

A novel anti-viral role for STAT3 in IFN- α signalling responses

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Abstract The cytokine, Interferon (IFN)- α , induces a wide spectrum of anti-viral mediators, via the Janus kinase/signal transducer and activator of transcription (JAK/STAT) pathway. STAT1 and STAT2 are well characterised to upregulate IFN-stimulated gene (ISG) expression; but even though STAT3 is also activated by IFN- α , its role in antiviral ISG induction is unclear. Several viruses, including Hepatitis C and Mumps, reduce cellular STAT3 protein levels, via the promotion of ubiquitin-mediated proteasomal degradation. This viral immune evasion mechanism suggests an undiscovered anti-viral role for STAT3 in IFN- α signalling. To investigate STAT3's functional involvement in this Type I IFN pathway, we first analysed its effect upon the replication of two viruses, Influenza and Vaccinia. Viral plaque assays, using Wild Type (WT) and STAT3-/- Murine Embryonic Fibroblasts (MEFs), revealed that STAT3 is required for the inhibition of Influenza and Vaccinia replication. Furthermore, STAT3 shRNA

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knockdown also enhanced Influenza replication and hindered induction of several, well characterised, anti-viral ISGs: PKR, OAS2, MxB and ISG15; while STAT3 expression had no effect upon induction of a separate ISG group: Viperin, IFI27, CXCL10 and CCL5. These discoveries reveal, for the first time, an anti-viral role for STAT3 in the IFN- α pathway and characterise a requirement for STAT3 in the expression of specific ISGs. These findings also identify STAT3 as a therapeutic target against viral infection and highlight it as an essential pathway component for endogenous and therapeutic IFN- α responsiveness.

Keywords

Janus kinase/signal transducer of activator of transcription (JAK/STAT) \cdot Interferon-alpha (IFN- α) \cdot STAT3 \cdot IFN-stimulated gene (ISG) \cdot Vaccinia virus \cdot Influenza virus \cdot Anti-viral

Abbreviations

AA	Amino acid
ANOVA	Analysis of variance
BCL6	B-cell lymphoma 6
DC	Dendritic cell
EIF2A	Eukaryotic translation initiation factor 2A
EMCV	Encephalomyocarditis virus
EMSA	Electrophoretic mobility shift assay
GAS	Gamma-activated sequence
HCV	Hepatitis C virus
HIF-1α	Hypoxia-inducible factor 1 alpha
HIV	Human immunodeficiency virus
IAV	Influenza A virus
IFN	Interferon
IFN-α	Interferon alpha

IFN-αR	IFN alpha receptor
IL	Interleukin
IRF	IFN regulatory factor
ISG	IFN-Stimulated Gene
ISGF3	IFN-stimulated gene factor 3
ISRE	IFN-stimulated response element
JAK	Janus kinase
MDCK	Madin–Darby canine kidney
MEF	Murine embryonic fibroblast
MHC	Major histocompatibility complex
MMP	Matrix metalloproteinase
MxA/MxB	Myxovirus resistance gene A/B
NK	Natural killer
OAS2	2'-5'-Oligoadenylate synthetase 2
PKR	Protein kinase R
PRR	Pathogen recognition receptor
qRT-PCR	Quantitative real time polymerase chain reaction
RIG-I	Retinoic acid-inducible gene I
RK13	Rabbit kidney 13
RPS15	Ribosomal protein \$15
SH2	Src Homology 2
shRNA	Short hairpin RNA
SIE	Sis-inducible element
SOCS	Suppressor of cytokine signalling
STAT	Signal transducer and activator of
01111	transcription
TNF-α	Tumour necrosis factor alpha
TGF-β	Transforming growth factor beta
Tvk2	Tyrosine kinase 2
VACV	Vaccinia virus
VSV	Vesicular stomatitis virus
WT	Wild type

Introduction

Type I IFNs (IFN-α, -β, -ω, -κ and -ε) are the first line of defence against viral infection and act through autocrine and paracrine pathways to induce an intracellular, anti-viral state [31]. Pathogen recognition receptor (PRR) detection of viral components, including RNA, DNA, protein and glycoprotein, stimulates Type I IFN synthesis. Once secreted by the virally infected cell, Type I IFNs activate the IFN-αR (composed of IFN-αR1 and IFN-αR2 chains), which initiates signalling via the JAK/STAT pathway. Intracellular JAK1 and Tyrosine kinase (Tyk)2 phosphorylate receptor tyrosine residues, which provide docking sites for cytoplasmic STAT proteins [42]. STATs are a family of seven mammalian transcription factor proteins: STAT1, STAT2, STAT3, STAT4, STAT5A, STAT5B and STAT6. STAT5 promote induction of a wide range of genes, encoding proteins with diverse activities, including cell growth, differentiation and immune defence. Receptor bound STATs are themselves phosphorylated by JAKs and subsequently dissociate from the receptor to form homoand hetero-dimers which translocate into the nucleus. STAT1 and STAT2 are well-established mediators of the anti-viral response, which bind IFN regulatory factor (IRF)9, thus forming the IFN-stimulated gene factor (ISGF)3 complex ([14]; [44]). Once ISGF3 translocates to the nucleus, it binds to the IFN-stimulated response element (ISRE) promoter site, which induces expression of several hundred ISGs, many of which are characterised as anti-viral, including protein kinase R (PKR), myxovirus resistance genes (Mx) and 2'-5'-oligoadenylate synthetase (OAS)2 [7, 11, 24, 28]. These genes use a spectrum of anti-viral processes to effectively eliminate infection, including apoptosis, suppression of cell growth and direct inhibition of virus transcription and translation [46]. ISGs are also involved in propagating innate and adaptive immunity via processes such as MHC class I synthesis and induction, which enhances CD8 cytotoxic T cell activity [16]; activation of NK cells, which selectively kills virally infected cells [43]; maturation of DCs, which enhances viral antigen presentation [34] and stimulation of B cell responses [35].

While the role of STAT1 and STAT2 in IFN- α signalling is extensively documented, a role for STAT3 is uncertain. IFN- α phosphorylates STAT3 and promotes the formation of STAT3:STAT3 and STAT1:STAT3 dimers [5]. The role of STAT3 has been historically limited to the transcription of oncogenes including c-Fos [52] and HIF- 1α [40]; pro-inflammatory genes, such as TGF- β [32], IL-6 [50] and TNF- α [9]; cell-cycle genes, including p21 [4] and Bcl-xL [6] and anti-apoptotic genes, including Bcl-2 [10] and Survivin [22]. While STAT3's role in cell proliferation and oncogenesis is clear, its contribution to anti-viral activity remains elusive. Recent studies have shown that STAT3 activity is targeted by Hepatitis C and Mumps viruses [47, 48]. Since both viruses actively deplete STAT3 expression, we hypothesised that this immune evasion strategy reveals an, as-of-yet, undiscovered anti-viral role for STAT3.

Here we show that inhibition of both Vaccinia virus (VACV) and Influenza A virus (IAV) replication is dependent on STAT3. A sub-group of ISGs (PKR, OAS2, MxB and ISG15), all with different anti-viral functions, is dependent on STAT3 for its induction. However, a separate group of ISGs (Viperin, IFI27, CXCL10 and CCL5) are unaffected by STAT3 knockdown. Together, these results reveal an essential, gene-specific anti-viral role for STAT3 in IFN- α signalling, which may be instrumental in developing new therapeutics that target viral infection.

Materials and methods

Cell culture

WT and STAT3–/– MEFs, Huh7 cells and A549 cells were grown in DMEM, supplemented with 10% FCS, 250U/ml penicillin and 250 µg/ml streptomycin. Cells were treated with 1000 IU/ml human IFN- α 2A (Roche) or 1000 IU/ml murine IFN- α (Bio-techne), as appropriate.

Immunoblotting

To prepare whole cell lysates, cells were harvested in RIPA lysis buffer, supplemented with protease and phosphatase inhibitors (1 mM Sodium orthovanadate (Na₃VO₄), 1 mM Phenylmethylsulfonylfluoride (PMSF), 1 mM leupeptin), 25U/ml benzonase and 1 mM DTT. Following 30 min incubation at 4 °C, SDS loading buffer was added and samples were boiled at 95 °C for 10 min. Equal quantities of whole cell lysates were resolved by electrophoresis on a denaturing SDS-polyacrylamide gel according to the method of Laemmli [33] and transferred to a PVDF membrane. Following Immunoblotting using anti-STAT1 (Cell signalling), anti-STAT2 (Santa Cruz), anti-phospho-STAT1 (Cell Signalling), anti-STAT3 (Santa Cruz) and anti- β -actin antibodies (Sigma), the membrane was developed using enhanced chemiluminescent horse radish

Table 1 Human primers used in study

peroxidase (HRP) substrate (BioRad) and analysed using the BioRad Imaging system.

shRNA

Huh7 or A549 cells were transfected in 6 well plates with 3 μ g of either scrambled or STAT3-specific shRNA (InvivoGen), using Lipofectamine 2000 transfection reagent (Life Technologies). Huh7 experiments: after 48 h transfection, cells were incubated in fresh media for 4 h, prior to IFN- α stimulation. A549 cell experiments: After 24 h transfection, cells were infected with IAV for 24 h.

qRT-PCR

RNA was isolated from cells using the TRI Reagent (Sigma) method. qRT-PCR was carried out as described previously [12]. Gene amplifications were normalised to Ribosomal protein S 15 (RPS15) (human) or 18S (murine). Primers used in the study are listed in Table 1 (human) and Table 2 (murine).

Viral infection and plaque assays

VACV Δ E3L (formerly called vP1080) was engineered by in vivo recombination to delete the E3L gene and

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Gene	Forward sequence	Reverse sequence	
PKR	TCTCAGCAGATACATCAGAGATA	AGTATACTTTGTTTCTTTCATGT	
OAS	GAAGCCCTACGAAGAATGTCAGA	TCGGAGTTGCCTCTTAAGACTGT	
MxB	AAGCAGTATCGAGGCAAGGA	TCGTGCTCTGAACAGTTTGG	
ISG15	TCCTGCTGGTGGTGGACAA	TTGTTATTCCTCACCAGGATGCT	
IFI27	GGCAGCCTTGTGGCTACTCT	CCCAGGATGAACTTGGTCAATC	
Viperin	CGTGGAAGAGGACATGACGGAAC	CCGCTCTACCAATCCAGCTTC	
CCL5	CAAGGAGCGGGTGGGGTAGGA	ATCCCCCAAACTGGCTGTCCCG	
CxCL10	GGAAGCACTGCATCGATTTTG	CAGAATCGAAGGCCATCAAGA	
STAT3	GAGAAGGACATCAGCGGTAAGAC	GCTCTCTGGCCGACAATACTTT	
RPS15	CGGACCAAAGCGATCTCTTC	CGCACTGTACAGCTGCATCA	

Table 2 Murine primers used in study

Gene	Forward sequence	Reverse sequence
MxB	GTCGCCTATTCACCAGGCTC	AGCATAACCTTTTGCGAAATTCT
ISG15	TGAGAGCAAGCAGCCAGAAG	ACGGACACCAGGAAATCGTT
Viperin	ATCGCTTCAACGTGGACGAA	GGAAAACCTTCCAGCGCAC
CxCL10	CTGAATCCGGAATCTAAGACCA	GAGGCTCTCTGCTGTCCATC
STAT3	AGATCATGGATGCGACCAACA	CCGTTATTTCCAAACTGCATCA
18S	GTAACCCGTTGAACCCCATT	CCATCCAATCGGTAGTAGCG
GAPDH	GATGCCCCCATGTTTGTGAT	GGTCATGAGCCCTTCCACAAT



Fig. 1 STAT3 is required for anti-viral activity against Vaccinia virus. **a** STAT3 mRNA levels were confirmed in MEFs derived from WT and STAT3–/– ("null") mice by qRT-PCR, before they were infected with vaccinia virus (strain VACV Δ E3L) and **b** treated with IFN- α for 0 and 24 h, after which viral plaque assays using RK13 cells were carried out and **c** quantified. The data shown in Fig. 1C are the mean plaque-forming units/mL \times 10³ obtained from 4 replicates carried out over 3 independent experiments

replace it with a lacZ gene at this locus [3]. MEFs were infected with VACV Δ E3L at an MOI of 0.1. After 1 h at 37 °C, inoculum was aspirated, replaced with 2 ml 2.5% FCS DMEM, treated with IFN- α and harvested after 24 h. Cells were washed with PBS, subjected to three freeze-thaw cycles and sonicated. Viral yields

were quantified by plaque titration on rabbit kidney (RK)13 cells. RK13 cells were infected with serial dilutions of VACV Δ E3L samples from MEFs. After 1.5 h, inoculum was replaced with 1.5% carboxycellulose/MEM (2.5% FCS) overlay for 48 h, before cells were stained with 0.1% crystal violet for 1 h, dried and plaques counted.

MEFs or STAT3 shRNA-transfected A549 cells were infected with IAV A/Puerto Rico/8/1934 (PR8) strain at an MOI of 0.3 and 0.01, respectively. After 1 h at 37 °C, inoculum was aspirated, cells were washed with PBS and overlayed with serum free DMEM with 0.3% BSA (Fraction V, GIBCO), plus TPCK trypsin (Worthington BioChem) and incubated at 37 °C for 24, 48 or 72 h, at which points supernatants were harvested. Supernatants were titrated by plaque assay on MDCK cells and plaque assay plates were incubated at 37 °C for 3 days before staining with crystal violet.

Statistical analysis

Statistical analysis was performed using GraphPad Prism version 5.01 for Windows. Groups were compared by paired or unpaired student's *t* test for parametric samples with Gaussian distributions, and two-way ANOVA with Bonferroni's multiple comparison post-test analysis, as appropriate *p < 0.05, **p < 0.01, ***p < 0.001, ***p < 0.001,

Results

STAT3 is required for anti-viral activity against Vaccinia virus

To determine if STAT3 has a functional anti-viral role, we initially examined VACV replication following IFN-a treatment of WT and STAT3-/- MEFs. VACV is a large dsDNA poxvirus, well known to elicit a robust Type I IFN response [41]. As the VACV-derived E3 protein is a potent inhibitor of the IFN response [23, 38], we used a VACV deletion mutant lacking its expression: VACV- $\Delta E3L$. After confirming the presence and absence of STAT3 mRNA in WT and STAT3-/- MEFs, respectively (Fig. 1a), we infected them with VACV Δ E3L, before treating with IFN- α for 24 h, following which viral titres were quantified by plaque assay titration on RK13 cells. We found that viral plaque formation was considerably increased in RK13 cells incubated with inoculum from STAT3-/- MEFs compared with WT, demonstrating, for the first time, that STAT3 suppresses viral activity (Fig. 1b, c). Furthermore, while treatment with the anti-

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Fig. 2 STAT3 is required for anti-viral activity against Influenza A virus. a WT and STAT3-/- MEF cells were infected with PR8 H1N1 virus. After incubation with Influenza, cells were washed 3 times. overlayed with media and incubated at 37 °C. At 24, 48 and 72 h post infection, samples of the supernatant were collected and titrated by plaque assay on MDCK cells (n = 3). **b** A549 cells were transfected with scrambled control or STAT3-specific shRNA for 24 h before infection with IAV for 24 h. Supernatants were titrated by plaque assay on MDCK cells. At 48 h post transfection c STAT3 mRNA and **d** protein levels were determined by qRT-PCR or immunoblotting, respectively (n = 3)



viral cytokine, IFN- α , reduced VACV replication in WT cells (STAT3 expressing cells), IFN- α had virtually no effect upon the formation of viral plaques without STAT3 expression (STAT3–/– cells) (Fig. 1b, c). These results demonstrate a lack of effective IFN- α -driven responses in the absence of STAT3 and thus reveal a new anti-viral role for STAT3 in IFN- α signalling.

STAT3 is required for anti-viral activity against Influenza A virus

While VACV is a DNA virus and IAV an RNA virus, PRR detection of either induces Type I IFN expression that subsequently mediates anti-viral activity through the JAK/ STAT pathway [2, 49]. Having discovered that STAT3 is **Fig. 3** STAT3 knockdown significantly reduces IFN-α induction of PKR, OAS, MxB and ISG15. Huh7 hepatocytes were transfected with scrambled control or STAT3-specific shRNA for 48 h. After confirming **a** STAT3 protein and **b** mRNA knockdown, cells were stimulated for 2 or 4 h with IFN-α. qRT-PCR was performed to measure **c** PKR, **d** OAS, **e** MxB and **f** ISG15 mRNA induction



required for IFN- α -mediated inhibition of VACV replication (Fig. 1b, c), we subsequently assessed the role of STAT3 in IAV infection, using WT and STAT3–/– MEFs. We found that IAV replicated to significantly higher titres at 24, 48 and 72 h post infection in STAT3–/– MEFs, compared with STAT3-expressing WT MEFs (Fig. 2a). To examine the cell-type specificity of this anti-viral effect, we also infected human alveolar basal epithelial (A549) cells with IAV following shRNA-mediated knockdown of STAT3 and investigated replication of the virus 24 h post infection. Suppression of endogenous STAT3 levels in A549 epithelial cells led to significantly higher titres of IAV (Fig. 2b).

Knockdown of STAT3 mRNA and protein in A549 cells was also confirmed by qRT-PCR (Fig. 2c) and immunoblotting (Fig. 2d), respectively. These observations further implicate STAT3 as an essential anti-viral mediator and demonstrate that its activity has immune regulatory implications for both RNA and DNA viral infection.

STAT3 is specifically required for induction of a subset of IFN-α-driven ISGs

Having established a functional role for STAT3 in blocking viral replication, we next assessed the molecular role for Fig. 4 STAT3 knockdown does not affect IFN- α induction of Viperin, IFI27, CCL5 and CxCL10. Huh7 hepatocytes were transfected with scrambled control or STAT3-specific shRNA for 48 h before stimulation for 2 or 4 h with IFN- α . qRT-PCR was performed to measure **a** Viperin, **b** IFI27, **c** CCL5 and **d** CxCL10 mRNA induction



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STAT3 in anti-viral ISG induction, using the IFN-responsive, human hepatocyte cell line, Huh7. Huh7 cells were transfected with either scrambled or STAT3 shRNA for 48 h, before being stimulated for 2 or 4 h with IFN- α . STAT1, 2 and 3 protein levels were also measured in Huh7 cells following STAT3-shRNA knockdown which confirmed that neither STAT1 nor STAT2 protein levels were altered in the absence of STAT3 (Supplementary Fig. 1). Knockdown of STAT3 protein (Fig. 3a) and mRNA (Fig. 3b) was confirmed by immunoblotting and qRT-PCR, respectively. Next, IFN- α -mediated induction of a group of well-defined anti-viral ISGs (PKR, OAS2, MxB, ISG15, Viperin, IFI27, CCL5 and CxCL10) was analysed by qRT-PCR. We found that induction of PKR, OAS2, ISG15 (4 h IFN- α) and MxB (2 and 4 h IFN- α) was significantly reduced upon STAT3 knockdown, compared to the scrambled control (Fig. 3c-f). Of particular significance was the dramatic suppression of ISG15 induction upon STAT3 shRNA-mediated depletion; in fact, this anti-viral gene had virtually no response to IFN- α in the absence of STAT3 (Fig. 3f). In contrast, qRT-PCR analysis showed that IFN-a-induced expression of Viperin, IFI27, CCL5 and CxCL10 (Fig. 4a-d), was unaffected by STAT3 knockdown, compared to scrambled controls. Having

observed that STAT3 was required for anti-viral activity against Vaccinia (Fig. 1b, c) and Influenza (Fig. 2a) in MEFs, we next analysed if, as in Huh7 cells, STAT3 was required for specific ISG induction in MEFs. WT and STAT3-/- MEFs were treated with IFN- α for 4 h before ISG15, MxB, Viperin and CxCL10 were measured by qRT-PCR. Interestingly, we found that, as in Huh7 cells (Figs. 3e, f, 4a, d), induction of ISG15 and MxB was reduced in the absence of STAT3, whereby Viperin and CxCL10 induction was not (Fig. 5a-d). Importantly, we also confirmed that STAT3 depletion in the MEF cells had no effect on STAT1 expression nor IFN-α-induced STAT1 phosphorylation (Supplementary Fig. 2). Together, these results reveal that induction of a specific ISG subset relies on STAT3 and indicate an essential role for STAT3 in IFN- α signalling that suppresses replication of both RNA and DNA viruses.

Discussion

STAT3 is well known to have essential roles in cell survival, differentiation and proliferation. It is often constitutively active in a wide range of cancers, including **Fig. 5** STAT3-null MEFs display significantly decreased IFN- α induction of ISG15 and MxB, but not CXCL10 or Viperin; MEFs derived from WT and STAT3-/- mice MEFs were stimulated for 4 h with murine IFN- α . qRT-PCR was performed to measure **a** ISG15, **b** MxB, **c** CXCL10 and **d** Viperin mRNA induction (*n* = 3)



leukaemia [20], lymphoma [39], multiple myeloma [25]; along with carcinomas of the breast [18], head and neck [37], ovary [30] and prostate [36]. While this proliferative function for STAT3 is extensively defined, its effect on anti-viral activity remains undecided. Here, we identify a new anti-viral role for STAT3 in IFN- α signalling, by demonstrating that STAT3 is required to block replication of both VACV and IAV and is essential for the optimum induction of a specific subset of anti-viral ISGs.

STAT1 and STAT2 are classically linked to anti-viral responses [29], with viruses such as Respiratory syncytial virus [15], Simian virus 5 (strain W3A) [1], Sendai virus [19] and HCV [47], targeting them for degradation to avoid the host immune response. Our study reveals STAT3 as another essential anti-viral mediator, thus explaining why it is also targeted for degradation by HCV and Mumps [47, 48]. Here, we have shown that loss of STAT3 specifically inhibits a unique group of ISGs (PKR, OAS2, ISG15 and MxB), well characterised to combat viral infection through a broad spectrum of mechanisms, including inhibition of viral mRNA translation (PKR) [17], ISGylation (ISG15) [13], viral RNA degradation (MxB) [21].

Although STAT3 is known to be activated by IFN- α [5], its anti-viral role in IFN- α signalling has remained poorly defined. The primary immunodeficiency disease, Hyper-immunoglobulin(Ig)-E recurrent-infection syndrome (or

Jobs), is caused by STAT3 mutations, that affect its DNAbinding domain and SH2 domain [8, 27]. While "Jobs" is commonly linked to bacterial skin disease and pneumonia, patients also have decreased ability to control Varicella zoster and Epstein-Barr viruses [45], further supporting STAT3's role as an essential anti-viral mediator.

Other groups have previously explored STAT3's antiviral role [26, 51], however, we are first to investigate these effects in human hepatocytes and specifically against VACV and IAV. Ho et al., suggested that STAT3 might sequester STAT1 in the THP-1 monocytic cell line, thus reducing IRF-1, CXCL-9 and CXCL-10 induction [26]. Due to their robust IFN- α responsiveness, we used the human Huh7 hepatoma cell line during our ISG analysis, which may explain the differences observed by Ho et al., in THP-1 cells and thus reveal cell type variation in STAT3's anti-viral activity. While Wang et al., found that knocking out STAT3 in MEFs suppressed titres of Encephalomyocarditis virus (EMCV) and vesicular stomatitis virus (VSV) [51], we found it enhanced VACV and IAV replication, possibly identifying a viral specificity for STAT3's activity.

In summary, our results demonstrate a previously undiscovered anti-viral function for STAT3 in IFN- α signalling, which is crucial for clearance of both VACV and IAV. These results reveal an anti-viral gene-specific role for STAT3, which is instrumental to our overall understanding of innate immunity against viruses. We anticipate these exciting findings to have wider implications against other viruses and be fundamental in the development of new immune regulatory therapeutics.

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