

Inflammatory Cytokine: IL-17A Signaling Pathway in Patients Present with COVID-19 and Current Treatment Strategy

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Abstract: Coronavirus disease 2019 (COVID-19) is a globally communicable public health disease caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). Eradication of COVID-19 appears practically impossible but, therefore, more effective pharmacotherapy is needed. The deteriorated clinical presentation of patients with COVID-19 is mainly associated with hypercytokinemia due to notoriously elevated pro-inflammatory cytokines such as interleukin (IL)-1B, IL-6, IL-8, IL-17, granulocyte-macrophage colony-stimulating factor (GM-CSF), granulocyte colony-stimulating factor (G-CSF), interferon- γ -inducible protein (IP10), monocyte chemoattractant protein (MCP1), and tumor necrosis factor- α (TNF α), and is usually responsible for cytokine release syndrome. In the cytokine storm, up-regulation of T-helper 17 cell cytokine IL-17A, and maybe also IL-17F, is mostly responsible for the immunopathology of COVID-19 and acute respiratory distress syndrome. Herein, I meticulously review the exuberant polarization mechanism of naïve CD4⁺ T cells toward Th17 cells in response to SARS-CoV-2 infection and its associated immunopathological sequelae. I also, propose, for clinical benefit, targeting IL-17A signaling and the synergic inflammatory cytokine IL-6 to manage COVID-19 patients, particularly those presenting with cytokine storm syndrome.

Keywords: IL-17A, inflammation, immunopathology, COVID-19, cytokine storm, Th17, IL-6, ARDS

Introduction

Coronavirus disease 2019 (COVID-19) is a current pandemic infectious disease caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), which first appeared in Wuhan, China, in December 2019, and has since spread globally.^{1–3} To date, official figures released by the World Health Organization (WHO) indicate over 21 million confirmed cases of COVID-19 worldwide, with 761,018 deaths.⁴ This 2019-nCoV is the third and the most lethal pathogenic human positive sense RNA coronavirus identified following the outbreak of zoonotic transmission of CoV that has been recognized as SARS-CoV (in 2003) and MERS-CoV (in 2012), affecting birds and a wide range of animals including humans.^{5–7} The hallmark of each of these infections is viral pneumonia accompanied by host inflammation, leading to pulmonary edema and a syndrome that resembles acute respiratory distress syndrome (ARDS).⁸

Following SARS-CoV-2 infection, patients present with exuberant activation of T cells, and Th17 cell infiltration leads to an elevation of inflammatory cytokines

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such as interleukin-17A (IL-17A) (also known as IL-17) and related families such as IL-17B, IL-17C, IL-17D, IL-17E (also known as IL-25), and IL-17F. In turn, inefficient production of type 1 interferons and an impaired antiviral response are seen, while increasing activation of NF- κ B contributes to the cytokine storm.^{9–12} Secondary to a large amount of inflammatory cell infiltration and the cytokine storm, the alveolar–capillary membrane becomes congested, damaged, and leaky, allowing increased movement of water and proteins from the intravascular space to the interstitial space.¹³ In turn, pulmonary edema, lung failure, and death ensue.⁹ Taking this a step further, severely infected patients with ARDS have elevated serum levels of IL-1B, IL1RA, IL-6, IL-7, IL-8, IL-17, IL-9, IL-10, fibroblast growth factor (FGF), granulocyte–macrophage colony-stimulating factor (GM-CSF), interferon- γ (IFN γ), granulocyte colony-stimulating factor (G-CSF), interferon- γ -inducible protein (IP10), monocyte chemoattractant protein (MCP1), macrophage inflammatory protein-1 α (MIP1 α), platelet-derived growth factor (PDGF), tumor necrosis factor (TNF α), and vascular endothelial growth factor (VEGF) (Table 1).^{9,14–17}

Because there is no specific antiviral therapy for COVID-19, understanding of the cytokine storm mechanisms in this disease could help to reveal possible therapeutic interventions. Therefore, among several inflammatory cells and cytokines, T-helper (Th)-17 cells are a unique subset of a cluster of differentiation 4 (CD4⁺) T-helper

cells characterized by the production of pro-inflammatory cytokines such as IL-17A, IL-17F, and IL-22. Differentiation of naïve CD4⁺ T cells to Th-17 cells is mediated by the activation of T-helper cells in the presence of a combination of TGF- β , IL-6, IL-1 β , and IL-23.^{10,11} In turn, IL-17A acts by activating inducible nitric oxide synthase (iNOS) and inducing the expression of macrophages, IL-1 β , IL-6, IL-8, TNF α , and several chemokines, which collaborate to potentiate the inflammatory process of ARDS.¹³ Elevated Th17 (as well as Th1) responses or enhanced IL-17-related pathways are also observed in MERS-CoV and SARS-CoV patients.^{14,18} ARDS is not commonly due to the viral load but due to an exuberant immune response, and results in cytokine release syndrome (CRS). Thus, there is an urgent need for anti-inflammatory drugs for COVID-19 patients presenting with CRS. This article provides the proposed immunopathological mechanism of IL-17A/F and drugs against IL-17–IL-17R signaling, including the synergistic IL-6–IL-6R axis, as a potential therapeutic option.

Th-17 Cell Response to SARS-CoV-2

Th17 cells are pro-inflammatory and were introduced in 2005 as a third subset of the CD4⁺ T-cell lineage.^{30,31} Functionally, Th17 cells play a role in host defense against extracellular pathogens by virtue of their production of IL-17 and IL-17F, and by mediating the recruitment of neutrophils and macrophages to infected tissues. Moreover, it

Table 1 Epidemiology and Common Inflammatory Cytokines in Patients Presenting with SARS-CoV-2,^{4,14,16,18–24} Compared to SARS-CoV^{18,23,25–27} and MERS-CoV^{12,14,18,25,28}

	SARS-CoV	MERS-CoV	SARS-CoV-2
Epidemiology			
Date of outbreak	November 2002	June 2012	December 2019
Disease	SARS	MERS	COVID-19
Origin	China	Saudi Arabia	Wuhan, China
Region covered	29 countries	28 countries	Worldwide
Confirmed cases	8096	2494	>21 million
Death toll	774	858	>700,000
Inflammatory cytokines			
IL-6	↑	↑	↑
IL-17	Unknown	↑	↑
IL-1	ns	ns	↑
TNF α	↑	↑	↑
MCP1	↑	Unknown	↑
CRP	↑	↑	↑

Abbreviations: SARS-CoV, severe acute respiratory syndrome coronavirus; MERS, Middle East respiratory syndrome; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2; IL, interleukin; TNF α , tumor necrosis factor- α ; MCP1, monocyte chemoattractant protein-1; CRP, C-reactive protein; ns, not significant; ↑, increase.

has become evident that aberrant regulation of Th17 cells may play a significant role in the pathogenesis of multiple inflammatory and autoimmune disorders.³¹ Th-17 cells are promoted by antigen-presenting cells (APCs) through transforming growth factor-beta (TGF- β) in mice, and by IL-6, IL-21, and IL-23 in both mice and humans.^{11,14,31–33} Most people seem to be affected less severely and either remain asymptomatic or develop only mild symptoms during COVID-19.³⁴ There is limited evidence regarding Th-17 cells and the related inflammatory cell profile in asymptomatic people; rather, elevated IL-17 serves as a biomarker of disease severity.³⁵ Moreover, in SARS-CoV-2 infection, APCs such as alveolar macrophages release IL-6, IL-23, and many more cytokines. Taking this a step further, following the binding of IL-6 and IL-23 with their respective receptors,³⁶ key factors in the polarization of naïve CD4⁺ T cells toward differentiation, as well as the maturation of Th17 cells, are signal transducer and activator of transcription-3 (STAT3) and retinoic acid receptor-related orphan receptors gamma (ROR γ) and alpha (ROR α).^{14,36,37} In the lung alveoli, IL-17A, IL-17F, IL-21, and IL-22 are produced as the signature cytokines by Th17 cells in response to polarizing cytokines secondary to presentation of viral infection (Figure 1).^{10,36,38,39} Furthermore, Xu et al showed in a patient with severe COVID-19 that the peripheral blood had a strikingly high number of CCR6⁺ Th17 cells,^{14,40} further supporting

a Th17-type cytokine storm in this disease. Elevated Th17 responses or enhanced IL-17-related pathways are also observed in MERS-CoV and SARS-CoV patients.^{18,41} Inflammatory cytokines continue to be disordered, perhaps leading to lymphocyte apoptosis. Basic research has confirmed that TNF α , IL-6, and other pro-inflammatory cytokines could induce lymphocyte deficiency.^{42,43} Collectively, the Th17-type response contributes to the cytokine storm in pulmonary viral infection, including SARS-CoV-2, which results in tissue damage and likely promotes pulmonary edema; therefore, targeting the Th17 pathway may benefit patients with a Th17-dominant immune profile.⁴⁴

IL-17A Signaling and Pathological Effects During COVID-19

In SARS-CoV, it is immune dysregulation, rather than viral load, which results in aberrant pro-inflammatory cytokine secretion by alveolar macrophages. As a new type of highly contagious disease in humans, the pathophysiology of the unusually high pathogenicity of COVID-19 is not yet completely understood,⁴⁵ but most likely ARDS in the case of SARS-CoV-2 infections is a result of exuberant infiltration of inflammatory cells responding to the SARS-CoV-2 viral infection, and the subsequent synthesis and release of inflammatory cytokines.⁴⁶ Among the many cytokines involved in the storm, IL-17 is a notable and predominant

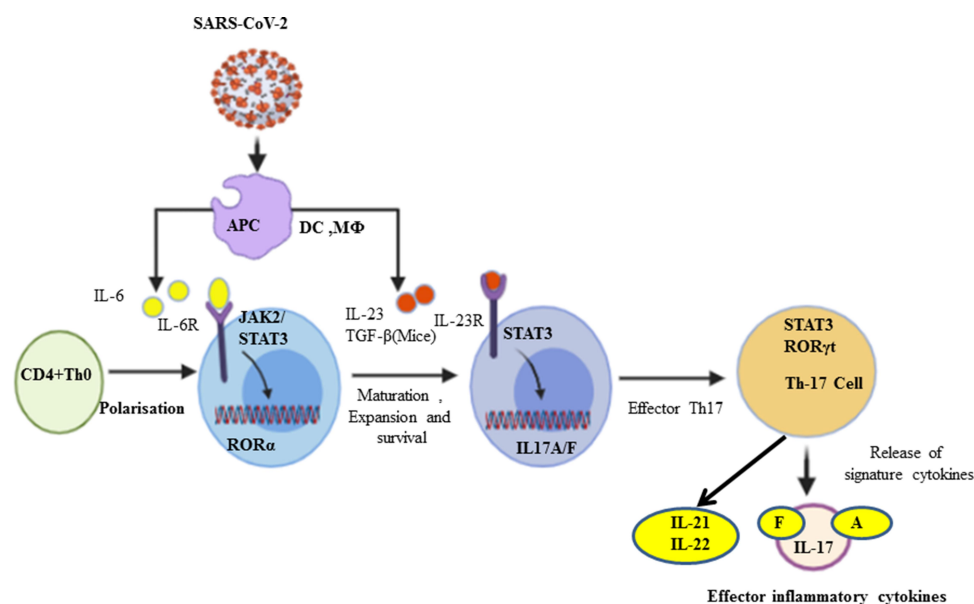


Figure 1 Th-17 cell polarization from naïve CD4⁺ T cells and activation. In response to SARS-CoV-2, antigen-presenting cells (APCs) such as dendritic cells as well as macrophages can present the fragments of the antigen to the naïve CD4⁺ T cells, and upon activation APCs release IL-6, TGF- β , and IL-23 polarizing cytokines. In turn, IL-6 binds with its receptor and through JAK-STAT3 causes polarization, maturation, and expansion of CD4⁺ T cells to Th17 cells with the expression of ROR γ t. In turn, the activated Th-17 cells produce inflammatory cytokines such as IL-17A, IL-17F, IL-21, and IL-22.

mediator of pulmonary inflammation. The pro-inflammatory properties of IL-17 also make it crucial to mediators of inflammation and immunopathology.^{47,48} IL-17 activates many signaling pathways, which in turn leads to the production of many other cytokines (such as IL-6, IL-1 β , TNF α , G-CSF, GM-CSF, and TGF- β) and chemokines (including IL-8 and MCP1) from many alveolar cell types (endothelial cells, epithelial cells, and macrophages).^{44,49} Among the chemokines, profoundly elevated G-CSF and IL-8 (CXCL8) lead to the recruitment of immune cells such as neutrophils to the inflamed area of the alveoli, and hence to dysregulated activation of immune cells, which results in a cytokine storm (Figure 2).^{44,50,51} During IL-17 signaling, IL-17 family cytokines (IL-17A, IL-17F, and others) bind with both IL-17RA and IL-17RC subunits to transduce signaling at the target cells and generate effector

molecules.⁵² Both receptors are engaged in Act1 as an adaptor molecule to recruit TRAF6. In turn, upon recruitment of TRAF6 and its ubiquitination (Act1 may indeed function as an E3 ubiquitin ligase), a cascade of molecular interactions is turned on, leading to the phosphorylation and consequent proteasomal degradation of I κ B, ultimately allowing the nuclear translocation of NF κ B and the activation of NF κ B targeted inflammatory cytokine and chemokine encoding genes.^{52,53} In turn, the gene products of inflammatory cytokines and chemokines are responsible for the cytokine storm.⁴⁹ Moreover, clinical symptoms of patients presenting with COVID-19 result from a consequent significant inflammatory cell infiltrate and release of pro-inflammatory cytokines.⁵⁴ In particular, IL-6, IL-17, and IL-8 are synergistically responsible for pulmonary fibrosis (promoting collagen deposition) secondary to

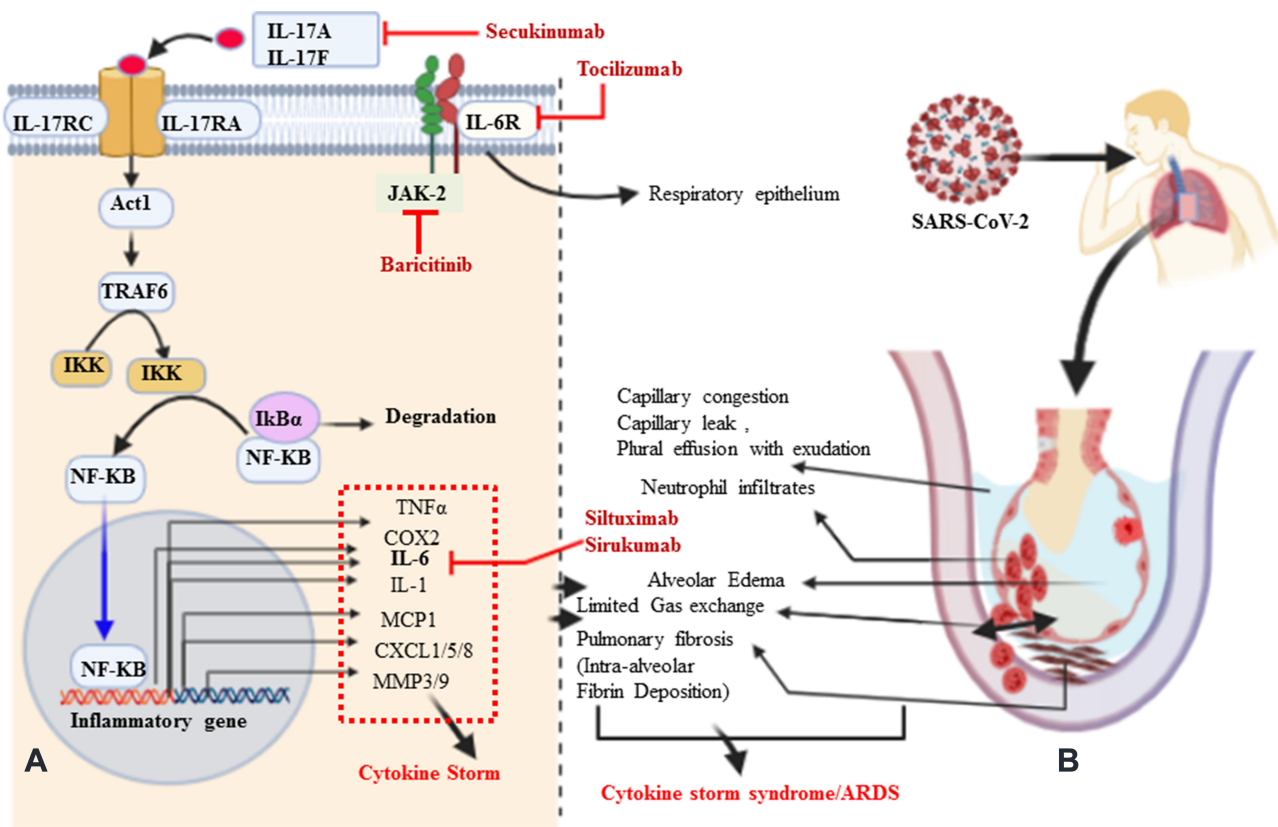


Figure 2 Signaling pathways of IL-17A, pathological effect, and potential drug targets. **(A)** Signaling transduction of IL-17A at type I epithelial cells of the alveoli of the lung. IL-17A and IL-17F are secreted predominantly by Th-17 cells; they are structurally very similar, bind the IL-17RA–IL-17RC receptor combination, and can form heterodimers together, signaling via the adaptor protein nuclear factor (NF)- κ activator (Act1). Many IL-17 target genes contain a promoter region that binds with NF- κ B. IL-17 is not a potent inducer of inflammation by itself. Its strong effects during inflammation are derived from its ability to recruit immune cells via chemokine expression such as CXCL1, CXCL5, and MCP1, as well as from its synergistic action with other cytokines such as IL-6, IL-1, and TNF α . Thus, IL-17, acting in synergy with IL-6 and TNF α , is a powerful inflammatory signal that results in the rapid recruitment and sustained presence of neutrophils and leads to a cytokine storm. **(B)** Schematic representation of IL-17A-mediated immunopathological effect of ARDS during SARS-CoV-2 infection. Pulmonary fibrosis is one of the pathological changes due to the activation of fibroblasts mediated by IL-6 and results in abnormal deposition of collagen. Moreover, the stimulation of fibroblasts can produce IL-8 and leads to the attraction of neutrophils to the site of injury. MMP3/9 (which causes tissue destruction) and PGE2 (increases capillary permeability) are also responsible for neutrophil infiltration, alveolar edema, and protein-positive (exudative) pleural effusion. Altogether, the proposed pathological mechanism suggests that IL-17 can mediate numerous immunopathological effects in CRS secondary to SARS-CoV-2 infection.

aberrant fibroblast and epithelial cell function or pleural effusion of the lung, and viral persistence results in dyspnea and provokes SARS (Figure 2).⁵⁵ I suggest that the presence of these cells may be a primary driver of the signature pathology observed in COVID-19 patients. Taken together, IL-17-induced dysregulated and exuberant immune responses have been shown to potentially cause stage 3 (characterized by a hyperinflammatory phase, cytokine storm) COVID-19 disease, likely through increased pulmonary pathology or lung damage and diminished survival.^{25,56}

IL-17A Targeting as a Treatment Strategy Against COVID-19

COVID-19 is a global public health problem.⁴ Currently, many different treatment approaches are under investigation and clinical trials are ongoing. There are currently no known specific treatment measures except for meticulous supportive care such as mechanical ventilation when indicated.⁵⁷ Taking the pathogenesis of this novel coronavirus from the experience of the first SARS-CoV, it is clear that the exuberant immune response and inflammatory cell infiltration are the pathological mechanisms most responsible for the catastrophic death toll.⁵⁸ Among the anti-cytokines, monoclonal antibodies are a potential and possible approach to the prevention of SARS, although none is yet available for clinical purposes. Anti-IL-17A humanized monoclonal antibodies such as secukinumab (AIN457) (approved fully human IL-17-specific IgG1k monoclonal

antibody) and Ixekizumab (LY2439821) (under a phase III clinical trial for psoriasis) are well-known and recognized treatment options for psoriasis.^{44,59–61} These anti-IL-17A monoclonal antibodies are also utilized in different inflammatory diseases, including ARDS,¹³ rheumatism,⁶¹ and pulmonary fibrosis.⁶² Moreover, patients presenting with COVID-19 show elevated serum IL-6, which correlates with respiratory failure.⁶³ As discussed earlier, IL-6 plays a key role not only in the polarization of Th17 cell from naïve CD4⁺ T cells but also in promoting pulmonary inflammation in synergy with IL-17A. Indirectly, tocilizumab is now approved by the US Food and Drug Administration (FDA) for the treatment of chimeric antigen receptor T cell (also known as CAR T cell)-induced CRS, with confirmed efficacy and minimal side effects in hundreds of patients.⁶³ Taking this a step further, tocilizumab against IL-6 has been shown to be efficient for patients presenting with inflammatory rheumatism (Figure 2).^{61,64,65} In addition, IL-23 and IL-6 are involved in Th17 differentiation, and they act through the JAK-STAT3 signaling pathway. This implies that STAT3 could be a potential target, at the convergence point of different upstream activators. On the other hand, the anti-JAK tofacitinib^{56,65} (oral inhibitor, selective for JAK1 and JAK3), is under a phase III clinical trial for the treatment of inflammatory rheumatic disorders as well as COVID-19.⁶⁶ JAK inhibitors interfere with phosphorylation of signal transducer and activator of transcription (STAT) proteins and the production of downstream inflammatory molecules. Thus, anti-IL-17 (secukinumab)

Table 2 Summary of Clinical Trials Investigating the Implications of Anti-Inflammatory Agents in COVID-19 Patients, Registered Under Clinicaltrials.gov, WHO Trial Registry Network, and NIH

Drugs	NCT Number	Clinical Trial Status	Sample (N)	Study Area	Outcome
Secukinumab	NCT04403243	Recruiting (P-II)	N=70	Russia	Time-frame: baseline, day 12, using 7 domains of laboratory parameters
Tocilizumab	NCT04320615	Recruiting (P-III)	N=300	USA	Clinical status assessed using a 7-category ordinal scale
Baricitinib	NCT04390464	Recruiting (P-IV)	N=1167	UK	Time to the incidence of the composite endpoint of: death, mechanical ventilation, cardiovascular organ support, or renal failure
Sirukumab	NCT04380961	Recruiting (P-II)	N=270	Belgium	Time-frame: up to day 28, using 6-category ordinal scale
Siltuximab	NCT04330638	Recruiting (P-III)	N=342	Belgium	ND
	NCT04322188	Completed	N=220	Italy	ND
	NCT04329650	Recruiting (P-II)	N=200	Spain	Proportion of patients requiring ICU admission

Abbreviations: COVID-19, disease related to SARS-CoV-2 infection; P, phase; N, number of COVID-19-positive study participants; ND, not defined; ICU, intensive care unit.

combined with blockade of IL-6 (sirukumab and siltuximab), IL-6R (tocilizumab), or JAK (baricitinib, selective for JAK1 and JAK2) may be beneficial in controlling the cytokine storm while boosting antiviral IFN-I responses during SARS-CoV-2 infection (Table 2).^{57,66}

Conclusion

In this review, I have discussed the current understanding of IL-17A signaling and the underlying immunopathological role as a pro-inflammatory cytokine during ARDS secondary to SARS-CoV-2 infection. Profound elevation of IL-17A and downstream synergetic effector molecules such as cytokines (IL-6, IL-1 β , TNF α) and chemoattractants (IL-8 and MCP1) are indicative of both pathological progression and the severity of COVID-19. The synergistic interaction of IL-17A and IL-6 (required during polarization of Th17 cell) is the central player in the development of pulmonary fibrosis and an impaired respiratory system. Regarding therapeutic strategies, specific therapies for SARS-CoV-2 infections and related complications remain challenging. Although there is limited research evidence, here it is proposed that anti-IL-17A (secukinumab), anti-IL-6, or anti-IL-6R antibody (tocilizumab), or an anti-JAK-2-STAT3 drug (baricitinib) as a promising therapeutic option to terminate pulmonary inflammation, could decrease dangerous cytokines and limit lung tissue pathology, and help to save life if used judiciously in appropriate COVID-19 cases.

Abbreviations

ARDS, acute respiratory distress syndrome; COVID-19, coronavirus disease 2019; CRS, cytokine release syndrome; IL-17, interleukin-17; JAK, janus kinase; ROR γ / α , retinoic acid receptor-related orphan receptors gamma and alpha; SARS, severe acute respiratory syndrome; STAT3, signal transducer and activator of transcription 3.

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Author Contributions

The author made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to

which the article has been submitted; and agree to be accountable for all aspects of the work.

Disclosure

The author reports no conflicts of interest in this work.

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