 (Fig. 3B and fig. S4, D and G). The T_H2 cytokine IL-5, which does not signal

through Stat6, did not promote MHV68 replication (Fig. 3B).

After treatment with IL-4, the majority of infected cells did not express the M2 markers CD206 or Arg1 (Fig. 3A and fig. S4A), suggesting that not all IL-4-induced changes in macrophage differentiation are required for enhanced MHV68 replication (20). Etomoxir blocks IL-4-induced changes in fatty acid oxidation (21) and upregulation of CD206 (fig. S5A) but did not block enhancement of MHV68 replication by IL-4 (fig. S5B). Moreover, IL-4 enhanced replication in the absence of peroxisome proliferator-activated receptor γ (PPAR γ) or ARG1, key proteins involved in M2 macrophage function, or inducible nitric oxide synthase, an essential protein in M1 macrophage function (fig. S5, C to G) (20). Importantly, IL-4 antagonized IFNy-mediated suppression of viral replication (Fig. 3C) (3). Because Stat6 antagonizes Stat1 (22), we tested whether IL-4 promoted virus replication in the absence of Stat1.



Fig. 4. MHV68 and KSHV reactivation from latency is regulated by IL-4. (A) C57BL6/J mice

were infected with MHV68-M3-FL intraperitoneally. Forty-two days later, mice were imaged for luciferase expression (d0) and then received isotype control (PIP), anti-IFNγ (H22), IL-4c, or both anti-IFNγ and IL-4c. On day 44, mice received a second dose of IL-4c or PBS. Mice were imaged 5 days after the first treatment, and total flux from the abdominal region was quantitated. Data from four representative mice are shown. (B) Quantitation of total flux from three individual experiments described in (A) is shown. Each symbol represents an individual mouse. Bars denote means; and error bars represent standard errors. (C) WT or Stat6KO mice were treated as in (A) with anti-IFNγ/IL-4c and imaged 5 days later. Bars denote the means of individual mice (symbols); error bars represent standard errors. (D) The experimental set-up was the same as that of Fig. 1A. Latently infected WT and Stat6KO mice infected with MHV68-M3-FL were infected with H. polygyrus or treated with PBS on day 42, and reactivation was quantitated by luciferase induction 7 days later. (E) BCBL-1 cells were treated with IL-4 for 3 or 5 days or 12-0tetradecanoylphorbol-13-acetate (TPA) for 48 hours, and viral gene expression was analyzed. The fold increase in gene expression over mock after normalization of GAPDH is shown. Data are from three independent experiments. (F) Supernatants from cells treated in (E) were collected, and virus was isolated by centrifugation. Viral genome copy number was assayed by qPCR using serial diluted LANA expression plasmid as a standard curve. One experiment representative of two independent experiments is shown. For luciferase experiments: *P < 0.05, **P < 0.01. n.s., not significant by two-way repeated measures ANOVA with Tukey's and Bonferroni's post-test. Error bars indicate SEM.

IL-4 increased virus replication in Stat1-deficient BMDMs (fig. S6).

Previously, we found that IFNy-mediated suppression of viral replication was associated with inhibition of promoters driving expression of the essential viral latent-to-lytic switch gene (gene 50) (3, 23). IL-4 antagonizes IFNy-mediated suppression of gene 50 transcription (Fig. 3D). This effect was specific to the viral promoter because IL-4 did not block IFNγ-mediated induction of Nos2, and IFNy did not inhibit IL-4-mediated induction of Arg1 and Relma/Fizz1 (fig. S7). Furthermore, IL-4 and IL-13 transactivated the gene 50 N4/N5 promoter (Fig. 3E) (24), and IL-4 antagonized IFNy-mediated suppression of the N4/N5 promoter (Fig. 3F). The effect of IL-4 on the N4/N5 promoter was diminished by mutation of two of four putative Stat-binding sites in the promoter (Fig. 3G and fig. S8). Further, chromatin immunoprecipitation (ChIP) experiments revealed that Stat6 bound to the N4/N5 promoter after IL-4 treatment of MHV68-infected cells (Fig. 3H). Taken together, these data suggest that activated Stat6 induced by IL-4/IL-13 promotes viral replication by binding to and acting on a viral promoter to induce expression of gene 50.

These counterbalancing effects of IFNy and IL-4 on virus replication and viral promoter activity suggest a potential mechanism by which IL-4-inducing pathogens such as helminths promote reactivation. Therefore, we treated mice infected with MHV68-M3-FL virus with a blocking antibody to IFN γ (anti-IFN γ) (clone H22) (25), an isotype control antibody (clone PIP), longlasting IL-4 complexes (IL-4c) (26), or a combination of anti-IFN γ and IL-4c. No reactivation was observed after treatment with anti-IFN γ , IL-4c, or PIP alone, indicating that a single signal was insufficient to reactivate virus in vivo. However, robust reactivation was observed in mice that received a combination of IL-4c and anti-IFN γ (Fig. 4, A and B, and fig. S9A). We next assayed reactivation using an independent assay (27, 28). Little or no preformed virus was detectable in tissues after treatment with PIP, IL-4c, or anti-IFN γ alone (27), whereas treatment with IL-4c plus anti-IFN γ increased infectious virus (fig. S9, B and C). Together, these data support a "twosignal" mechanism by which coinfections could induce reactivation via induction of IL-4 and inhibition of $T_{\rm H}1$ responses (12).

Increased reactivation after treatment with both IL-4c/anti-IFN γ required Stat6 (Fig. 4C). We did not test the role of Stat1 or the IFN γ receptor because both are required to establish latency (6). To assess whether the effects of helminth infection on MHV68 reactivation also required Stat6, we challenged MHV68-infected Stat6KO mice with *H. polygyrus*. We found that helminth infection did not reactivate MHV68 from latency in Stat6KO mice, further supporting a two-signal model for control of γ -herpesvirus reactivation in vivo (Fig. 4D).

Our results suggested a possible role for IL-4 in human γ -herpesvirus reactivation. We therefore tested whether IL-4 could reactivate the human γ -herpesvirus KSHV in the BCBL-1 human

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B cell lymphoma cell line. We found that treatment with IL-4 increased immediate early (RTA, ORF45, and ORF57) and late (ORF19) viral transcripts (29) (Fig. 4E). RTA is the homolog in KSHV of MHV68 gene 50, and ORF45 and ORF57 are both transactivators, indicating a common role of IL-4 in regulating important viral transcriptional transactivators. Furthermore, IL-4 treatment of cells increased virus production (Fig. 4F), indicating that IL-4 is capable of inducing reactivation of KSHV.

A notable aspect of herpesvirus infection is its permanence despite ongoing immunity combined with the capacity to reactivate and spread to new hosts. This work illuminates one potential mechanism by which a γ -herpesvirus exhibits these two apparently disparate functions. Our data suggest that the virus evolved cytokineresponsive promoters to remain latent under some conditions (IFN γ -dominant) and reactivate under other conditions (IL-4-dominant). In this setting, coinfection may govern the outcome of reactivation by changing the balance in IL-4 and IFNy, thus raising a potential issue with herpesvirus reactivation and proposed live helminth therapies (12). Additionally, our data illustrate one potential mechanism by which helminths and other type 2 immune response-inducing parasites influence host control of another pathogen through M2 macrophage polarization (13). The fact that viral promoters for an essential gene are responsive to host cytokines implies that the viral genome evolved to sense the infection status of the host. We speculate that a similar mechanism for IL-4-induced reactivation of KSHV could also be true. Although not extensively studied, seroprevalence to KSHV is associated with hookworm and other parasitic infections in Uganda (30). Intriguingly, certain Burkitt's lymphoma cell lines are reported to express EBV transcripts in response to IL-4 (31).

Although mouse studies are done in specific pathogen-free animals, our data suggest that there is added complexity when multiple pathogens infect the same host, particularly in situations where one pathogen has the capacity to respond to specific immune signals generated to another pathogen to regulate chronic infection. Previously, we showed that herpesvirus infection, a component of the mammalian virome (1), enhances resistance to some pathogens (7). Here, we demonstrate the opposite effect: that coinfection regulates herpesvirus reactivation. These studies emphasize that the spectrum of immunity to chronic infection is a dynamic equilibrium regulated by coinfections, in part through highly evolved pathogen genomes with the capacity to sense host cytokines.

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ACKNOWLEDGMENTS

T.A.R. was supported by a Damon Runyon Postdoctoral Fellowship. This work was supported by grants U54 AI057160 and AI084887 and Crohn's and Colitis Foundation grant 274415 to H.W.V., grants AI032573 and CA164062 to E.J.P., grant CA119917 to R.R., and grant CA52004 to S.H.S. We thank R. Schreiber and K. Sheehan for supplying anti-IFN γ and PIP, G. Randolph and E. Gautier for PPARy f/f x LyzMcre mice and helpful discussion, J. Urban for help setting up the H. polygyrus system, D. Kreamalmeyer for animal care and breeding, members of the Virgin lab for manuscript review and discussion, the Genome Technology Access Center at Washington University for sequencing, and the Flow Cytometry Core at Washington University for assistance with sorting. The data presented in this manuscript are tabulated in the main paper and in the supplementary materials. The GEO accession numbers for RNAseq data on virus-infected cells and for the BMDMs are GSE58116 and GSE21895, respectively, T.A.R. designed the project, performed experiments, and wrote the paper. H.W.V. assisted with project design and paper writing. B.S.W. and S.H.S. defined promoters and performed promoter assays, X.Z. created and characterized MHV68-cre virus and performed limiting dilution PCR and Western blots. H.S.C. and R.R. performed KSHV experiments, M.M.H. and M.H.K. performed ChIP experiments. S.C.H. and E.J.P. helped design helminth experiments and supplied reagents. M.D.B. did RT-PCR for gene 50 and host genes. A.J. did flow cytometry with IL-4 in RAW264.7 cells. A.K. did Western blots. C.Y.L. did plaque assays. G.G. and R.J.X. did initial bioinformatic analysis of gene expression by array, later replaced with RNASeq. P.J.M. provided arginase mice. M.N.A. analyzed sequencing data

SUPPLEMENTARY MATERIALS

www.sciencemag.org/content/345/6196/573/suppl/DC1 Materials and Methods Figs. S1 to S9 Table S1 References (*32*-53)

8 April 2014; accepted 16 June 2014 Published online 26 June 2014; 10.1126/science.1254517



Helminth infection reactivates latent γ -herpesvirus via cytokine competition at a viral promoter

T. A. Reese, B. S. Wakeman, H. S. Choi, M. M. Hufford, S. C. Huang, X. Zhang, M. D. Buck, A. Jezewski, A. Kambal, C. Y. Liu, G. Goel, P. J. Murray, R. J. Xavier, M. H. Kaplan, R. Renne, S. H. Speck, M. N. Artyomov, E. J. Pearce and H. W. Virgin

Science **345** (6196), 573-577. DOI: 10.1126/science.1254517originally published online June 26, 2014

Parasites make it hard to fight viruses

Microbial co-infections challenge the immune system—different pathogens often require different flavors of immune responses for their elimination or containment (see the Perspective by Maizels and Gause). Two teams studied what happens when parasitic worms and viruses infect mice at the same time. Reese *et al.* found that parasite co-infection woke up a dormant virus. Osborne *et al.* found that mice already infected with parasitic worms were worse at fighting off viruses. In both cases, worms skewed the immune response so that the immune cells and the molecules they secreted created an environment favorable for the worm at the expense of antiviral immunity. *Science*, this issue p. 573 and p. 578; see also p. 517

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