

Activation of Host Pattern Recognition Receptors by Viruses

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Summary

Viral recognition by the host innate immune system has become an exciting and growing area of research focus in recent years. It is now apparent that multiple pattern recognition receptor (PRR) families, including Toll-like receptors (TLRs), RIG-I-like receptors (RLRs) and NOD-like receptors (NLRs), contribute significantly to viral detection by sensing viral proteins and nucleic acids, leading to induction of cytokines and type I interferons (IFNs). Of particular current interest is the sensing of viral DNA within infected cells, since the PRRs responsible for this are only partially defined. Recently RNA polymerase III (Pol III) was shown to transcribe some viral DNAs into RNA for detection by RIG-I, leading to IFN induction. Another novel mechanism of viral DNA recognition unveiled, leading to proinflammatory cytokine production, involves the PYHIN family member AIM2.

Introduction

Viral infection of host cells gives rise to IFN β and interleukin-1 β (IL-1 β) production, both of which are essential for host immunity to viruses[1]. IFN β induction requires activation of the transcription factors IFN regulatory factor 3 (IRF3) and NF κ B, while IL-1 β production

involves inflammasome formation[1]. The PRRs involved in eliciting these responses to viral infection had been poorly characterised, however, within the last 2-3 years the knowledge surrounding recognition of viral PAMPS by PRRs in host cells has increased dramatically (see Figure 1). Recent studies have clarified the roles played by TLRs, RLRs and NLRs in recognising specific viruses[2], and have also identified entirely novel receptors for viral nucleic acids. Further, current understanding of host-virus interactions suggests that there are more PRRs for viral nucleic acid yet to be discovered. Here we review these recent developments and highlight some particularly surprising discoveries.

Cell Surface TLRs and Viruses

TLRs are well known to be involved in the detection of multiple pathogens. TLR2 and TLR4 are two cell surface TLRs best known for their role in sensing extracellular pathogens. For example, TLR2 recognises bacterial lipopeptides and various fungal PAMPs [3,4], while TLR4 senses Gram-negative lipopolysaccharide. However recent studies have highlighted the importance of TLR2 & TLR4 in the anti-viral response. Barbalat et al. [5] used cells deficient in TLR2 and TLR9 to show that TLR2 plays a major role in viral recognition in cells derived from the spleen and bone marrow, and that TLR2 activation by an unidentified virus-derived ligand led to production of type I IFN, a classic anti-viral response. The cells responsible for this anti-viral response were identified as inflammatory monocytes, and were shown to be capable of discriminating between bacterial and viral ligands and activating separate sets of genes depending on the stimulus [5].

A novel role for TLR4 in vaccinia virus (VACV) immunity has also been uncovered. Following VACV infection, mice lacking functional TLR4 had greater viral replication, hypothermia and mortality than wild type mice [6]. A novel VACV ligand, rather than an endogenous ligand for TLR4 was invoked here, and it will be of interest both in the case of this and the TLR2 study to see what viral PAMPs are identified for these PRRs in the future.

Endosomal TLRs and viruses

The endosomal TLRs (TLR3, TLR7, TLR8 and TLR9) have traditionally been more strongly linked to anti-viral immunity than the cell surface TLRs, recognising as they do viral nucleic acids, and their roles in sensing viruses are further emphasised by recent studies.

Several studies have highlighted a sensing role for TLR3 in virus infection as a receptor for viral dsRNA (reviewed in [7]). Oddly, and in contrast to TLR4, TLR3 has actually been shown to play a positive role for VACV during infection, since TLR3-deficient mice showed reduced disease morbidity and improved survival relative to wild-type mice following VACV infection [8]. However TLR3 has become one of the few TLRs to be implicated with a protective role in human viral infections: children with herpes simplex virus-1 (HSV-1) encephalitis were found to have a dominant-negative TLR3 allele [9]. Although TLR3 was seen to be redundant in host defense to HSV-1 in epithelial and dendritic cells, expression of TLR3 in the CNS was essential for immunity to HSV-1 [9].

Another recent human study involving TLR3 and relating to Epstein-Barr virus (EBV) has shown that TLR3 can recognise single-stranded non-coding RNA from EBV [10]. The EBV-encoded small RNA (EBER) found in EBV-infected cells can form a stem loop structure which gives rise to dsRNA-like molecules. RNA purified from the sera of patients with chronic active EBV infection (CAEBV) and EBV-associated hemophagocytic lymphohistiocytosis (EBV-HLH) was capable of inducing TLR3 signalling in EBV-transformed lymphocytes and peripheral mononuclear cells. This study highlighted the importance of TLR3 in recognition of EBER, but also the involvement of TLR3 in immunopathologic diseases caused by active EBV infection [10].

TLR7 is known to respond to ssRNA, and another example of a link between TLRs and human disease comes from data showing that responses to HIV-1-encoded TLR7 ligands differ markedly between male and female plasmacytoid dendritic cells[11]. This is now

thought to be one reason for known differences in HIV-1 disease progression between men and women [11]. A further human pathogen that has been added to the list of viruses sensed by TLR7 is West Nile virus (WNV). Two studies using murine models of WNV infection showed increased susceptibility to lethal WNV encephalitis, and an impaired host immune response to the virus [12] [13].

Viral RNA, RLRs and MAVS

The RLRs retinoic acid-inducible gene I (RIG-I) and melanoma differentiation-associated gene 5 (MDA5) are RNA helicases that have previously been identified as critical receptors for the activation of antiviral immunity by viral RNA [14], and recently the exact nature of the RNA ligands recognised by RLRs have been uncovered.

In 2006, two groups uncovered important information regarding the ligand for RIG-I. Pichlmair et al [15] demonstrated that influenza A virus infection, which does not generate dsRNA, can still activate RIG-I. RIG-I was shown to be activated by viral genomic single-stranded RNA (ssRNA) bearing a 5'-triphosphate [15]. Hornung et al [16] simultaneously showed that 5'-triphosphate RNA bound directly to RIG-I. These studies highlighted that uncapped 5'-triphosphate RNA, which is present in viruses detected by RIG-I, but absent in viruses recognised by MDA-5 such as the picornaviruses, is a molecular signature for the detection of viral infection by RIG-I [16]. The exact viral RNA motifs capable of initiating an antiviral response through RIG-I were further characterised by Schmidt et al [17] and Schlee et al [18]. These studies found that a 5'-triphosphate group on RNA was not sufficient to cause antiviral signalling through RIG-I but that RNA structures also need a blunt ended region of dsRNA adjacent to the 5'-triphosphate to initiate signalling [17,18]. Importantly, this explains how RIG-I recognises negative-stranded RNA viruses such as rabies, since such viruses have these blunt end 5'-triphosphate motifs at the end of their ssRNA genomes.

Further, a recent study showed definitively that RIG-I directly recognises viral genomes: using Influenza A and Sendai virus the authors showed that RIG-I agonists are exclusively generated during viral replication and correspond to full-length viral genomes [19].

In the case of MDA5, the ligand had been assumed to be long linear dsRNA, and indeed Pichlmair et al showed that dsRNA extracted from encephalomyocarditis virus- or VACV-infected cells could induce MDA5-dependent Type I IFN [20]. However the stimulatory activity of the RNA resided in higher order structures containing both ss and dsRNA, leading to the conclusion that MDA5 recognises a mesh of web-like RNA web generated during the viral life cycle rather than long dsRNA [20].

Both RIG-I and MDA-5 signal through the adaptor MAVS via caspase recruitment domain (CARD) homotypic interactions [21-24]. As such, MAVS is a critical anti-viral signalling adaptor protein, and in fact has been recently shown for the first time to also mediate a non-RLR anti-viral response: NOD2, a receptor that was previously believed to be restricted to the sensing of bacterial ligands, was shown to bind and respond to single-stranded RNA and activate IRF3, leading to IFN β production [25]. NOD2, which has a CARD, was shown signal via MAVS, and was required in vivo for host defense against RSV [25]. Thus NOD2 has a novel function as an anti-viral PRR.

Viral Sensing, inflammasome activation and IL-1

Many viruses are known to stimulate IL-1 β production, which is important for the host inflammatory response to viruses and also to the induction of fever. After transcriptional upregulation of the *IL1B* gene, IL-1 β production is mediated by caspase 1 which cleaves the precursor inactive protein pro-IL-1 β into the active form. Caspase 1 functions in large complexes termed inflammasomes, the best characterised of which contain NALP3. For

example, NALP3 has recently been shown to be necessary for IL-1 β production in response to adenovirus infection [26] and to RNA derived from influenza [27].

NALP3-independent viral stimulation of IL-1 β production has also recently been shown to occur, leading to the discovery of yet another novel viral PRR, absent in melanoma 2 (AIM2). AIM2 was identified as an IFN-inducible protein that can activate the inflammasome in response to VACV [28] and cytosolic DNA [28-31]. This is consistent with the domain structure of AIM2, which contains a HIN domain capable of binding DNA, and like NALP3, a pyrin domain that is required for inflammasome activation. Expression of AIM2 caused inflammasome formation, and knockdown of endogenous AIM2 siRNA verified its importance for dsDNA-induced inflammasome activation and IL-1 β production in human and murine cells [28-30] [31].

Cytosolic Sensing of Viral DNA for IFN β induction

It is now well known that cytoplasmic DNA sensing pathways exist to detect exogenous DNA, which involve activation of the transcription factor IRF3, via TBK1, leading to induction of IFN β [32], and that such response pathways are critical both for anti-viral immunity, and as triggers of autoimmunity [33]. Although novel downstream signalling components involved in DNA sensing pathways, such as DDX3 [34-36] and STING [37], continue to be identified, and although high mobility group proteins have recently been implicated in delivering nucleic acid to PRRs for detection [38], identification of cytoplasmic DNA PRRs (that could bind DNA and transmit a signal to TBK1) remains elusive.

In 2007, DAI (DNA-dependent activator of IRFs) was implicated as such a PRR [39]. That study showed that overexpression of DAI (DLM-1/ZBP1) in murine fibroblasts enhanced induction of type I IFN and other innate immune genes associated with DNA responses. Silencing of the protein inhibited DNA-induced signalling, and the binding of DAI to dsDNA

induced association of the protein with IRF3 and TBK1 [39]. Further work showed that the DNA binding domains of DAI, and dimerisation were critical for full activation of the pathway [40]. DAI has now been shown to be important for the antiviral immune response to Human cytomegalovirus (HCMV). It was observed that DAI was essential for IRF3 activation and IFN β expression triggered by HCMV, and that overexpression of DAI could inhibit replication of HCMV [36]. From this work, the authors suggested that DAI is the principal PRR for the IFN β response to HCMV infection. However, further work showed that DAI is only responsible for IFN β responses in restricted cells since knockdown of DAI in several cell types had little effect on the IFN β response to exogenous DNA [41]. Importantly, another study showed that adjuvant effect of DNA vaccines in vivo, which occurs through PRR sensing of exogenous DNA, was dependent on TBK1, but independent of DAI [42].

Two groups identified an additional cytoplasmic DNA sensing pathway in 2009, which surprisingly for a DNA response, involved RIG-I and MAVS [43,44]. One type of DNA known to induce IFN β when transfected in the cytoplasm is the B-form synthetic dsDNA poly(dA-dT)*poly(dA-dT) (poly(dA-dT)). This is used as a surrogate synthetic mimic for AT-rich pathogen DNA, in much the same way that Poly(I:C) is used to mimic viral RNA responses. It was shown that Pol III transcribed poly(dA-dT) into an RNA-containing a 5'-triphosphate moiety, which could then be sensed by RIG-I leading to IFN β induction [43,44]. This pathway was also shown to be important in the sensing of EBV EBERs, which were transcribed by RNA polymerase III and then triggered RIG-I activation [43]. Thus EBERs are sensed by both TLR3 (see above) and RIG-I.

Many different types of pathogen DNA, and not just that which is AT-rich, have been shown to induce IFN β through cytoplasmic PRR sensing [33]. Given that the role of DAI is very cell type specific, and that the Pol III-RIG-I pathway only senses AT-rich DNA, additional cytoplasmic DNA PRRs remain to be discovered.

Conclusions and future perspectives

This review has illustrated the often dramatic and surprising discoveries made over the past few years, in terms of how host PRRs sense viruses. As well as heightened understanding of how known viral PRRs function, completely new PRRs have been discovered, and knowledge of the relevance of PRRs for distinct viruses, and in human disease, has increased.

In the near future, we can look forward to gaining further insights into the role of viral PRRs not only in human viral disease, but also in autoimmunity. Importantly, the role of novel PRRs such as AIM2 remains to be determined *in vivo*, through the use of knock-out mice. It will also be important to determine which viruses are sensed by the Pol-III-RIG-I pathway *in vivo*. The identification of cytoplasmic PRRs for non-AT rich pathogen DNA also remains a challenge [33], while another challenge will be to explain the ability of host cytoplasmic PRRs to discriminate between self and non-self DNA, which is now well-defined in the case of RNA.

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Conflict of Interest

The authors declare no conflict of interest.

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- of special interest
- of outstanding interest

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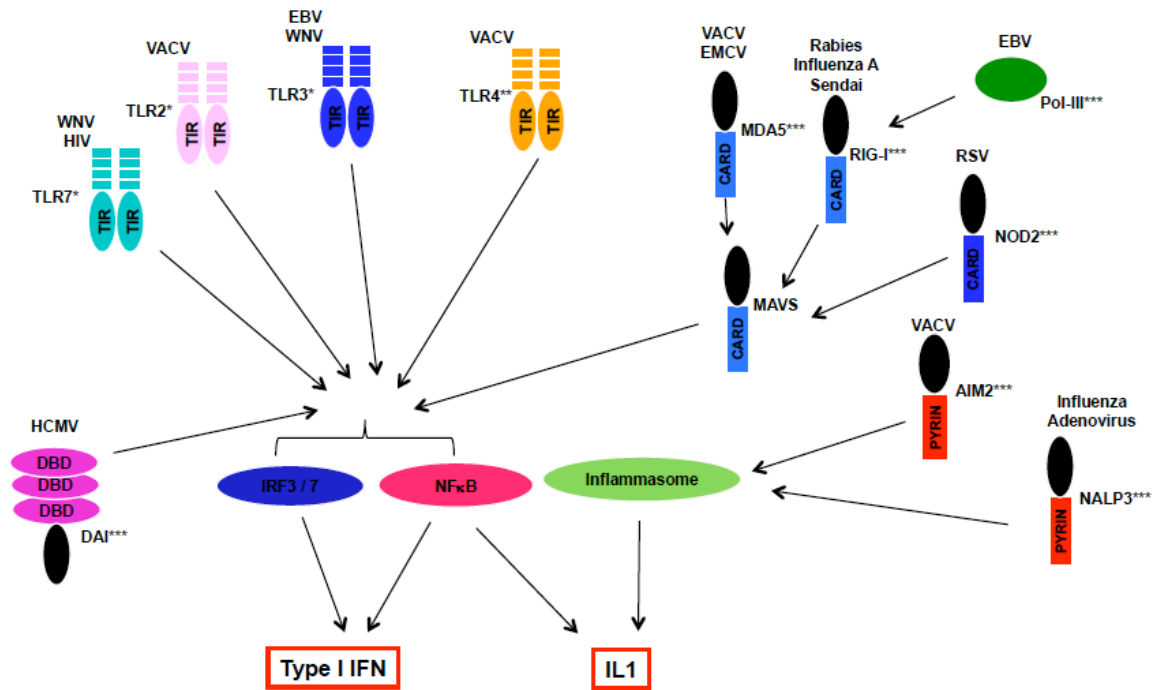


Figure 1. Recognition of viruses by host pattern recognition receptors. Recent advances in understanding the role of TLRs, RLRs and novel viral PRRs in sensing viruses are shown. Arrows indicate signal transduction pathways leading to activation of the transcription factors IRF3 and NFκB, and to the inflammasome, giving rise to type I IFN and IL-1β production respectively. DBD, DNA binding domain; TIR, Toll-interleukin-1 receptor domain; Asterisks indicate the location of each PRR within the cell: *, endosomal; **, cell surface; ***, cytoplasmic.