

A review on COVID-19 effective pharmaceuticals considering their molecular targets and Single nucleotide polymorphisms effects

zahra saadatian¹, Saba Seyedi¹, Mohammadreza Gerami¹, Shadan Navid¹, Samira Ezi¹, Ziba Nariman-Saleh-Fam¹, Lida Nariman-Saleh-Fam¹, and Zahra Saadatian¹

¹Affiliation not available

June 27, 2023

A review on COVID-19 effective pharmaceuticals considering their molecular targets and
Single nucleotide polymorphisms effects

Saba Seyedi^a, Mohammadreza Gerami^b, Shadan Navid^c, Samira Ezi^d, Ziba Nariman-Saleh-Fam^e, lida Nariman-Saleh-Fam^f, Zahra Saadatian^{g*}

- a) Department of Medical laboratory sciences, School of Medicine, Gonabad University of Medical Sciences, Gonabad, Iran.
- b) Department of architecture, Eqbal lahoori institute of higher education, Mashhad, Iran.
- c) Department of Anatomy, Faculty of Medicine, Social Determinants of Health Research Center, Gonabad University of Medical Sciences, Gonabad, Iran.
- d) Department of Anatomy, Faculty of Medicine, Gonabad University of Medical Sciences, Gonabad, Iran.
- e) Women's Reproductive Health Research Center, Tabriz University of Medical Sciences, Tabriz, Iran.
- f) Faculty of Medicine, Ardabil University of Medical Sciences, Ardabil, Iran.
- g) Department of Physiology, Faculty of Medicine, Infectious Diseases Research Center, Gonabad University of Medical Sciences, Gonabad, Iran.

Corresponding author: Zahra Saadatian. Department of Physiology, School of Medicine, Gonabad University of Medical Sciences, Gonabad, Iran.

z.saadatian@yahoo.com saadatian.z@gmu.ac.ir

A review on COVID-19 effective pharmaceuticals considering their molecular targets and Single nucleotide polymorphisms effects

Abstract

COVID-19 is a highly contagious viral disorder which declared a global pandemic and results in more than 6 million mortalities worldwide since the late December 2019. Considering it continues to be a major health problem, finding the best medicines to concur with the effects of COVID-19 is essential and various drugs are suggested and used in clinical trials against covid-19. This review article provided an overview of the potential therapeutics in the management of COVID-19 based on the current disseminated scientific documents. After categorizing pharmaceuticals in to the anti-interleukin drugs, Antiviral factors, Monoclonal antibodies, Corticosteroids and Anticoagulant drugs we presented a comprehensive description of their molecular mechanisms and clearly demonstrated the function on the target cells in COVID-19 virus infection. Moreover, we reviewed the single nucleotide polymorphisms located at direct target of these drugs and may be interfered with their functions. Our intention in this review attempted to supply beneficial therapeutic drugs to treat COVID-19 patients. We hope that this Review shed light on the field of current COVID-19 research, raise awareness and help researchers to select the best treatment protocols against COVID-19.

Keywords: COVID-19, treatment, anti-interleukin, Antiviral, drugs, SNP

Introduction

The Coronavirus disorder 2019 (COVID-19), is investigated as a superlative tension global pandemic that is derived from severe acute respiratory syndrome coronavirus-2 (SARS-CoV-2) (1). Coronaviruses are an assortment of RNA viruses that contaminate humans, and even also animals (2). based on their morphology as round virions with a core-shell and surface layouts that resemble a solar corona, they were denominated coronaviruses (Latin: corona = crown) (3). They are classified in to four subfamilies, including alpha-, beta-, gamma-, and delta-coronaviruses. Pneumonia, asymptomatic infections and gastrointestinal manifestations were identified as primary clinical signs of COVID-19 disease (4). SARS-CoV-2 targets lung alveolar epithelial cells by receptor-mediated endocytosis utilizing the angiotensin-converting enzyme II (ACE2) as an entrance receptor (5). Common symptoms of this disease include cough, fever, dyspnea, musculoskeletal symptoms, gastrointestinal symptoms, and anosmia/dysgeusia (6).

The World Health Organization (WHO) states there is currently no specific treatment for coronavirus. furthermore, considering standard care, treatment of covid 19, just be carried out in approved, randomized, controlled trials (7). Threatening agents for patients infected with COVID-19 is underlying diseases such as cardiovascular disease, chronic lung disease, diabetes, and older age and obesity (8, 9). Management of COVID-19 relies on severity of the condition (8). mild COVID-19 typically cause common cold symptoms. cough with or without hyposmia and sputum are it's the most typical symptoms (8, 10). Homecare, rest and consumption of enough water and adequate calorie, usually ameliorate Patients with mild symptoms (11). However moderate or severe form of the disease needs intensive care and sometimes hospitalization(8). probably antiviral medications like remdesivir and antibody-based treatments are most impressive when prescribed early (12). anti-inflammatory medications, immunomodulators, and anticoagulants are the other effective medications (8, 13).

Single nucleotide polymorphisms (SNPs) are the most common variations in genome (14). They can be used to anticipate how an individual might react to specific medications, their vulnerability to environmental hazards like toxins, and their likelihood of developing certain illnesses(14).

Therefore, single nucleotide variations in the drugs target genes could affect their impression and maybe it would be important to take them in to consideration in the treatment process of COVID-19. In this review, we summarize effective pharmacological treatments for COVID-19 considering their molecular targets and effective single nucleotide polymorphisms.

1. anti-interleukin drugs:

cytokines are some mini proteins that are playing a major role in cell signaling pathways (15). chemokines, interferons, interleukins, lymphokines, and tumor necrosis factors, are included in the cytokine family that is produced by a wide range of cells, involving immune cells such as macrophages, T lymphocytes, B lymphocytes, and mast cells as well as fibroblasts, endothelial cells, and various stromal cells (16, 17). Cytokine storm is associated with the expansion of unrestrained systemic inflammation and further hyperproduction of cytokines (18). Regulation of immune responses caused by cytokine release is disrupted and causes epithelial cell apoptosis and endothelial and vascular leakage, and reduction of the T cell responses which results in inefficiency and hyperactivity of virus-infected macrophages and disturbance of tissue homeostasis followed by pathogenesis and severity of macrophage activation storm (MAS) resulting in MODS (multiple organ dysfunction syndromes) and ARDS(Acute respiratory distress syndrome) (19).

SARSCoV-2 infection motivates hyperinflammation of the innate and adaptive immune systems, which results in a cytokine storm. hypercytokinemia, or “Cytokine Storm (CS)” identified via a severe hyperinflammatory immune response is the principal symptom of covid-19. It is initiated by activation of macrophages, T-cells, and discharge of other cytokines leading to activation and recruitment of other cells involved in the immune system. The increase in the serum levels of pro-inflammatory cytokines and chemokines mainly IL-6, IL-1, IL-12, IL-17, TNF- α , IFN- γ is characteristic of the cytokine storm (20). As well as cytokines raised serum levels of C-reactive protein, lactate dehydrogenase, procalcitonin, creatinine, d-dimer, ferritin, and White Blood Cell count are serious parameters for the prediction of respiratory failure in COVID-19 patients (21). In the following paragraphs we will explain more about Interleukin 1, Interleukin 6, Interleukin 17 and Interleukin 23 proteins.

a) Interleukin 1:

Increased levels of IL-1 that is released in viral diseases cause inflammation of the lung tissue, fibrosis and fever (19). Excessive expression of interleukin 1 by activating some factors such as nuclear and transcription factors, activator protein 1, and activating factor 2 develops viral disease (22). It stimulates pro-interleukin-1 and the regulative cells in the innate and adaptive immune systems producing certain immune responses (19). As a result, interleukin 1B is produced and leads to lung damage and respiratory complications in the host infected with the virus

(19). IL-1a and IL-1b are mediators of inflammatory responses to tissue damage. They are secreted from damaged epithelial and endothelial cells and permeate to macrophages, neutrophils, and monocytes (23). IL-1 receptor antagonist is the basic innate regulatory mechanism that prevents inordinate Inflammation caused by IL-1(24).

A recombinant, intravenous drug, named Anakinra is a non-glycosylated form of the human interleukin-1 receptor antagonist which is expressed in Escherichia coli expression system (25). As the principal advantage, short half-life of this medicine permits quick disposal from circulation. Anakinra previously approved by the US Food and Drug Administration is available as a safe treatment to decrease adverse inflammation in patients suffering from COVID-19 (19). Anakinra as an interleukin IL-1 receptor antagonist inhibits activation of pro-inflammatory cytokines such as IL-1 α and IL-1 β and reduce hyperinflammation (19) (Figure 1). Studies have shown that early treatment with this drug attuned to the soluble plasminogen urokinase activator receptor (suPAR) reduces respiratory failure and amends the inflammatory balance (26).

b) Interleukin 6:

IL-6 is a crucial pro-inflammatory cytokine and is involved in activating Janus kinase (JAK) signal by binding the transmembrane (cis-signaling) or soluble form (trans-signaling) of the IL-6 receptor and linking with membrane-bound gp130 (21). it is involved in two signaling pathways called Classical-signaling and trans-signaling. In the classical signaling pathway with the production of C-reactive protein (CRP), it plays an essential role in the acute immune response against pathogens. while the trans-signaling pathway is involved in long-term inflammation (21). Excessive IL-6 signaling causes numerous effects that leads to organ damage, such as transforming naive T cells into efficacy T cells, inducing vascular endothelial growth factor (VEGF) expression in epithelial cells, expanding vessel permeability, and diminishing myocardial contractility (27).

After Coronavirus infection, a cytokine storm occurs that triggers the release of inflammatory cytokines such as IL-6, Tumor Necrosis Factor- α (TNF $-\alpha$), and IL-12 (20) . most probably IL-6 has a major part in a cytokine storm, thus the drugs which their object is the IL-6 receptor suggested for severe disease COVID-19 patients (28). Tocilizumab is a recombinant humanized monoclonal anti-IL-6R antibody. It is prescribed intravenously and attaches to both soluble and membrane-bound IL-6 receptors to suppress IL-6 cis-signaling and trans-signaling (27) (Figure 2). Infection reduction, diminish of fever and a subtracted

demand for supportive oxygen is appeared a few days after receiving tocilizumab (27).

c) Interleukin 17 and Interleukin 23:

Interleukin 17 is one of the members of cytokine storm produced by Th17, Tc17, and other lymphoid cells (29). Interleukin 17 is mainly secreted from T helper 17 cells in response to any viral respiratory infection such as COVID-19 in the lungs (30). The first function of interleukin 17 is the initiation of neutrophil penetration into infected tissues and the tissue response prompt to extracellular pathogens, and the second function is indirect, such as the induction of chemokines (31). Interleukin-17 stimulates the production of cytokines, chemokines, other inflammatory mediators, matrix metalloproteinases, and growth factors (30). Interleukin 17 can directly activate fibroblasts and indirectly increase viral mediators in the inflammatory process. In addition, it can stimulate fibrogenesis by activating pro-coagulation pathways. So its inhibitors can be used for acute stages of COVID-19 and are also useful to prevent long-term fibrotic consequences (31). There are three ways to reduce the effects of interleukin 17, including blocking interleukin 17, its receptor and its pathway (32). (Figure 3).

Netakimab, is a monoclonal antibody against interleukin 17-A that is used in diseases such as moderate-to-severe plaque psoriasis, ankylosing spondylitis, and psoriatic arthritis (29). Moreover, it can be a possible target for COVID-19 therapy, and reduction of the inflammatory response (29, 33). Secukinumab may suppress T-helper 17 cytokine storm (33) and Brodalumab hinders IL-17R (29). (Figure 3).

similar to interleukin 17, interleukin 23 is a major cytokine in the maintenance of helper T cells, which mediates defense mechanisms against pathogens. Interleukin 23 inhibitors may play an important role in attenuating important cytokines through their effect on T helper 17 (34). Risankizumab, a humanized IgG monoclonal antibody binds with high affinity to the p19 and inhibits IL-23 (34) (Figure 3).

2. Janus kinase inhibitor

The Janus kinase (JAK) family plays a critical role in the immune system's response to viral infections such as COVID-19 (35). Upon viral infection, the host immune system activates cytokines and other signaling molecules to initiate an immune response (35). The JAK family of proteins is involved in the downstream signaling of these cytokines, transmitting signals from the cell surface to the nucleus, where they activate transcription factors such as STAT (Signal Transducers and Activators of Transcription) proteins (36). Thus, the Janus

kinase family, specifically JAK1 and JAK2, have emerged as potential therapeutic targets for COVID-19, and JAK inhibitors have shown promising results in reducing inflammation and improving outcomes in COVID-19 patients.

a) Baricitinib

Baricitinib is a small molecule inhibitor that selectively targets the Janus kinase family, with specificity towards JAK1 and JAK2 (37, 38). The inhibition of Janus kinases by Baricitinib results in the prevention of downstream phosphorylation and activation of Signal Transducers and Activators of Transcription (STAT) proteins (39). Consequently, JAK inhibitors such as Baricitinib can modify the signaling pathways of various interleukins, interferons, and growth factors (36). In addition to its anti-inflammatory profile, Baricitinib exhibits antiviral effects by blocking the entry of SARS-CoV-2 into lung cells by reducing the endocytosis of SARS-CoV-2 through the inhibition of AP2-associated protein kinase 1 and cyclin G associated kinase (39, 40) (41) (Figure 4). The recommended dosage of Baricitinib is 4 mg once daily for up to 14 days. Baricitinib has been suggested as a potential therapeutic option for COVID-19 due to its significant immunosuppressive and antiviral properties (39).

b) Tofacitinib

Tofacitinib is a Janus kinase inhibitor, exhibiting partial selectivity towards Janus kinase 2, and possesses immunomodulatory and anti-inflammatory properties (42). Tofacitinib functions by binding to Janus kinases, which inhibits the activation of the JAK-STAT signaling pathway, leading to a potential reduction in the production of pro-inflammatory cytokines (42) (Figure 5). Tofacitinib is administered orally at a dosage of 10 mg twice daily. Its effectiveness has been observed in Covid-19 patients admitted to hospitals (43) (44, 45).

3. Antiviral factors (RNA-dependent RNA polymerase inhibitor)

a) Remdesivir:

In the class of RNA-dependent RNA polymerase inhibitor, Remdesivir is classified as a prodrug of a monophosphate nucleoside analog and it was developed for the treatment and antiviral function against some RNA viruses including coronaviruses (SARS-CoV, MERS-Co-V, SARS-CoV-2), filoviruses (Ebola viruses, Marburg virus), paramyxoviruses (parainfluenza type III virus, Nipah virus, Hendra virus, measles, and mumps virus), and Pneumovirus (respiratory syncytial virus) (46).

In theory, nucleoside analogs permeate through the cell wall barely. On their next entry into the host cell, they should undergo phosphorylation to produce nucleoside triphosphate (NTP), this is similar to adenosine triphosphate (ATP) and can be adopted in genome replication by the RNA-dependent RNA polymerase (RdRp) enzymes (46). After Remdesivir metabolization into the pharmacologic active analog adenosine triphosphate it participates in a competition with ATP and disrupts RNA synthesis (46) (Figure 6). Remdesivir constrains viral replication in human airway epithelial cell culture by interrupting the first stages of viral replication (47). World Health Organization(WHO), Food and Drug Administration(FDA) and the Infectious Disease Society of America(IDSA), approved and recommended this drug (47). However, like any other antiviral medicine, there are some concerns about the stability of mutant viruses (46).

Remdesivir is prescribed for patients with severe forms of the disease suffered from respiratory failure aged ≥ 12 years, with a body weight ≥ 40 kg (48) and can decrease time of hospitalization (46). The recommended treatment period is between 5-10 days (46), by Intravenous injection of 200 mg in the first day and 100 mg daily for 9 days (total 10 days of treatment) (48).

The metabolism of remdesivir is by cytochrome P450 (CYP450), thus possibly has drug-drug interaction (46). Remdesivir could disturb Cardiovascular, Pulmonary, Hematological, Endocrine, Gastrointestinal, Neurological, skin, Renal and Metabolic normal functions (46).

b) Favipiravir:

Favipiravir as an oral RNA-dependent RNA polymerase inhibitor is a purine nucleotide, or a guanine analog (47). It was used against RNA viral diseases like influenza and Ebola viruses, but it is now used for other RNA viruses like coronaviruses (47). Favipiravir is a purine base analog and intracellular phosphoribosylation converts it to its active form, ribofuranosyl-5B-triphosphate (favipiravir-RTP) (49). Favipiravir with high affinity to bind RNA-dependent RNA polymerase, inhibits RNA-dependent RNA polymerase (RdRp) of RNA viruses intently, which results in ending the chain and viral mutagenesis. favipiravir-RTP integrated with viral genome, along with mutagenesis leads to subtraction of RNA virus (49) (Figure 7).

Favipiravir is prescribed orally (47) in different dosages for patients with mild to moderate hepatic disorder(at the first day 1200 mg twice daily; Days 2-4: 800 mg twice a day) and patients suffering from severe hepatic disease at first day 800 mg twice daily; Days 2 to 3: 400 mg twice a day) (49). There is a possible

drug interaction between Favipiravir and drugs that inhibit aldehyde oxidase (49). The common side effects of it includes gastrointestinal troubling, increased uric acid, Neutropenia, Anemia, diarrhea, increase of aspartate aminotransferase (AST) and alanine transaminase (ALT), psychiatric symptom reactions, and excess triglycerides in the blood (49). It has contraindicated in pregnant women and lactating women (49).

c) Nirmatrelvir

Nirmatrelvir is a selective inhibitor of RNA-dependent RNA polymerase that specifically targets the 3C-like protease enzyme required for COVID-19 viral replication (50-52). This inhibition results in the prevention of virus replication. However, due to its short half-life, nirmatrelvir is co-administered with ritonavir, which is a potent inhibitor of cytochrome P450 (CYP) 3A4, leading to the inhibition of nirmatrelvir metabolism and an increase in its plasma concentration, thereby enhancing its pharmacokinetic profile (50, 53) (Figure 8). Ritonavir has no direct effect on SARS-CoV-2 (53). The recommended dosage for nirmatrelvir/ritonavir is two 150 mg tablets of nirmatrelvir and one 100 mg tablet of ritonavir taken together twice daily for five days (53). The FDA has granted emergency use authorization for nirmatrelvir/ritonavir for the treatment of COVID-19 patients (51).

d) Molnupiravir

Molnupiravir is a prodrug that is a ribonucleoside small molecule derivative known as β -D-N4-hydroxy cytidine (NHC) (54, 55). It has been shown to possess antiviral activity against RNA viruses (54). NHC circulates in the body and undergoes intracellular phosphorylation to form NHC triphosphate (54). Subsequently, viral RNA polymerase combines with NHC triphosphate to misdirect the virus to bind to either guanosine or adenosine when replicating (54). This results in the accumulation of destructive mistakes in the viral genome, ultimately rendering the virus non-infectious and unable to replicate (54) (Figure 9). Molnupiravir is administered at doses of 600 and 800 mg twice daily for five days (54). The FDA has granted emergency use authorization for Molnupiravir in the treatment of COVID-19 (56).

4. Corticosteroids:

Corticosteroids have two subtypes called glucocorticoids and mineralocorticoids. Glucocorticoids have anti-inflammatory effects and mostly are involved in attenuating immune responses (57). Glucocorticoids based medicines are among the most widely used drugs and are used for various diseases such as autoimmune diseases and allergies. These drugs include betamethasone, dexamethasone,

hydrocortisone, triamcinolone, methylprednisolone, prednisone, clobetasol, beclomethasone, fludrocortisone, fluocinolone (57). Corticosteroids, mainly prednisolone or methylprednisolone, due to their immunosuppressive properties, have strong anti-inflammatory effects and are mainly used to reduce pneumonia, prevent the progression of respiratory failure and death, and are useful for patients with covid 19 (47, 58).

Corticosteroids alter the functions of dermal cells and leukocytes in inflammatory diseases. they can pass the cell membrane and react with receptor proteins in the cytoplasm to make a steroid-receptor complex. This complex needs to bind to DNA; thus, it moves into the nucleus. transcription of messenger RNA (mRNA) is changed following binding process. in conclusion, corticosteroids can stimulate or inhibit the synthesis of specific proteins (59) (Figure 10).

Corticosteroids should be used just in cases of chronic obstructive pulmonary disease exacerbation or septic shock, as recommended by the WHO and the Centers for Disease Control and Prevention (CDC) and Infectious Disease Society of America (IDSA) (47). In addition the dosage should be low-to-moderate (≤ 0.5 – 1 mg/kg per day methylprednisolone or equivalent) and the duration should be short (≤ 7 days) (60). adverse effects such as e.g., hyperglycemia, neuropsychiatric symptoms, secondary infections (57). The current treatment with corticosteroid in COVID-19 is specialized to patients with fatal conditions that are correlated with cytokine storm such as ARDS, renal failure, and acute cardiac injury (57).

5. Monoclonal antibodies therapy:

In addition to anti interleukin Monoclonal antibodies that we described above several monoclonal antibodies have been developed to target the spike protein of SARS-CoV-2, including Bamlanivimab plus etesevimab, sotrovimab, and Casirivimab plus imdevimab. However, they are not approved for inpatient use and are prescribed intravenously or subcutaneously for individuals who are at least 12 years old and weigh at least 40 kg (61, 62) (Figure 11). Sotrovimab and Regdanvimab are recombinant human monoclonal antibodies that target the spike protein receptor-binding domain and receptor-binding domain (RBD) of the SARS-CoV-2 spike protein, respectively (63). Bebtelovimab binds to the viral spike protein to prevent it from attaching to the human ACE2 receptor (64). All three antibodies have received Emergency Use Authorization from the FDA for the treatment of mild-to-moderate COVID-19 (63). Sotrovimab is effective in treating mild-to-moderate COVID-19 based on clinical studies, with a recommended dosage of a single 500 mg intravenous infusion (65). Regdanvimab is administered as a single intravenous infusion at a dose of 40 mg/kg, while

Bebtelovimab is administered via IV injection at a recommended dosage of 175 mg over a minimum of 30 seconds (66, 67).

6. Anticoagulant drugs:

Many patients with COVID-19 develop clinical coagulation identified with the following signs: Thrombocytopenia, Prolonged prothrombin time (PT), and partial thromboplastin time (aPTT), Increased serum D-dimer and fibrinogen, etc. and are at hazard for major vascular thrombosis (68), therefore anticoagulants are recommended for the prevention and treatment of thrombosis (69, 70). There are many anticoagulant drugs that are classified into wide categories as well as Heparin or enoxaparin, dalteparin, and rivaroxaban (70, 71). Heparin as the first anticoagulant encompasses unfractionated heparin (UFH) and low-molecular-weight heparin (LMWH) (70).

Heparin can reduce inflammation through impeding neutrophil infiltration and inhibiting the production of inflammatory factors, such as IL-8, IL-6, and TNF- α and (68). It can also attach to the spike protein of SARS-CoV-2 and serve as a competitive suppressor for virus entry, thus reduce infectivity(69). Moreover, both unfractionated heparin (UFH) and low-molecular-weight heparin (LMWH) can bind to antithrombin III which inhibits the activation of FX. Unfractionated heparin (UFH) also can blocks thrombin by consisting heparin-antithrombin-thrombin complex and leading anticoagulation (70).

So as mentioned above, heparin has notable features such as anti-inflammatory, antiviral, and anticoagulant properties (69-71) (Figure 12). Bleeding and thrombocytopenia are major disadvantages of heparin, as well as alopecia, and hyperkalemia (69, 70).

Single nucleotide polymorphisms (SNPs) located at direct targets of anti-COVID-19 drugs and may interact with their functions:

As we discussed above inflammatory elements such as IL-1R, IL-6R, IL17A, IL17RA, JAK1, JAK2 could be possible targets for anti-COVID-19 medicines. But all of these genes have some SNPs which may impact on gene expression, associated with other inflammatory diseases and interferes with drug function. We provide the complete information about SNPs of these genes based on snpedia and dbsnp, in table1. Although there are limited studies about interactions between their SNPs and drugs, several studies demonstrated association of interleukin 6 two SNPS, rs12083537 and rs2228145 with tocilizumab function.

the rs2228145 SNP (single nucleotide polymorphism), which has been associated with variation in the response to tocilizumab treatment in patients with rheumatoid arthritis (RA), results in a change from guanine (G) to adenine

(A) at position -174 in the promoter region of the IL-6R gene. The A allele has been associated with increased IL-6R production and activity, whereas the G allele is associated with reduced IL-6R production and activity. One study found that patients carrying the AA genotype had a slower and less pronounced response to tocilizumab compared to those carrying the GG or GA genotypes (72). Another study found that patients with the AA genotype had higher levels of IL-6R and C-reactive protein (CRP) at baseline, and were less likely to achieve remission after tocilizumab treatment (73). Moreover, another study demonstrated better response of IL-6R rs12083537 AA compared to GA and GG (74). However, in this investigation no association was found between rs4329505 and tocilizumab therapy, AAC haplotype of three polymorphisms (rs2228145 A allele, rs12083537A and rs4329505 C allele) was associated with poor response to tocilizumab treatment (73). And GAT-haplotype was related with good response (73).

Conclusion

COVID-19 is known as an inflammatory disorder. With due attention to its high rate of mortality finding a best treatment protocol and appropriate therapeutics is imperative. In this mini-Review we described various efficient drugs prescribed in patients suffering from COVID-19 with mild to severe symptoms. We tried to collect and classify drugs and express their molecular target briefly. However, this review didn't conclude all medicine used for COVID-19 treatment and we mentioned just the medicines involving prominent evidences of high efficiency and performance. In addition we provided SNPs related to drugs prescribed against covid-19. In conclusion this review presents beneficial information about covid19 appropriate therapeutics according to the molecular targets.

References

- .1 Choi SW, Shin JS, Park SJ, Jung E, Park YG, Lee J, et al. Antiviral activity and safety of remdesivir against SARS-CoV-2 infection in human pluripotent stem cell-derived cardiomyocytes. *Antiviral Res.* 2020;184:104955.
- .2 V'kovski P, Kratzel A, Steiner S, Stalder H, Thiel V. Coronavirus biology and replication: implications for SARS-CoV-2. *Nature Reviews Microbiology.* 2021;19(3):155-70.
- .3 Pal M, Berhanu G, Desalegn C, Kandi V. Severe Acute Respiratory Syndrome Coronavirus-2 (SARS-CoV-2): An Update. *Cureus.* 2020;12(3):e7423.
- .4 Hassan SA, Sheikh FN, Jamal S, Ezeh JK, Akhtar A. Coronavirus (COVID-19): A Review of Clinical Features, Diagnosis, and Treatment. *Cureus.* 2020;12(3):e7355.
- .5 Velavan TP, Meyer CG. The COVID-19 epidemic. *Tropical medicine & international health.* 2020;25(3):278.

- .6 Carfi A, Bernabei R, Landi F. Persistent symptoms in patients after acute COVID-19. *Jama*. 2020;324(6):603-5.
- .7 Xu X, Ong YK, Wang Y. Role of adjunctive treatment strategies in COVID-19 and a review of international and national clinical guidelines. *Mil Med Res*. 2020;7(1):22.
- .8 Gandhi RT, Lynch JB, del Rio C. Mild or Moderate Covid-19. *New England Journal of Medicine*. 2020;383(18):1757-66.
- .9 Ejaz H, Alsrhani A, Zafar A, Javed H, Junaid K, Abdalla AE, et al. COVID-19 and comorbidities: Deleterious impact on infected patients. *J Infect Public Health*. 2020;13(12):1833-9.
- .10 Baj J, Karakuła-Juchnowicz H, Teresiński G, Buszewicz G, Ciesielka M, Sitarz R, et al. COVID-19: Specific and Non-Specific Clinical Manifestations and Symptoms: The Current State of Knowledge. *J Clin Med*. 2020;9.(6)
- .11 Hafeez A, Ahmad S, Siddqui SA, Ahmad M, Mishra S. A review of COVID-19 (Coronavirus Disease-2019) diagnosis, treatments and prevention. *EJMO*. 2020;4(2):116-25.
- .12 Aleem A, Slenker AK. Monoclonal Antibody Therapy For High-Risk Coronavirus (COVID 19) Patients With Mild To Moderate Disease Presentations. *StatPearls*. Treasure Island (FL): StatPearls Publishing
- Copyright © 2022, StatPearls Publishing LLC.; 2022.
- .13 Zhang W, Zhao Y, Zhang F, Wang Q, Li T, Liu Z, et al. The use of anti-inflammatory drugs in the treatment of people with severe coronavirus disease 2019 (COVID-19): The Perspectives of clinical immunologists from China. *Clin Immunol*. 2020;214:108393.
- .14 Shastry BS. SNPs: impact on gene function and phenotype. *Methods Mol Biol*. 2009;578:3-22.
- .15 Morris R, Kershaw NJ, Babon JJ. The molecular details of cytokine signaling via the JAK/STAT pathway. *Protein Sci*. 2018;27(12):1984-2009.
- .16 Zhang JM, An J. Cytokines, inflammation, and pain. *Int Anesthesiol Clin*. 2007;45(2):27-37.
- .17 Mukai K, Tsai M, Saito H, Galli SJ. Mast cells as sources of cytokines, chemokines, and growth factors. *Immunol Rev*. 2018;282(1):121-50.
- .18 Fajgenbaum DC, June CH. Cytokine Storm. *N Engl J Med*. 2020.73-2255:(23)383;
- .19 Cavalli G, De Luca G, Campochiaro C, Della-Torre E, Ripa M, Canetti D, et al. Interleukin-1 blockade with high-dose anakinra in patients with COVID-19, acute respiratory distress syndrome, and hyperinflammation: a retrospective cohort study. *The Lancet Rheumatology*. 2020;2(6):e325-e31.
- .20 Ragab D, Salah Eldin H, Taeimah M, Khattab R, Salem R. The COVID-19 Cytokine Storm; What We Know So Far. *Frontiers in Immunology*. 2020;11.
- .21 Sebbar EH, Choukri M. Interleukin 6: A biomarker for COVID-19 progression. *Mater Today Proc*. 2022.
- .22 Kaneko N, Kurata M, Yamamoto T, Morikawa S, Masumoto J. The role of interleukin-1 in general pathology. *Inflammation and Regeneration*. 2019;39(1):12.
- .23 Chen L, Deng H, Cui H, Fang J, Zuo Z, Deng J, et al. Inflammatory responses and inflammation-associated diseases in organs. *Oncotarget*. 2018;9(6):7204-18.
- .24 Dinarello CA. Overview of the IL-1 family in innate inflammation and acquired immunity. *Immunol Rev*. 2018;281(1):8-27.
- .25 Giat E, Ben-Zvi I, Lidar M, Livneh A. The Preferential Use of Anakinra in Various Settings of FMF: A Review Applied to an Updated Treatment-Related Perspective of the Disease. *Int J Mol Sci*. 2022;23.(7)
- .26 Kyriazopoulou E, Poulakou G, Milionis H, Metallidis S, Adamis G, Tsiakos K, et al. Early treatment of COVID-19 with anakinra guided by soluble urokinase plasminogen receptor plasma levels: a double-blind, randomized controlled phase 3 trial. *Nat Med*. 2021;27(10):1752-60.
- .27 Maes B, Bosteels C, De Leeuw E, Declercq J, Van Damme K, Delporte A, et al. Treatment of severely ill COVID-19 patients with anti-interleukin drugs (COV-AID): A structured summary of a study protocol for a randomised controlled trial. *Trials*. 2020;21(1):468.

- .28 Shekhawat J, Gauba K, Gupta S, Purohit P, Mitra P, Garg M, et al. Interleukin-6 Perpetrator of the COVID-19 Cytokine Storm. *Indian J Clin Biochem.* 2021;36(4):440-50.
- .29 Bryushkova EA, Skatova VD, Mutovina ZY, Zagrebneva AI, Fomina DS, Kruglova TS, et al. Tocilizumab, netakimab, and baricitinib in patients with mild-to-moderate COVID-19: An observational study. *PLoS one.* 2022;17(8):e0273340.
- .30 Coskun Benlidayi I, Kurtaran B, Tirasci E, Guzel R. Coronavirus disease 2019 (COVID-19) in a patient with ankylosing spondylitis treated with secukinumab :a case-based review. *Rheumatol Int.* 2020;40(10):1707-16.
- .31 Ayhan E, Öztürk M, An İ, Abdelmaksoud A, Araç E. Potential role of anti-interleukin-17 in COVID-19 treatment. *Dermatologic Therapy.* 2020;33(4):e13715.
- .32 Mendoza VMM. Interleukin-17: A potential therapeutic target in COVID-19. *J Infect.* 2020;81(2):e136-e8.
- .33 Cafarotti S. Severe Acute Respiratory Syndrome-Coronavirus-2 Infection and Patients With Lung Cancer: The Potential Role of Interleukin-17 Target Therapy. *J Thorac Oncol.* 2020;15(7):e101-e3.
- .34 Singh S, Kroe-Barrett RR, Canada KA, Zhu X, Sepulveda E, Wu H, et al. Selective targeting of the IL23 pathway: Generation and characterization of a novel high-affinity humanized anti-IL23A antibody. *MAbs.* 2015;7(4):778-91.
- .35 Stebbing J, Sánchez Nieves G, Falcone M, Youhanna S, Richardson P, Ottaviani S, et al. JAK inhibition reduces SARS-CoV-2 liver infectivity and modulates inflammatory responses to reduce morbidity and mortality. *Sci Adv.* 2021;7.(1)
- .36 Assadiasl S, Fatahi Y, Mosharmovahed B ,Mohebbi B, Nicknam MH. Baricitinib: From Rheumatoid Arthritis to COVID-19. *J Clin Pharmacol.* 2021;61(10):1274-85.
- .37 Baricitinib in patients admitted to hospital with COVID-19 (RECOVERY): a randomised, controlled, open-label, platform trial and updated meta-analysis. *Lancet.* 2022;400(10349):359-68.
- .38 Shi JG, Chen X, Lee F, Emm T, Scherle PA, Lo Y, et al. The pharmacokinetics, pharmacodynamics, and safety of baricitinib, an oral JAK 1/2 inhibitor, in healthy volunteers. *The Journal of Clinical Pharmacology.* 2014;54(12):1354-61.
- .39 Jorgensen SCJ, Tse CLY, Burry L, Dresser LD. Baricitinib: A Review of Pharmacology, Safety, and Emerging Clinical Experience in COVID-19. *Pharmacotherapy.* 2020;40(8):843-56.
- .40 Ahmad A, Zaheer M, Balis FJ. Baricitinib. *StatPearls. Treasure Island (FL): StatPearls Publishing*
- Copyright © 2022, StatPearls Publishing LLC.; 2022.
- .41 Richardson P, Griffin I, Tucker C, Smith D, Oechsle O, Phelan A, et al. Baricitinib as potential treatment for 2019-nCoV acute respiratory disease .*Lancet.* 2020;395(10223):e30-e1.
- .42 Levy G, Guglielmelli P, Langmuir P, Constantinescu SN. JAK inhibitors and COVID-19. *J Immunother Cancer.* 2022;10.(4)
- .43 Bande BD. Use of Tofacitinib in the Management of COVID-19 Pneumonia. *Indian J Crit Care Med.* 202.90-1089:(10)25;1
- .44 Guimarães PO, Quirk D, Furtado RH, Maia LN, Saraiva JF, Antunes MO, et al. Tofacitinib in Patients Hospitalized with Covid-19 Pneumonia. *N Engl J Med.* 2021;385(5):406-15.
- .45 Satarker S, Tom AA, Shaji RA, Alosious A, Luvis M, Nampoothiri M. JAK-STAT Pathway Inhibition and their Implications in COVID-19 Therapy. *Postgrad Med.* 2021;133(5):489-507.
- .46 Aleem A, Kothadia J. Remdesivir. *StatPearls.* 2021.
- .47 Kapoor M, Panda PK, Mohanty V. Pharmacotherapy for COVID-19: A Ray of Hope. 2021.
- .48 Lamb YN. Remdesivir: First Approval. *Drugs.* 2020;80(13):1355-63.
- .49 Joshi S, Parkar J, Ansari A, Vora A, Talwar D, Tiwaskar M, et al. Role of favipiravir in the treatment of COVID-19. *Int J Infect Dis.* 2021;102:501-8.
- .50 Buxeraud J, Faure S, Fougere É. [Nirmatrelvir/ritonavir (Paxlovid®), a treatment for Covid-19]. *Actual Pharm.* 2022;61(617):10-2.

- .51 Amani B, Amani B. Efficacy and safety of nirmatrelvir/ritonavir (Paxlovid) for COVID-19 : a rapid review and meta-analysis. *J Med Virol.* 2022;95:(2)
- .52 Hammond J, Leister-Tebbe H, Gardner A, Abreu P, Bao W, Wisemandle W, et al. Oral Nirmatrelvir for High-Risk, Nonhospitalized Adults with Covid-19. *N Engl J Med.* 2022;386(15):1397-408.
- .53 Nirmatrelvir and ritonavir for COVID-19. *Aust Prescr.* 2022;45(2):61:(
- .54 Jayk Bernal A, Gomes da Silva MM, Musungaie DB, Kovalchuk E, Gonzalez A, Delos Reyes V, et al. Molnupiravir for Oral Treatment of Covid-19 in Nonhospitalized Patients. *N Engl J Med.* 2022;386(6):509-20.
- .55 Wen W, Chen C, Tang J, Wang C, Zhou M ,Cheng Y, et al. Efficacy and safety of three new oral antiviral treatment (molnupiravir, flvoxamine and Paxlovid) for COVID-19 : a meta-analysis. *Ann Med.* 2022;54(1):516-23.
- .56 Imran M, Kumar Arora M, Asdaq SMB, Khan SA, Alaqel SI, Alshammari MK, et al. Discovery, Development, and Patent Trends on Molnupiravir: A Prospective Oral Treatment for COVID-19. *Molecules.* 2021;26:(19)
- .57 Liu D, Ahmet A, Ward L, Krishnamoorthy P, Mandelcorn ED, Leigh R, et al. A practical guide to the monitoring and management of the complications of systemic corticosteroid therapy. *Allergy Asthma Clin Immunol.* 2013;9(1):30.
- .58 Sahebnaasagh A, Avan R, Saghafi F, Mojtahedzadeh M, Sadremomtaz A, Arasteh O, et al. Pharmacological treatments of COVID-19. *Pharmacological Reports.* 2020;78-1446:(6)72;
- .59 Kragballe K. Topical corticosteroids: mechanisms of action. *Acta Derm Venereol Suppl (Stockh).* 1989;151:7-10; discussion 47-52.
- .60 Shang L, Zhao J, Hu Y, Du R, Cao B. On the use of corticosteroids for 2019-nCoV pneumonia. *The Lancet.* 4-683:(10225)395;2020 .
- .61 Lloyd EC, Gandhi TN, Petty LA. Monoclonal Antibodies for COVID-19. *JAMA.* 2021;325(10):1015-.
- .62 Deb P, Molla MMA, Saif-Ur-Rahman KM. An update to monoclonal antibody as therapeutic option against COVID-19. *Biosafety and Health.* 91-87:(02)03;2021 .
- .63 Tuccori M, Ferraro S, Convertino I, Cappello E, Valdiserra G, Blandizzi C, et al. Anti-SARS-CoV-2 neutralizing monoclonal antibodies: clinical pipeline. *MAbs.* 2020;12(1):1854149.
- .64 Westendorf K, Žentelis S, Wang L, Foster D, Vaillancourt P, Wiggin M, et al. LY-CoV1404 (bebtelovimab) potently neutralizes SARS-CoV-2 variants. *bioRxiv.* 2022.
- .65 Gupta A, Gonzalez-Rojas Y, Juarez E, Crespo Casal M, Moya J, Falci DR, et al. Early Treatment for Covid-19 with SARS-CoV-2 Neutralizing Antibody Sotrovimab. *New England Journal of Medicine.* 2021;385(21):1941-50.
- .66 Syed YY. Regdanvimab: First Approval. *Drugs.* 2021;81(18):2133-7.
- .67 Orders M. An EUA for bebtelovimab for treatment of COVID-19. *Med Lett Drugs Ther.* 2022;64(1646):41-2.
- .68 Liu H, Hu T, Zhang C, Chen X, Zhang S, Li M, et al. Mechanisms of COVID-19 thrombosis in an inflammatory environment and new anticoagulant targets. *Am J Transl Res.* 2021;13(5):3925-41.
- .69 Hippensteel JA, LaRiviere WB, Colbert JF, Langouët-Astrié CJ, Schmidt EP. Heparin as a therapy for COVID-19: current evidence and future possibilities. *American Journal of Physiology-Lung Cellular and Molecular Physiology.* 2020;319(2):L211-L7.
- .70 Chandra A, Chakraborty U, Ghosh S, Dasgupta S. Anticoagulation in COVID-19 :current concepts and controversies. *Postgraduate Medical Journal.* 2021;98(1159):395-402.
- .71 Hirsh J, Anand SS, Halperin JL, Fuster V. Mechanism of Action and Pharmacology of Unfractionated Heparin. *Arteriosclerosis, Thrombosis, and Vascular Biology.* 2001;6-1094:(7)21;
- .72 Enevold C, Baslund B, Linde L, Josephsen NL, Tarp U, Lindegaard H, et al. Interleukin-6-receptor polymorphisms rs12083537, rs2228145, and rs4329505 as predictors of response to tocilizumab in rheumatoid arthritis. *Pharmacogenet Genomics.* 5-401:(8)24;2014 .

- .73 Janahiraman S, Too CL, Lee KW, Shahril NS, Leong CO. Genetic Biomarkers as Predictors of Response to Tocilizumab in Rheumatoid Arthritis: A Systematic Review and Meta-Analysis. *Genes (Basel)*. 2022;13.(7)
- .74 Lee YH, Song GG. Associations between the interleukin-6 rs1800795 G/C and interleukin-6 receptor rs12083537 A/G polymorphisms and response to disease-modifying antirheumatic drugs in rheumatoid arthritis: A meta-analysis. *Int Immunopharmacol*. 2022;112:109184.
- .75 Liu X, Peng L, Li D, He C, Xing S, Wang Y, et al. The Impacts of IL1R1 and IL1R2 Genetic Variants on Rheumatoid Arthritis Risk in the Chinese Han Population: A Case-Control Study. *Int J Gen Med*. 2021;14:2147-59.
- .76 An F, Wang J, Gao H, Liu C, Tian Y, Jin T, et al. Impact of IL1R1 and IL1R2 gene polymorphisms on risk of osteonecrosis of the femoral head from a case-control study. *Mol Genet Genomic Med*. 2019;7(3):e00557.
- .77 Zhu Y, Li S, Sun Y, Wu J, Xiong Z, Jin T, et al. IL1R1 Polymorphisms are Associated with Lumbar Disc Herniation Risk in the Northwestern Chinese Han Population. *Med Sci Monit*. 2019;25:3728-38.
- .78 Wang J, Shi Y, Wang G, Dong S, Yang D, Zuo X. The association between interleukin-1 polymorphisms and their protein expression in Chinese Han patients with breast cancer. *Mol Genet Genomic Med*. 2019;7(8):e804.
- .79 Xie M, Zhang D, Zhang Y, Yang X, Su Y, Wang Y, et al. Association of genetic polymorphisms in IL-1R1 and IL-1R2 genes with IgA nephropathy in the Han Chinese population. *Oncotarget*. 2017;8(31):50.9-673
- .80 Karaesmen E, Hahn T, Dile AJ, Rizvi AA, Wang J, Wang T, et al. Multiple functional variants in the IL1RL1 region are pretransplant markers for risk of GVHD and infection deaths. *Blood Adv*. 2019;3(16):2512-24.
- .81 Chen J, Zhang J, Hu H, Jin Y, Xue M. Polymorphisms of RAD50, IL33 and IL1RL1 are associated with atopic asthma in Chinese population. *Tissue Antigens*. 2015;86(6):443-7.
- .82 Yang GS, Barnes NM, Lyon DE, Dorsey SG. Genetic Variants Associated with Cancer Pain and Response to Opioid Analgesics: Implications for Precision Pain Management. *Semin Oncol Nurs*. 2019;35(3):291-9.
- .83 Sangil A, Arranz MJ, Güerri-Fernández R, Pérez M, Monzón H, Payeras A, et al. Genetic susceptibility to invasive pneumococcal disease. *Infect Genet Evol*. 2018;59:12.31-6
- .84 Park SW, Kim MK, Kwon KH, Kim J. Association between a promoter polymorphism (rs2192752, -1028A/C) of interleukin 1 receptor, type I (IL1R1) and location of papillary thyroid carcinoma in a Korean population. *Int J Immunogenet*. 2012;39(6):501-7.
- .85 Chien CY, Tai SY, Li KH, Yang HL, Chan LP, Hsi E, et al. The association of genetic polymorphisms in interleukin-1 receptors type 1 and type 2 with sudden sensorineural hearing loss in a Taiwanese population: a case control study. *J Otolaryngol Head Neck Surg*. 2021;50(1):69.
- .86 Ahir-Bist S, Chavan V, Samant-Mavani P, Nanavati R, Mehta P, Mania-Pramanik J. Polymorphisms in TH1-TH2 cytokine and receptor genes associated with risk of vertical HIV transmission, in Mumbai, India. *J Gene Med*. 2018;20(10-1):(1e3047.
- .87 Abtahi S, Farazmand A, Mahmoudi M, Ashraf-Ganjouei A, Javinani A, Nazari B, et al. IL-1A rs1800587, IL-1B rs1143634 and IL-1R1 rs2234650 polymorphisms in Iranian patients with systemic sclerosis. *Int J Immunogenet*. 2015;42(6):423-7.
- .88 Näkki A, Kouhia ST, Saarela J, Harilainen A, Tallroth K, Videman T, et al. Allelic variants of IL1R1 gene associate with severe hand osteoarthritis. *BMC Med Genet*. 2010;11:50.
- .89 Lingappa JR, Dumitrescu L, Zimmer SM, Lynfield R, McNicholl JM, Messonnier NE, et al. Identifying host genetic risk factors in the context of public health surveillance for invasive pneumococcal disease. *PLoS One*. 2011;6(8):e23413.
- .90 Zhang L, Zhou Q, Wu Z, Zhu X, Geng T. The effect of IL-1R1 and IL-1RN polymorphisms on osteoporosis predisposition in a Chinese Han population. *Int Immunopharmacol*. 2020;87:106833.

- .91 Xiong Z, Sun Y, Wu J, Niu F, Jin T, Li B. Genetic polymorphisms in IL1R1 and IL1R2 are associated with susceptibility to thyroid cancer in the Chinese Han population. *J Gene Med.* 2019;21(6):e3093.
- .92 Na Y, Bai R, Ren Y, Zhao Z, Kong L, Li R, et al. IL1R1 polymorphisms are associated with ankylosing spondylitis in the Han Chinese population: a case-control study. *Int J Clin Exp Pathol.* 2018;11(7):3759-64.
- .93 Fan J, Cai Y, Huang X, Wang Y, Mu L, Zhou L. Variations in IL-1R1 Gene Influence Risk for Hepatitis B Virus Infection of Children in a Han Chinese population. *Int J Infect Dis.* 2017;55:45-50.
- .94 Leung G, Baggott C, West C, Elboim C, Paul SM, Cooper BA, et al. Cytokine candidate genes predict the development of secondary lymphedema following breast cancer surgery. *Lymphat Res Biol.* 2014;12(1):10-22.
- .95 Akhabir L, Sandford A. Genetics of interleukin 1 receptor-like 1 in immune and inflammatory diseases. *Curr Genomics.* 2010;11(8):591-606.
- .96 Lin H, Sinner MF, Brody JA, Arking DE, Lunetta KL, Rienstra M, et al. Targeted sequencing in candidate genes for atrial fibrillation: the Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Targeted Sequencing Study. *Heart Rhythm.* 2014;11(3):452-7.
- .97 Maldonado-Montoro M, Cañadas-Garre M, González-Utrilla A, Ángel Calleja-Hernández M. Influence of IL6R gene polymorphisms in the effectiveness to treatment with tocilizumab in rheumatoid arthritis. *Pharmacogenomics J.* 2018;18(1):167-72.
- .98 Zubiaur P, Koller D, Saiz-Rodríguez M, Navares-Gómez M, Abad-Santos F. Important Pharmacogenetic Information for Drugs Prescribed During the SARS-CoV-2 Infection (COVID-19). *Clin Transl Sci.* 2020;13(6):1023-33.
- .99 Luxembourger C, Ruysen-Witrand A, Ladhari C, Rittore C, Degboe Y, Maillefert JF, et al. A single nucleotide polymorphism of IL6-receptor is associated with response to tocilizumab in rheumatoid arthritis patients. *Pharmacogenomics J.* 2019;19(4):368-74.
- .100 Huang X, Ye Q, Zhu Z, Chen W, Chen Y, Li J, et al. Polymorphism of IL6 receptor gene is associated with ischaemic stroke in patients with metabolic syndrome. *Brain Res.* 2020;1728:146594.
- .101 Hansen PR, Nelveg-Kristensen KE, Rasmussen HB, Torp-Pedersen C, Køber L, Nielsen CH, et al. Prognostic role of genetic polymorphisms of the interleukin-6 signaling pathway in patients with severe heart failure. *Pharmacogenomics J.* 2019;19(5):428-37.
- .102 Wang Y, Hu H, Wu J, Zhao X, Zhen Y, Wang S, et al. The IL6R gene polymorphisms are associated with sIL-6R, IgE and lung function in Chinese patients with asthma. *Gene.* 2016;585(1):51-7.
- .103 Gigante B, Strawbridge RJ, Velasquez IM, Golabkesh Z, Silveira A, Goel A, et al. Analysis of the role of interleukin 6 receptor haplotypes in the regulation of circulating levels of inflammatory biomarkers and risk of coronary heart disease. *PLoS One.* 2015;10(3):e0119980.
- .104 Puchenkova OA, Soldatov VO, Belykh AE, Bushueva O, Piavchenko GA, Venediktov AA, et al. Cytokines in Abdominal Aortic Aneurysm: Master Regulators With Clinical Application. *Biomark Insights.* 2022;17:11772719221095676.
- .105 Padyukov L. Genetics of rheumatoid arthritis. *Semin Immunopathol.* 2022;44(1):47-62.
- .106 Zhang M, Bai Y, Wang Y, Cui H, Tang M, Wang L, et al. Cumulative Evidence for Associations Between Genetic Variants in Interleukin 6 Receptor Gene and Human Diseases and Phenotypes. *Front Immunol.* 2022;13:860703.
- .107 Rokni M, Sarhadi M, Heidari Nia M, Mohamed Khosroshahi L, Asghari S, Sargazi S, et al. Single nucleotide polymorphisms located in TNFA, IL1RN, IL6R, and IL6 genes are associated with COVID-19 risk and severity in an Iranian population. *Cell Biol Int.* 2022;46(7):1109-27.
- .108 Xu H, Liu J, Niu M, Song S, Wei L, Chen G, et al. Soluble IL-6R-mediated IL-6 trans-signaling activation contributes to the pathological development of psoriasis. *J Mol Med (Berl).* 2021;99(7):1009-20.

- .109 Ribeiro de Andrade Ramos B, Cosi Bento GF, Navascues Bernardino RA, Miot HA, Guimarães da Silva M. Influence of single nucleotide polymorphisms (SNPs) in immunoregulatory genes in the morbidity of preterm newborns. *J Matern Fetal Neonatal Med.* 2021;34(22):3684-9.
- .110 Vogrinc D, Goričar K, Dolžan V. Genetic Variability in Molecular Pathways Implicated in Alzheimer's Disease: A Comprehensive Review. *Front Aging Neurosci.* 2021;13:646901.
- .111 Mikhailova SV, Ivanoshchuk DE. Innate-Immunity Genes in Obesity. *J Pers Med.* 2021;11.(11)
- .112 Coller JK, Tuke J, Wain TJ, Quinn E, Steele L, Abreu M, et al. Associations of Immune Genetic Variability with Gulf War Illness in 1990-1991 Gulf War Veterans from the Gulf War Illness Consortium (GWIC) Multisite Case-Control Study. *Brain Sci.* 2021;11.(11)
- .113 Lin J, Wang Y, Wang Y, Pan Y. Inflammatory biomarkers and risk of ischemic stroke and subtypes: A 2-sample Mendelian randomization study. *Neurol Res.* 2020;42(2):118-25.
- .114 Aparicio-Siegmund S, Garbers Y, Flynn CM, Waetzig GH, Gouni-Berthold I, Krone W, et al. The IL-6-neutralizing sIL-6R-sgp130 buffer system is disturbed in patients with type 2 diabetes. *Am J Physiol Endocrinol Metab.* 2019;317(2):E411-e20.
- .115 Sundaresh A, Oliveira J, Chinnadurai RK, Rajkumar RP, Hani L, Krishnamoorthy R, et al. IL6/IL6R genetic diversity and plasma IL6 levels in bipolar disorder: An Indo-French study. *Heliyon.* 2019;5(1):e01124.
- .116 Wu W, Yang H, Feng Y, Zhang P, Li S, Wang X, et al. Polymorphisms in Inflammatory Mediator Genes and Risk of Preeclampsia in Taiyuan, China. *Reprod Sci.* 2017;24(4):539-47.
- .117 Ferreira MA, Matheson MC, Duffy DL, Marks GB, Hui J, Le Souëf P, et al. Identification of IL6R and chromosome 11q13.5 as risk loci for asthma. *Lancet.* 2011;378(9795):1006-14.
- .118 Ruan WF, Xie JT, Jin Q, Wang WD, Ping AS. The Diagnostic and Prognostic Role of Interleukin 12B and Interleukin 6R Gene Polymorphism in Patients With Ankylosing Spondylitis. *J Clin Rheumatol.* 2018;24(1):18-24.
- .119 Liu K, Xie Y, Zhao Q, Peng W, Guo C, Zhang J, et al. Polymorphisms and Gene-Gene Interaction in AGER/IL6 Pathway Might Be Associated with Diabetic Ischemic Heart Disease. *J Pers Med.* 2022;12.(3)
- .120 Lopez-Lasanta M, Julià A, Maymó J, Fernández-Gutierrez B, Ureña-Garnica I, Blanco FJ, et al. Variation at interleukin-6 receptor gene is associated to joint damage in rheumatoid arthritis. *Arthritis Res Ther.* 2024;(1)17;15
- .121 Chu NF, Lin FH, Chin HC, Hong YJ. Association between interleukin-6 receptor gene variations and atherosclerotic lipid profiles among young adolescents in Taiwan. *Lipids Health Dis.* 2011;10:136.
- .122 Kim DH, Yoo SD, Chon J, Yun DH, Kim HS, Park HJ, et al. Interleukin-6 Receptor Polymorphisms Contribute to the Neurological Status of Korean Patients with Ischemic Stroke. *J Korean Med Sci.* 2016;31(3):430-4.
- .123 Gu F, Qureshi AA, Niu T, Kraft P, Guo Q, Hunter DJ, et al. Interleukin and interleukin receptor gene polymorphisms and susceptibility to melanoma. *Melanoma Res.* 2008;18(5):330-5.
- .124 Abe S, Tokoro F, Matsuoka R, Arai M, Noda T, Watanabe S, et al. Association of genetic variants with dyslipidemia. *Mol Med Rep.* 2015;12(4):5429-36.
- .125 Horibe H, Fujimaki T, Oguri M, Kato K, Matsuoka R, Abe S, et al. Association of a polymorphism of the interleukin 6 receptor gene with chronic kidney disease in Japanese individuals. *Nephrology (Carlton).* 2015;20(4):273-8.
- .126 Kessler T, Vilne B, Schunkert H. The impact of genome-wide association studies on the pathophysiology and therapy of cardiovascular disease. *EMBO Mol Med.* 2016;8(7):688-701.
- .127 Guo Y, Garcia-Barrio MT, Wang L, Chen YE. Experimental Biology for the Identification of Causal Pathways in Atherosclerosis. *Cardiovasc Drugs Ther.* 2016;30(1):1-11.
- .128 Birmann BM, Tamimi RM, Giovannucci E, Rosner B, Hunter DJ, Kraft P, et al. Insulin-like growth factor-1- and interleukin-6-related gene variation and risk of multiple myeloma. *Cancer Epidemiol Biomarkers Prev.* 2009;18(1):282-8.

- .129 Marquet S, Conte I, Poudiougou B, Argiro L, Cabantous S, Dessein H, et al. The IL17F and IL17RA Genetic Variants Increase Risk of Cerebral Malaria in Two African Populations. *Infect Immun*. 2016;84(2):590-7.
- .130 McCaughan JA, McKnight AJ, Maxwell AP. Genetics of new-onset diabetes after transplantation. *J Am Soc Nephrol*. 2014;25(5):1037-49.
- .131 Somers J, Ruttens D, Verleden SE, Vandermeulen E, Piloni D, Wauters E, et al. Interleukin-17 receptor polymorphism predisposes to primary graft dysfunction after lung transplantation. *J Heart Lung Transplant*. 2015;34(7):941-9.
- .132 Ozkol HU, Gorgisen G, Ates C, Özkol H, Tülüce Y, Savas H, et al. Evaluation of the relationship of IL-17A and IL-17F gene polymorphisms with the response to treatment in psoriatic patients using biological drugs: a case-control study in patients in Eastern Turkey. *Postepy Dermatol Alergol*. 2021;38(5):780-7.
- .133 Caputo V, Strafella C, Cosio T, Lanna C, Campione E, Novelli G, et al. Pharmacogenomics: An Update on Biologics and Small-Molecule Drugs in the Treatment of Psoriasis. *Genes (Basel)*. 2021;12.(9)
- .134 Murdaca G, Negrini S, Magnani O, Penza E, Pellicchio M, Puppo F. Impact of pharmacogenomics upon the therapeutic response to etanercept in psoriasis and psoriatic arthritis. *Expert Opin Drug Saf*. 2017;16(10):1173-9.
- .135 Zhang X, Ye T, Li M, Yan H, Lin H, Lu H, et al. Association of Polymorphisms in Inflammation Genes With the Prognosis of Advanced Non-Small Cell Lung Cancer Patients Receiving Epidermal Growth Factor Receptor Tyrosine Kinase Inhibitors. *Front Oncol*. 2022;12:836117.
- .136 Cai T, Wang G, Yang Y, Mu K, Zhang J, Jiang Y, et al. Association Between Polymorphisms of IL-23/IL-17 Pathway and Clinical Phenotypes of Autoimmune Thyroid Diseases. *Iran J Immunol*. 2022;19(2):139-49.
- .137 Chen A, Zhao H, Wang J, Zhang R, Liu J, Zhao X, et al. Haplotype Analysis of Candidate Genes Involved in Inflammation and Oxidative Stress and the Susceptibility to Preeclampsia. *J Immunol Res*. 2020;2020:4683798;
- .138 Andrade JMA, de Oliveira CBS, Meurer Y, Santana JE, de Almeida YGB, Vilela Dos Santos P, et al. Genetic polymorphism in IL17RA induces susceptibility to *Toxoplasma gondii* infection in Brazilian pregnant women. *Acta Trop*. 2020;211:10559.4
- .139 Sandip C, Tan L, Huang J, Li Q, Ni L, Cianflone K, et al. Common variants in IL-17A/IL-17RA axis contribute to predisposition to and progression of congestive heart failure. *Medicine (Baltimore)*. 2016;95(27):e4105.
- .140 Ruttens D, Wauters E, Kiciński M, Verleden SE, Vandermeulen E, Vos R, et al. Genetic variation in interleukin-17 receptor A is functionally associated with chronic rejection after lung transplantation. *J Heart Lung Transplant*. 2013;32(12):1233-40.
- .141 McGovern DP, Rotter JI, Mei L, Haritunians T, Landers C, Derkowski C, et al. Genetic epistasis of IL23/IL17 pathway genes in Crohn's disease. *Inflamm Bowel Dis*. 2009;15(6):883-9.
- .142 Bedoui S, Dallel M, Barbirou M, Stayoussef M, Mokrani A, Mezlini A, et al. Interleukin-17A polymorphisms predict the response and development of tolerance to FOLFOX chemotherapy in colorectal cancer treatment. *Cancer Gene Ther*. 2020;27(5):311-8.
- .143 Popp NA, Yu D, Green B, Chew EY, Ning B, Chan CC, et al. Functional single nucleotide polymorphism in IL17-A 3' untranslated region is targeted by miR-4480 in vitro and may be associated with age-related macular degeneration. *Environ Mol Mutagen*. 2016;57(1):58-64.
- .144 Borilova Linhartova P, Kastovsky J, Lucanova S, Bartova J, Poskerova H, Vokurka J, et al. Interleukin-17A Gene Variability in Patients with Type 1 Diabetes Mellitus and Chronic Periodontitis: Its Correlation with IL-17 Levels and the Occurrence of Periodontopathic Bacteria. *Mediators Inflamm*. 2016;2016:2979846.
- .145 Li W, Shi W, Yin Y, Chen J, Luo L. Association of IL-17 and IL-23 Gene Variants with Plasma Levels and Risk of Vulvovaginal Candidiasis in a Chinese Han Population. *Pharmgenomics Pers Med*. 2020;13:725-33.

- .146 Tayefinasrabadi H, Mohebbsi SR, Hosseini SM, Azimzadeh P, Pourhoseingholi MA, Ghaemi A, et al. Association of Interleukin-17 gene polymorphisms with susceptibility to chronic hepatitis B virus infection and clearance in Iranian population. *Microb Pathog.* 2020;144:104195.
- .147 Bedoui SA, Barbirou M, Stayoussef M, Dallel M, Mokrani A, Makni L, et al. Association of interleukin-17A polymorphisms with the risk of colorectal cancer: A case-control study. *Cytokine.* 2018;110:18-23.
- .148 Silva MJ, de Santana MBR, Tosta BR, Espinheira RP, Alcantara-Neves NM, Barreto ML, et al. Variants in the IL17 pathway genes are associated with atopic asthma and atopy makers in a South American population. *Allergy Asthma Clin Immunol.* 2019;15:28.
- .149 Korytina GF, Akhmadishina LZ, Kochetova OV, Aznabaeva YG, Zagidullin SZ, Victorova TV. Inflammatory and Immune Response Genes Polymorphisms are Associated with Susceptibility to Chronic Obstructive Pulmonary Disease in Tatars Population from Russia. *Biochem Genet.* 2016;54(4):388-412.
- .150 Nakada TA, Russell JA, Boyd JH, Walley KR. IL17A genetic variation is associated with altered susceptibility to Gram-positive infection and mortality of severe sepsis. *Crit Care.* 2011;15(5):R254.
- .151 Li W, Li L, He L, Du Y, Fu HD, Peng ZY, et al. Cytokine Gene Polymorphisms in Chinese Children with Idiopathic Nephrotic Syndrome. *Iran J Immunol.* 2022;19(1):9.
- .152 Salvador-Martín S, Bossacoma F, Pujol-Muncunill G, Navas-López VM, Gallego-Fernández C, Viada J, et al. Genetic Predictors of Long-term Response to Antitumor Necrosis Factor Agents in Pediatric Inflammatory Bowel Disease. *J Pediatr Gastroenterol Nutr.* 2020;71(4):508-15.
- .153 Marwa OS, Kalthoum T, Wajih K, Kamel H. Association of IL17A and IL17F genes with rheumatoid arthritis disease and the impact of genetic polymorphisms on response to treatment. *Immunol Lett.* 2017;183:24-36.
- .154 Bank S, Andersen PS, Burisch J, Pedersen N, Roug S, Galsgaard J, et al. Associations between functional polymorphisms in the NFκB signaling pathway and response to anti-TNF treatment in Danish patients with inflammatory bowel disease. *Pharmacogenomics J.* 2014;14(6):526-34.
- .155 Liao H, Huang Z, Zhang J, Yang B. Association of genetic polymorphisms in IL-23R and IL-17A with the susceptibility to IgA nephropathy in a Chinese Han population. *Genes Immun.* 2022;23(1):33-41.
- .156 Zou L, Cheng Y, Yang L, Zhang F, Zhao H, Nian L, et al. Association of IL-17A gene polymorphism rs2275913 with the polycystic ovary syndrome in Yunnan Province, China. *Eur J Obstet Gynecol Reprod Biol.* 2022;271:27-30.
- .157 Das AP, Saini S, Agarwal SM. A comprehensive meta-analysis of non-coding polymorphisms associated with precancerous lesions and cervical cancer. *Genomics.* 2022;114(3):110323.
- .158 Padhi S, Sarangi S, Nayak N, Barik D, Pati A, Panda AK. Interleukin 17A rs2275913 polymorphism is associated with susceptibility to systemic lupus erythematosus: A meta and trial sequential analysis. *Lupus.* 2022;31(6):674-83.
- .159 Subbanna M, Shivakumar V, Bhalerao G, Varambally S, Venkatasubramanian G, Debnath M. Variants of Th17 pathway-related genes influence brain morphometric changes and the risk of schizophrenia through epistatic interactions. *Psychiatr Genet.* 2022;32(4):146-55.
- .160 Li G, Ma J, Zhang N, Li X, Li F, Jiang Y. The associations between interleukin-17 single-nucleotide polymorphism and colorectal cancer susceptibility: a systematic review and meta-analysis. *World J Surg Oncol.* 2022;20(1):116.
- .161 Shao M, Xu W, Yang H, Chen Y, Gao X, Xu S, et al. Interleukin-17 Gene Polymorphism (Rs2275913 G/A, Rs763780 C/T) in Rheumatoid arthritis : Meta-analysis Based on Ethnicity. *Immunol Invest.* 2021;50(6):685-99.
- .162 Mostafa AM, Saafan HA, Al-Tawashi AS, Kasem MH, Alaa AM, Eltobgy MM, et al. Interleukin-17 haplotyping predicts hepatocellular carcinoma in sofosbuvir, pegylated interferon-alpha-2a & ribavirin treated chronic hepatitis C patients. *Virus Res.* 2021;292:198226.

- .163 Chen L, Li XG, Wang JF, Hao RS, Xiang WY, Tan PF, et al. Potential effects of IL-17A rs2275913 and IL-17F rs763780 polymorphisms on susceptibility to gastric cancer in Chinese population: a meta-analysis. *Eur Rev Med Pharmacol Sci.* 2020;24(7):3633-41.
- .164 Strauss M, Palma-Vega M, Casares-Marfil D, Bosch-Nicolau P, Lo Presti MS, Molina I, et al. Genetic polymorphisms of IL17A associated with Chagas disease: results from a meta-analysis in Latin American populations. *Sci Rep.* 2020;10(1):5015.
- .165 Dimberg J, Rubér M, Skarstedt M, Andersson M, Andersson RE. Genetic polymorphism patterns suggest a genetic driven inflammatory response as pathogenesis in appendicitis. *Int J Colorectal Dis.* 84-277:(2)35;2020 .
- .166 Aguín N, Rodríguez-Alonso A, López-Trigo N, Castuera IP, Luis JR, Caeiro B. Association Between the Interleukin-17 Gene Polymorphism -197G>A and the Risk of Prostate Cancer in a Galician Population. *Pathol Oncol Res.* 2020;26(1):483-9-
- .167 Hu WL, Ren H, Xu BF, Zhang JP, Zhang RL, Wang QQ, et al. Evaluation of IL-17A, IL-17F, IL-23R, VDR, CCL2, CCL5, CCR2, and CCR5 gene polymorphisms and expression in Chinese individuals with syphilis. *J Cell Biochem.* 2018;119(12):10151-64.
- .168 Keshavarz M, Namdari H, Farahmand M, Mehrbod P, Mokhtari-Azad T, Rezaei F. Association of polymorphisms in inflammatory cytokines encoding genes with severe cases of influenza A/H1N1 and B in an Iranian population. *Virologia.* 2019;16(1):79.
- .169 Xie M, Cheng B, Ding Y, Wang C, Chen J. Correlations of IL-17 and NF- κ B gene polymorphisms with susceptibility and prognosis in acute respiratory distress syndrome in a Chinese population. *Biosci Rep.* 2019;39:(2)
- .170 Aquino JS, Ambrosio-Albuquerque EP, Alves HV, Macedo LC, Visentainer L, Sell AM, et al. IL8 and IL17A polymorphisms associated with multibacillary leprosy and reaction type 1 in a mixed population from southern Brazil. *Ann Hum Genet.* 2019;83(2):110-4.
- .171 Akbulut UE, Emeksiz HC, Citli S, Cebi AH, Korkmaz HAA, Baki G. IL-17A, MCP-1, CCR-2, and ABCA1 polymorphisms in children with non-alcoholic fatty liver disease. *J Pediatr (Rio J).* 2019;95(3):350-7.
- .172 Xiang H, Cheng D, Guo H, Wang Y, Jia Z, Gao Q. Relationships of interleukin-17 polymorphisms with recurrent aphthous ulcer risk in a Han Chinese population. *J Int Med Res.* 2020;48(12):300060520976833.
- .173 Yang HY, Liu YZ, Zhou XD, Huang Y, Xu NW. Role of IL-17 gene polymorphisms in osteoarthritis: A meta-analysis based on observational studies. *World J Clin Cases.* 2020;8(11):2280-93.
- .174 Krajewski W, Karabon L, Partyka A, Tomkiewicz A, Poletajew S, Tukiendorf A, et al. Polymorphisms of genes encoding cytokines predict the risk of high-grade bladder cancer and outcomes of BCG immunotherapy. *Cent Eur J Immunol.* 2020;45(1):37-47.
- .175 Du J, Han JC, Zhang YJ, Qi GB, Li HB, Zhang YJ, et al. Single-Nucleotide Polymorphisms of IL-17 Gene Are Associated with Asthma Susceptibility in an Asian Population. *Med Sci Monit.* 2016;22:780-7.
- .176 Huang HT, Lu YL, Wang R, Qin HM, Wang CF, Wang JL, et al. The association of IL-17A polymorphisms with IL-17A serum levels and risk of ischemic stroke. *Oncotarget.* 2017;8(61):103499-508.
- .177 Yang B, Xu Y, Liu X, Huang Z, Wang L. IL-23R and IL-17A polymorphisms correlate with susceptibility of ankylosing spondylitis in a Southwest Chinese population. *Oncotarget.* 2017;8(41):70310-6.
- .178 Akbulut UE, Cebi AH, Sağ E, İkbāl M, Çakır M. Interleukin-6 and interleukin-17 gene polymorphism association with celiac disease in children. *Turk J Gastroenterol.* 2017;28(6):471-5.
- .179 Si FZ, Feng YQ, Han M. Association between interleukin-17 gene polymorphisms and the risk of laryngeal cancer in a Chinese population. *Genet Mol Res.* 2017;16:(1)

- .180 Zhang M, Xu J, Bao X, Niu W, Wang L, Du L, et al. Association between Genetic Polymorphisms in Interleukin Genes and Recurrent Pregnancy Loss - A Systematic Review and Meta-Analysis. *PLoS One*. 2017;12(1):e0169891.
- .181 Pinto LA, LA DEAL, Mocellin M, Acosta P, Caballero MT, Libster R, et al. IL-8/IL-17 gene variations and the susceptibility to severe viral bronchiolitis. *Epidemiol Infect*. 2017;145(4):642-6.
- .182 He Y, Du Y, Wei S, Shi J, Mei Z, Qian L, et al. IL-17A and IL-17F single nucleotide polymorphisms associated with lung cancer in Chinese population. *Clin Respir J*. 2017;11(2):230-42.
- .183 Kim ES, Kim SW, Moon CM, Park JJ, Kim TI, Kim WH, et al. Interactions between IL17A, IL23R, and STAT4 polymorphisms confer susceptibility to intestinal Behcet's disease in Korean population. *Life Sci*. 2012.6-740:(20-19)90;
- .184 Wang L, Jiang Y, Zhang Y, Wang Y, Huang S, Wang Z, et al. Association analysis of IL-17A and IL-17F polymorphisms in Chinese Han women with breast cancer. *PLoS One*. 2012;7(3):e34400.
- .185 Saraiva AM, Alves e Silva MR, Correia Silva Jde F, da Costa JE, Gollob KJ, Dutra WO, et al. Evaluation of IL17A expression and of IL17A, IL17F and IL23R gene polymorphisms in Brazilian individuals with periodontitis. *Hum Immunol*. 2013;74(2):207-14.
- .186 Mucientes A, Márquez A, Cordero-Coma M, Martín-Villa JM, Gorroño-Echebarría MB, Blanco R, et al. Specific association of IL17A genetic variants with panuveitis. *Br J Ophthalmol*. 2015;99(4):566-70.
- .187 Li N, Zhang C, Chen Z, Bai L, Nie M, Zhou B, et al. Interleukin 17A and interleukin 17F polymorphisms are associated with oral squamous cell carcinoma susceptibility in a Chinese population. *J Oral Maxillofac Surg*. 2015;73(2):267-73.
- .188 Zhang X, Zheng L, Sun Y, Zhang X. Analysis of the association of interleukin-17 gene polymorphisms with gastric cancer risk and interaction with *Helicobacter pylori* infection in a Chinese population. *Tumour Biol*. 2014;35(2):1575-80.
- .189 Qi Y, Zheng H, Liu N, Guo T, Zhu W, Wang S, et al. Genetic association between Interleukin-17A gene polymorphisms and the pathogenesis of Graves' disease in the Han Chinese population. *Clin Endocrinol (Oxf)*. 2016;84(2):265-70.
- .190 Yu ZG, Wang BZ, Li J, Ding ZL, Wang K. Association between interleukin-17 genetic polymorphisms and tuberculosis susceptibility: an updated meta-analysis. *Int J Tuberc Lung Dis*. 2017;21(12):1307-13.
- .191 Ben Jmaa M, Abida O, Fakhfakh R, Bahloul E, Sellami K, Gaddour L, et al. Involvement of the IL23/Th17 Pathway in the Pathogenesis of Tunisian Pemphigus Foliaceus. *Mediators Inflamm*. 2018;2018:8206983.
- .192 Liang T, Xu YT, Zhang Y, Cai PC, Hu LH. Interleukin-17A and -17F single nucleotide polymorphisms associate with susceptibility of asthma in Chinese Han population. *Hum Immunol*. 2018;79(10):736-42.
- .193 Keramat F, Kazemi S, Saidijam M, Zamani A, Kohan HF, Mamani M, et al. Association of interleukin-17 gene polymorphisms and susceptibility to brucellosis in Hamadan, western Iran. *Microbiol Immunol*. 2019;63(3-4):139-46.
- .194 He B, Pan B, Pan Y, Wang X, Zhou L, Sun H, et al. Polymorphisms of IL-23R predict survival of gastric cancer patients in a Chinese population. *Cytokine*. 2019;117:79-83.
- .195 de Moura EL, Dos Santos ACM, da Silva DM, Dos Santos BB, Figueredo DS, Moura AWA, et al. Association of Polymorphisms in Cytokine genes with susceptibility to Precancerous Lesions and Cervical Cancer: A systematic review with meta-analysis. *Immunol Invest*. 2021;50(5):492-526.
- .196 Yang J. Role and mechanism of IL - 17 and its gene polymorphisms in dyslipidemia caused by obstructive sleep apnea syndrome in children. *Cell Mol Biol (Noisy-le-grand)*. 2022;68(2):208-12.
- .197 Zhang X, Yu P, Wang Y, Jiang W, Shen F, Wang Y, et al. Genetic polymorphisms of interleukin 17A and interleukin 17F and their association with inflammatory bowel disease in a Chinese Han population. *Inflamm Res*. 2013;62(8):743-50.

- .198 Ren Z, Li M, Liu R, Wang Y, Gu H. Interleukin 17A rs3819024 A>G polymorphism is associated with an increased risk of gastric cardia adenocarcinoma in a Chinese population. *Biomarkers*. 2014;19(5):411-6.
- .199 Vargas-Alarcón G, Angeles-Martínez J, Villarreal-Molina T, Alvarez-León E, Posadas-Sánchez R, Cardoso-Saldaña G, et al. Interleukin-17A gene haplotypes are associated with risk of premature coronary artery disease in Mexican patients from the Genetics of Atherosclerotic Disease (GEA) study. *PLoS One*. 2015;10(1):e0114943.
- .200 Overton NL, Denning DW, Bowyer P, Simpson A. Genetic susceptibility to allergic bronchopulmonary aspergillosis in asthma: a genetic association study. *Allergy Asthma Clin Immunol*. 2016;12:47.
- .201 Rimachi Hidalgo MA, Cirelli T, da Silva BR, Nicchio IG, Nepomuceno R, Orrico SRP, et al. Polymorphisms and haplotypes in the Interleukin 17 Alfa gene: potential effect of smoking habits in the association with periodontitis and type 2 diabetes mellitus. *Mol Biol Rep*. 2021;48(2):1103-14.
- .202 Lee YH, Bae SC. Associations between circulating IL-17 levels and rheumatoid arthritis and between IL-17 gene polymorphisms and disease susceptibility: a meta-analysis. *Postgrad Med J*. 2017;93(1102):465-71.
- .203 Xu H, Pan Y, Li W, Fu H, Zhang J, Shen H, et al. Association between IL17A and IL17F polymorphisms and risk of Henoch-Schonlein purpura in Chinese children. *Rheumatol Int*. 2016;36(6):829-35.
- .204 Shen L, Zhang H, Yan T, Zhou G, Liu R. Association between interleukin 17A polymorphisms and susceptibility to rheumatoid arthritis in a Chinese population. *Gene*. 2015;566(1):18-22.
- .205 Jeyakumar N, Aldoss I, Yang D, Mokhtari S, Gendzekhadze K, Khaled S, et al. Cytokine gene polymorphisms are associated with response to blinatumomab in B-cell acute lymphoblastic leukemia. *Eur J Haematol*. 2021;106(6):851-8.
- .206 Ju H, Liu H, Tian ZB, Jiang YP, Zhang CP, Liu XS. Association of polymorphisms in key Th-17 immune response genes with HBeAg-positive chronic hepatitis B susceptibility and response to PEG-IFNa-2α. *Virology*. 2017;509:35-41.
- .207 Wu W, Zeng Y, Lin J, Chen T, Xun Z, Li B, et al. IL-17 and IL-21 polymorphisms in relation to HBV related hepatocellular carcinoma in Chinese Han population. *Infect Genet Evol*. 2021;87:104638;1
- .208 Korppi M, Teräsjärvi J, Liehu-Martiskainen M, Lauhkonen E, Vuononvirta J, Nuolivirta K, et al. Haplotype of the Interleukin 17A gene is associated with osteitis after Bacillus Calmette-Guerin vaccination. *Sci Rep*. 2017;7(1):11691.
- .209 Wang J, Liu Y, Xie L, Li S, Qin X. Association of IL-17A and IL-17F gene polymorphisms with chronic hepatitis B and hepatitis B virus-related liver cirrhosis in a Chinese population: A case-control study. *Clin Res Hepatol Gastroenterol*. 2016;40(3):288-96.
- .210 Márquez A, Hernández-Rodríguez J, Cid MC, Solans R, Castañeda S, Fernández-Contreras ME, et al. Influence of the IL17A locus in giant cell arteritis susceptibility. *Ann Rheum Dis*. 2014;73(9):1742-5.
- .211 Ponce-Gallegos MA, González-Pérez MI, Mejía M, Nava-Quiroz KJ, Pérez-Rubio G, Buendía-Roldán I, et al. Single Nucleotide Polymorphism in the IL17A Gene Is Associated with Interstitial Lung Disease Positive to Anti-Jo1 Antisynthetase Autoantibodies. *Life (Basel)*. 2021;11.(2)
- .212 Hennessy MD, Zak RS, Gay CL, Pullinger CR, Lee KA, Aouizerat BE. Polymorphisms of interleukin-1 Beta and interleukin-17Alpha genes are associated with restless legs syndrome. *Biol Res Nurs*. 2014;16(2):143-51.
- .213 Stappers MH, Thys Y, Oosting M, Plantinga TS, Ioana M, Reimnitz P, et al. Polymorphisms in cytokine genes IL6, TNF, IL10, IL17A and IFNG influence susceptibility to complicated skin and skin structure infections. *Eur J Clin Microbiol Infect Dis*. 2014;33(12):2267-74.
- .214 Nie K, Zhang Y, Gan R, Wang L, Zhao J, Huang Z, et al. Polymorphisms in immune/inflammatory cytokine genes are related to Parkinson's disease with cognitive impairment in the Han Chinese population. *Neurosci Lett*. 2013;541:111-5.

- .215 Slattery ML, Herrick JS, Torres-Mejia G, John EM, Giuliano AR, Hines LM, et al. Genetic variants in interleukin genes are associated with breast cancer risk and survival in a genetically admixed population: the Breast Cancer Health Disparities Study. *Carcinogenesis*. 2014;35(8):1750-9.
- .216 Zhou Z, Li X, Li H, Guo M, Liu S, Li C. Genetic Analysis of IL-17 Gene Polymorphisms in Gout in a Male Chinese Han Population. *PLoS One*. 2016;11(2):e0148082.
- .217 Bao MH, Luo HQ, Xiang J, Tang L, Dong LP, Li GY, et al. Meta-Analysis for the Association between Polymorphisms in Interleukin-17A and Risk of Coronary Artery Disease. *Int J Environ Res Public Health*. 2016;13.(7)
- .218 Cheng S, Shao Z, Liu X, Guo L, Zhang X, Na Q, et al. Interleukin 17A polymorphism elevates gene expression and is associated with increased risk of nonsmall cell lung cancer. *DNA Cell Biol*. 2015;34(1):63-8.
- .219 Rasouli M, Asaei S, Kalani M, Kiany S, Moravej A. Interleukin-17A genetic variants can confer resistance to brucellosis in Iranian population. *Cytokine*. 2013;61(1):297-303.
- .220 Silva LK, Blanton RE, Parrado AR, Melo PS, Morato VG, Reis EA, et al. Dengue hemorrhagic fever is associated with polymorphisms in JAK1. *Eur J Hum Genet*. 2010;18(11):1221-7.
- .221 Lan NT, Hirayama K. Host genetic susceptibility to severe dengue infection. *Trop Med Health*. 2011;39(4 Suppl):73-81.
- .222 Hou S, Qi J, Zhang Q, Liao D, Li Q, Hu K, et al. Genetic variants in the JAK1 gene confer higher risk of Behcet's disease with ocular involvement in Han Chinese. *Hum Genet*. 2013;132(9):1049-58.
- .223 Hu K, Hou S, Li F, Xiang Q, Kijlstra A, Yang P. JAK1, but not JAK2 and STAT3, confers susceptibility to Vogt-Koyanagi-Harada (VKH) syndrome in a Han Chinese population. *Invest Ophthalmol Vis Sci*. 2013;54(5):3360-5.
- .224 Shen Y, Liu Y, Ke X, Kang HY, Hu GH, Hong SL. Association between JAK1 gene polymorphisms and susceptibility to allergic rhinitis. *Asian Pac J Allergy Immunol*. 2016;34(2):124-9.
- .225 Sayed KS, El-Komy MHM, Shehata H, ElShazly SH, El Desouky ED, Amr KS, et al. JAK1 rs310241 and JAK3 rs3008 Genotypes May Increase Susceptibility to Psoriasis: A Case Control Study. *Skin Pharmacol Physiol*. 2020;33(4):207-12.
- .226 Miao L, Wang L, Yuan H, Hang D, Zhu L, Du J, et al. MicroRNA-101 polymorphisms and risk of head and neck squamous cell carcinoma in a Chinese population. *Tumour Biol*. 74-4169:(3)37;2016 .
- .227 Chen C, Zhang X, Wang Y. Analysis of JAK2 and STAT3 polymorphisms in patients with ankylosing spondylitis in Chinese Han population. *Clin Immunol*. 2010;136(3):442-6.
- .228 Macedo LC, Santos BC, Pagliarini-e-Silva S, Pagnano KB, Rodrigues C, Quintero FC, et al. JAK2 46/1 haplotype is associated with JAK2 V617F--positive myeloproliferative neoplasms in Brazilian patients. *Int J Lab Hematol*. 2015;37(5):654-60.
- .229 Hsiao HH, Liu YC, Tsai HJ, Lee CP, Hsu JF, Lin SF. JAK2V617F mutation is associated with special alleles in essential thrombocythemia. *Leuk Lymphoma*. 2011;52(3):478-82.
- .230 Sarlos P, Kovessi E, Magyari L, Banfai Z, Szabo A, Javorhazy A, et al. Genetic update on inflammatory factors in ulcerative colitis: Review of the current literature. *World J Gastrointest Pathophysiol*. 2014;5(3):304-21.
- .231 Koh SP, Yip SP, Lee KK, Chan CC, Lau SM, Kho CS, et al. Genetic association between germline JAK2 polymorphisms and myeloproliferative neoplasms in Hong Kong Chinese population: a case-control study. *BMC Genet*. 2014;15:147.
- .232 Pardanani A, Fridley BL, Lasho TL, Gilliland DG, Tefferi A. Host genetic variation contributes to phenotypic diversity in myeloproliferative disorders. *Blood*. 2008;111(5):2785-9.
- .233 Bank S, Andersen PS, Burisch J, Pedersen N, Roug S, Galsgaard J, et al. Genetically determined high activity of IL-12 and IL-18 in ulcerative colitis and TLR5 in Crohns disease were associated with non-response to anti-TNF therapy. *Pharmacogenomics J*. 2018;18(1):87-97.
- .234 Anelli L, Zagaria A, Specchia G, Albano F. The JAK2 GGCC (46/1) Haplotype in Myeloproliferative Neoplasms: Causal or Random? *Int J Mol Sci*. 2018;19.(4)

- .235 Nahajevszky S, Andrikovics H, Batai A, Adam E, Bors A, Csomor J, et al. The prognostic impact of germline 46/1 haplotype of Janus kinase 2 in cytogenetically normal acute myeloid leukemia. *Haematologica*. 2011;96(11):1613-8.
- .236 Colaizzo D, Tiscia GL, Bafunno V, Amitrano L, Vergura P, Grandone E, et al. The JAK2 rs12343867 CC genotype frequently occurs in patients with splanchnic venous thrombosis without the JAK2V617F mutation: a retrospective study. *J Thromb Haemost*. 2010;8(2):413-6.
- .237 Messaoudi S, Hizem S, Al-Sulaiti MA, Al-Busaidi AS, Magdoud K, Dendana M, et al. Contribution of JAK2 and STAT3 variants to the genetic susceptibility of recurrent spontaneous miscarriage in a Tunisian population. *Genet Test Mol Biomarkers*. 2013;17(1):35-9.
- .238 Yang L, Liu D, Liang S, Guo R, Zhang Z, Xu H, et al. Janus kinase 2 polymorphisms are associated with risk in patients with gastric cancer in a Chinese population. *PLoS One*. 2013;8(5):e64628.
- .239 Lee JJ, Wu X, Hildebrandt MA, Yang H, Khuri FR, Kim E, et al. Global assessment of genetic variation influencing response to retinoid chemoprevention in head and neck cancer patients. *Cancer Prev Res (Phila)*. 2011;4(2):185-93.
- .240 Kim SY, Hur MS, Choi BG, Kim MJ, Lee YW, Choe YB, et al. A preliminary study of new single polymorphisms in the T helper type 17 pathway for psoriasis in the Korean population. *Clin Exp Immunol*. 2017;187(2):251-8.
- .241 Slattery ML, Lundgreen A, Kadlubar SA, Bondurant KL, Wolff RK. JAK/STAT/SOCS-signaling pathway and colon and rectal cancer. *Mol Carcinog*. 2013;52(2):155-66.
- .242 Ohyashiki JH, Yoneta M, Hisatomi H, Iwabuchi T, Umezumi T, Ohyashiki K. The C allele of JAK2 rs4495487 is an additional candidate locus that contributes to myeloproliferative neoplasm predisposition in the Japanese population. *BMC Med Genet*. 2012;13:6.
- .243 Chan CHT, Munusamy P, Loke SY, Koh GL, Wong ESY, Law HY, et al. Identification of Novel Breast Cancer Risk Loci. *Cancer Res*. 2017;77(19):5428-37.
- .244 Zhong Y, Wu J, Ma R, Cao H, Wang Z, Ding J, et al. Association of Janus kinase 2 (JAK2) polymorphisms with acute leukemia susceptibility. *Int J Lab Hematol*. 2012;34:53-248:(3)
- .245 Pritchard AL, Johansson PA, Nathan V, Howlie M, Symmons J, Palmer JM, et al. Germline mutations in candidate predisposition genes in individuals with cutaneous melanoma and at least two independent additional primary cancers. *PLoS One*. 2014;9(13):e0194098.
- .246 Núñez-Marrero A, Arroyo N, Godoy L, Rahman MZ, Matta JL, Dutil J. SNPs in the interleukin-12 signaling pathway are associated with breast cancer risk in Puerto Rican women. *Oncotarget*. 2020;11(37):3420-31.
- .247 Saadi A, Dang J, Shan S, Ladjouze-Rezig A, Lefkir-Tafiani S, Gong Y, et al. Ankylosing spondylitis: analysis of gene-gene interactions between IL-12 β , JAK2, and STAT3 in Han Chinese and Algerian cohorts. *Cent Eur J Immunol*. 2019;44(1):65-74.

Author Contributions

Zahra Saadatian and Saba Seyedi designed the study.

Saba Seyedi, Ziba Nariman-Saleh-Fam and Shadan Navid, Samira Ezi and Lida Nariman-Saleh-Fam carried out the search.

Zahra Saadatian and Saba Seyedi wrote the manuscript with support from Samira Ezi and Shadan Navid.

Mohammadreza Gerami drew the Figures.

Zahra Saadatian, Lida Nariman-Saleh-Fam and Ziba Nariman-Saleh-Fam revised the first draft.

All authors commented on the manuscript.

Acknowledgements

This study was approved by the ethics committee of the infectious diseases center, Gonabad university of medical sciences, Gonabad, Iran.

IR.GMU.REC.1401.139

This study received no funding.

Competing Interests

All authors declare no financial or non-financial competing interests.

Data Availability

Data sharing is not applicable to this article as no datasets were generated or analysed during the current study.

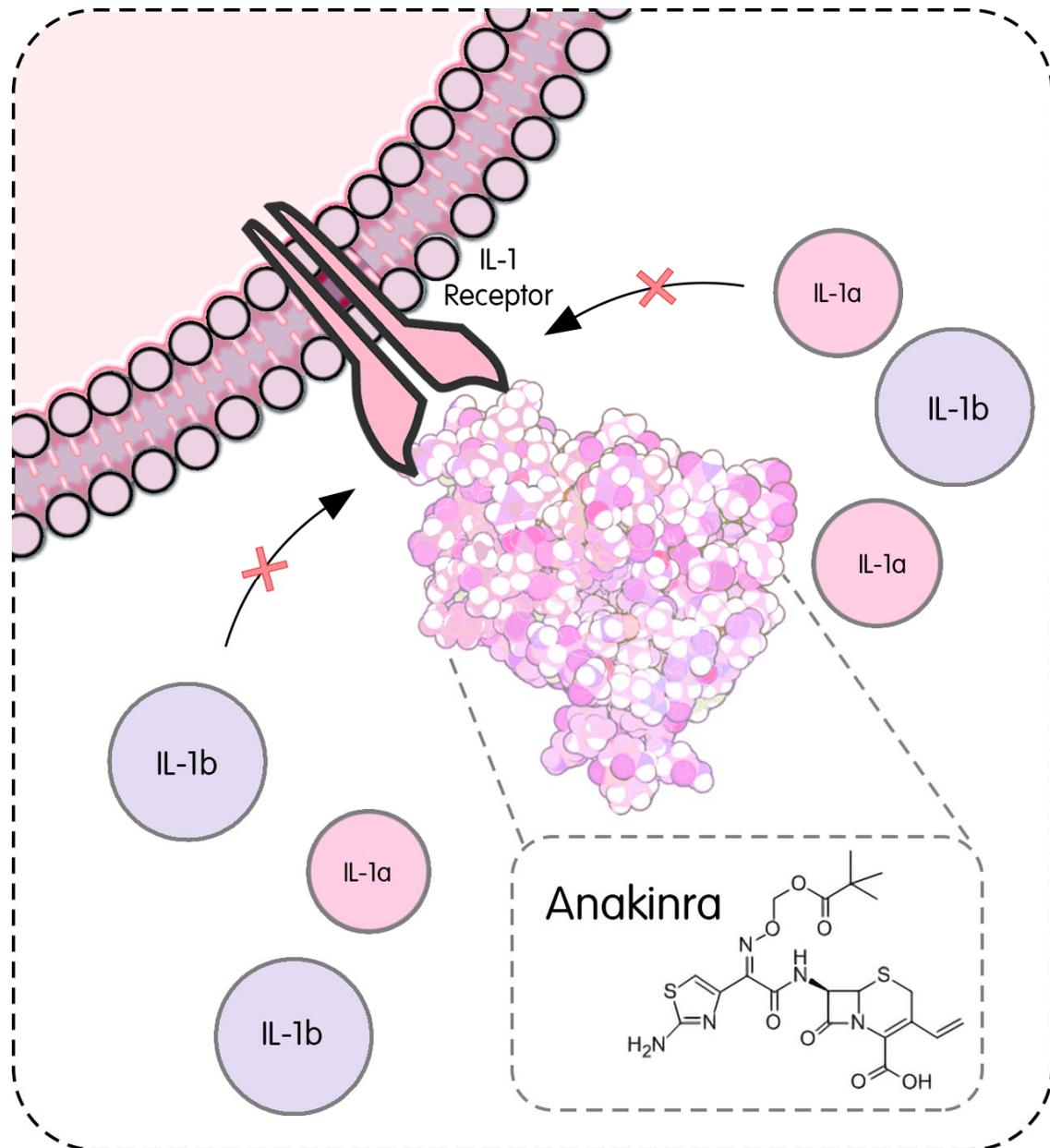


Figure 1: Anakinra

Anakinra as an interleukin-1 receptor antagonist, blocks the activity of IL-1 α and IL-1 β and inhibits binding of the IL-1 to its receptor.

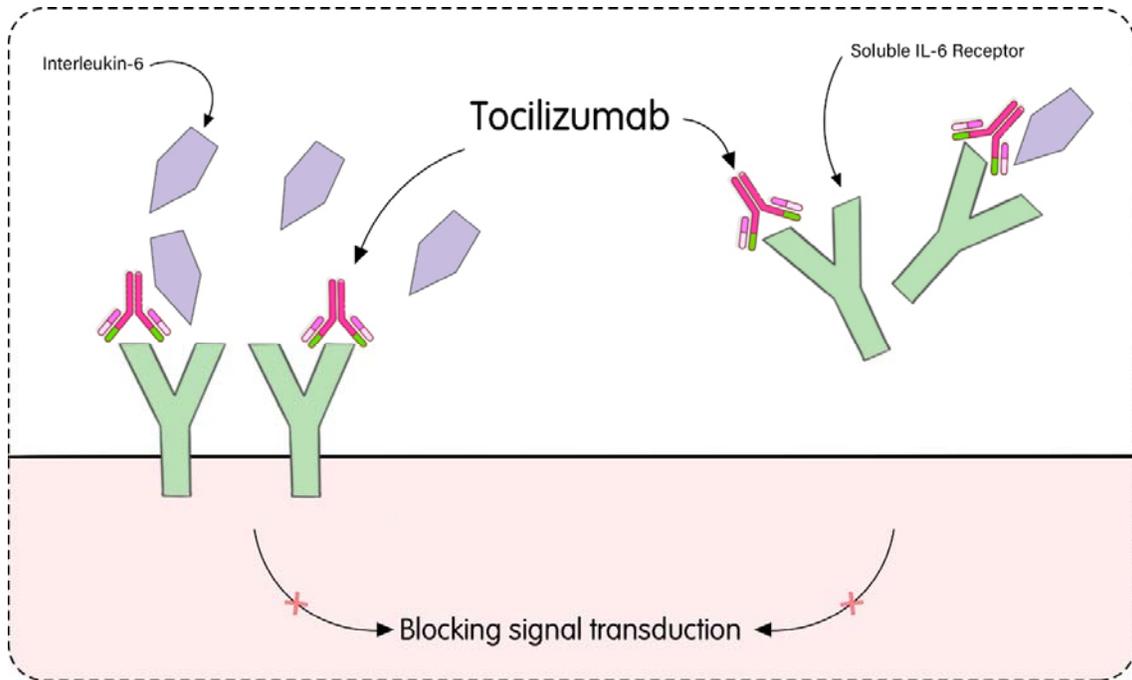


Figure 2: Tocilizumab

Tocilizumab suppresses the cytokine storm by blocking IL-6 receptors.

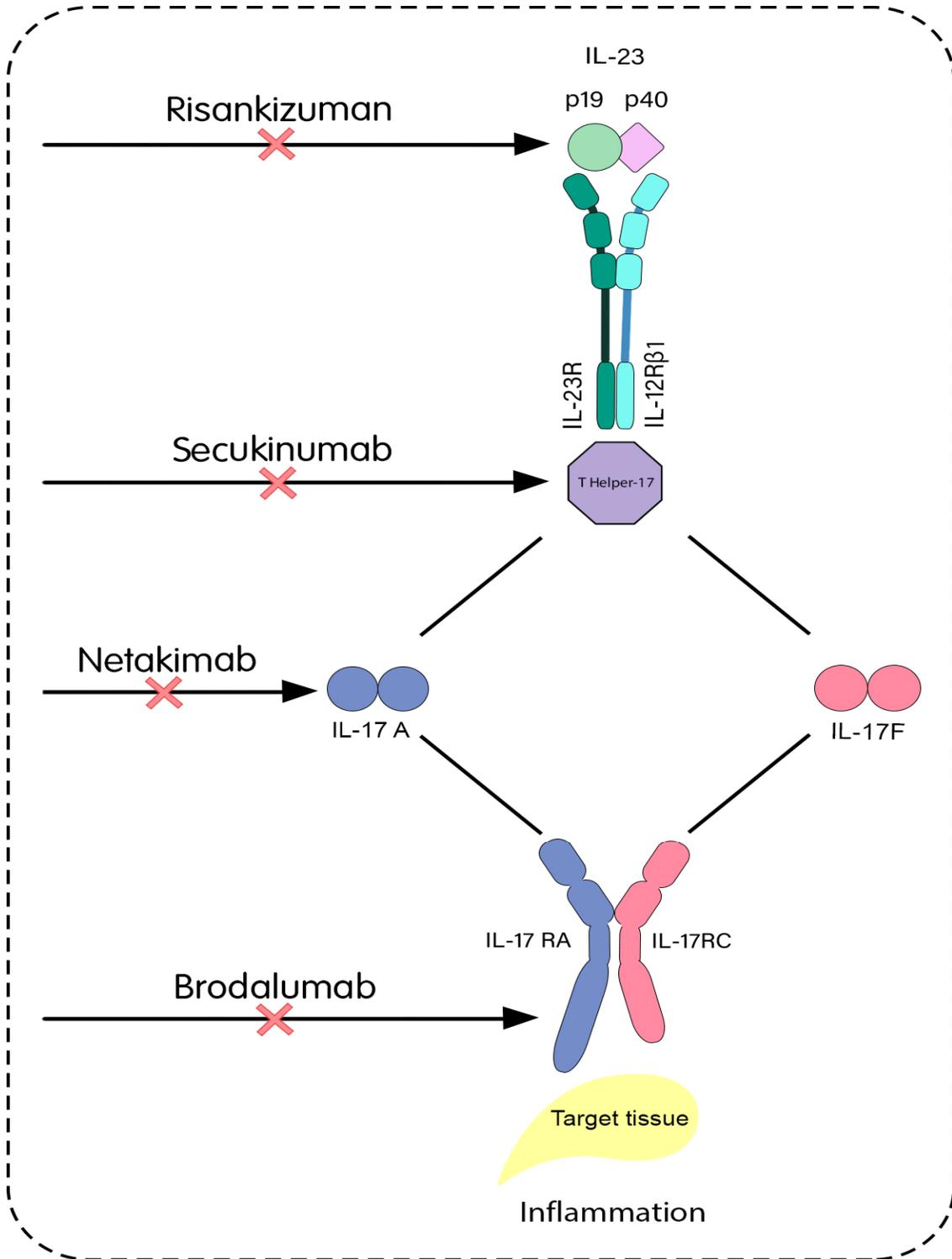


Figure 3: Netakimab

Netakimab, hinders interleukin 17-A, and Brodalumab blocks its receptor of IL-17A (IL-17RA), Secukinumab suppresses T-helper 17, Risankizumab, binds to the p19 and inhibits IL-23.

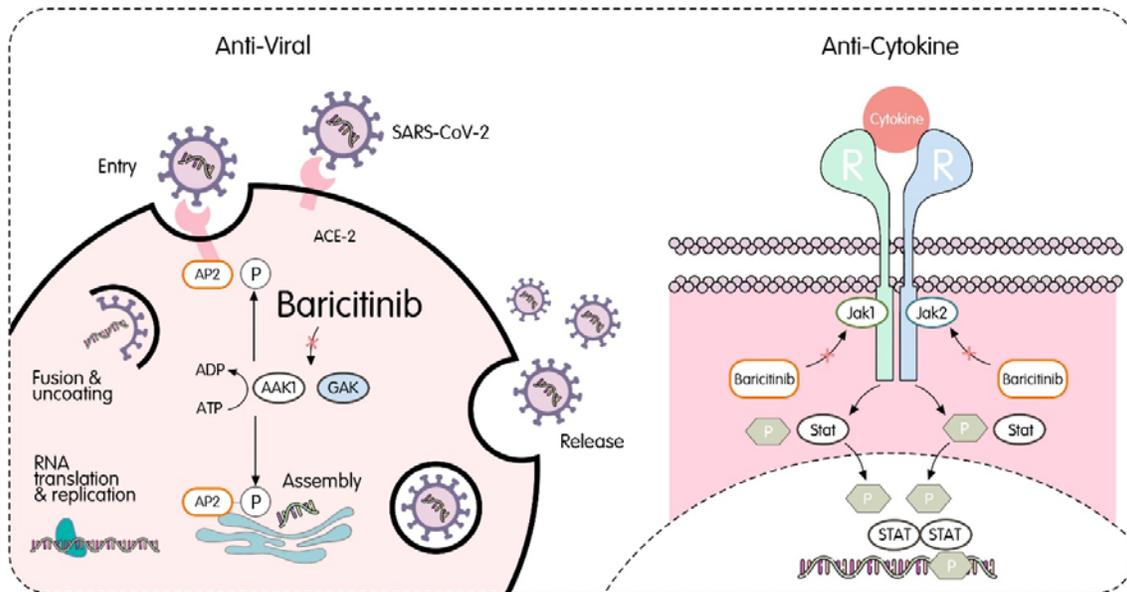


Figure 4: Baricitinib

Baricitinib targets the Janus kinase family, with specificity towards JAK1 and JAK2 and results in the prevention of downstream phosphorylation and activation of Signal Transducers and Activators of Transcription (STAT) proteins. In addition to its anti-inflammatory profile, Baricitinib exhibits antiviral effects by blocking the entry of SARS-CoV-2 into lung cells by reducing the endocytosis of SARS-CoV-2 through the inhibition of AP2-associated protein kinase 1 and cyclin G associated kinase.

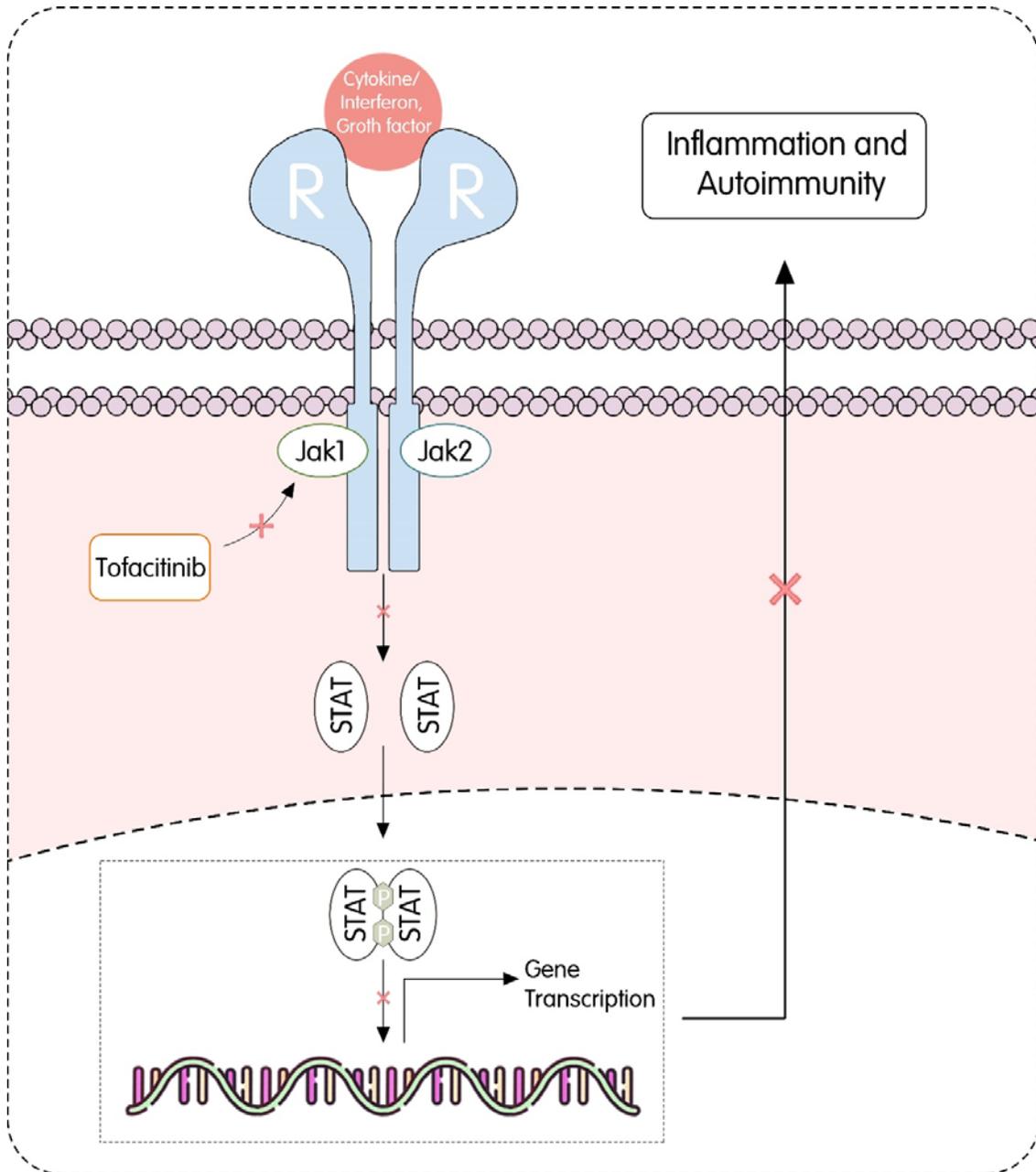


Figure 5: Tofacitinib

Tofacitinib functions by binding to Janus kinases, which inhibits the activation of the JAK-STAT signaling pathway, leading to a potential reduction in the production of pro-inflammatory cytokines.

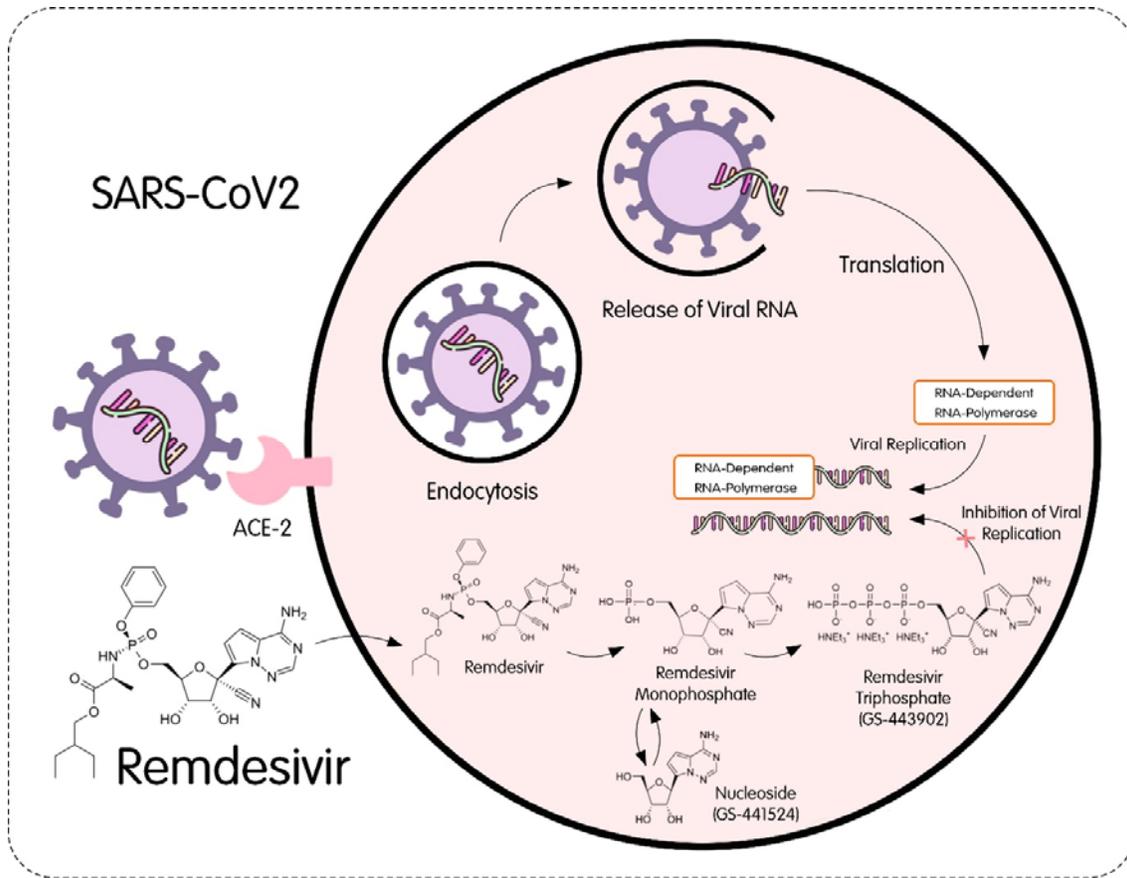


Figure 6: Remdesivir

Remdesivir is metabolized into the pharmacologic active analog adenosine triphosphate and in a competition with ATP interrupts RNA-dependent RNA polymerase (RdRp) function and RNA synthesis.

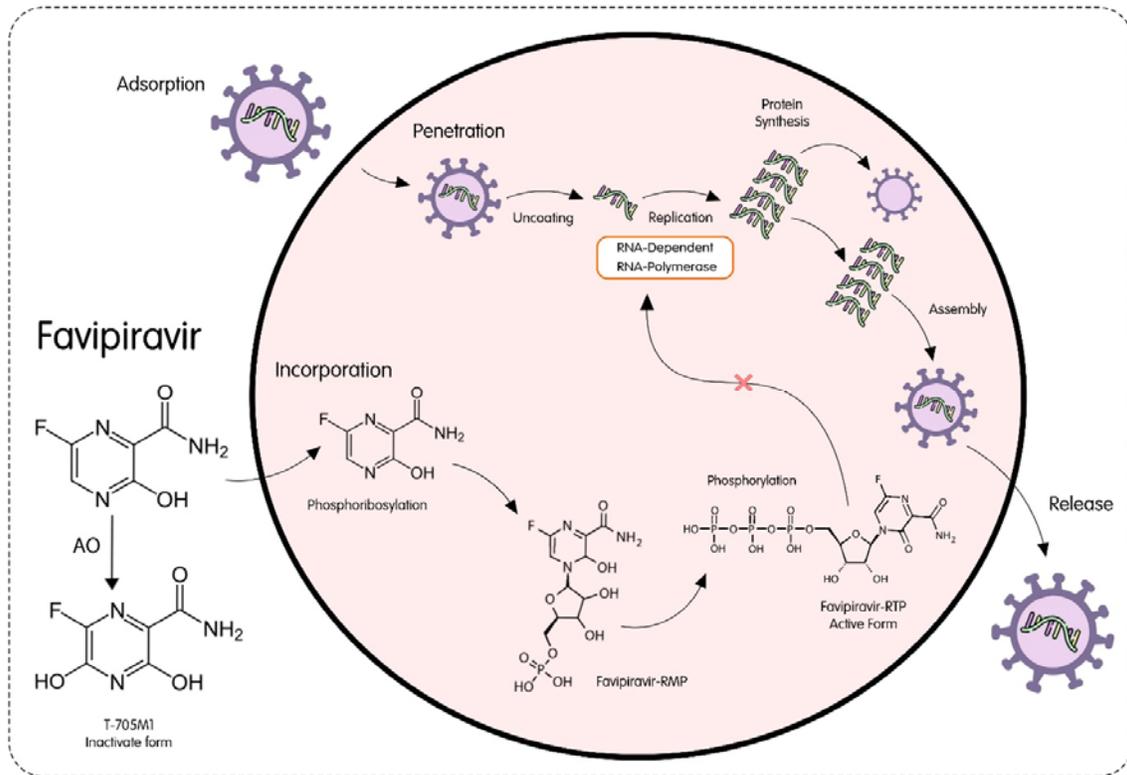


Figure 7: Favipiravir

Favipiravir is metabolized into the active form, ribofuranosyl-5B-triphosphate (favipiravir-RTP) and interferes with RNA-dependent RNA polymerase (RdRp) function and RNA synthesis.

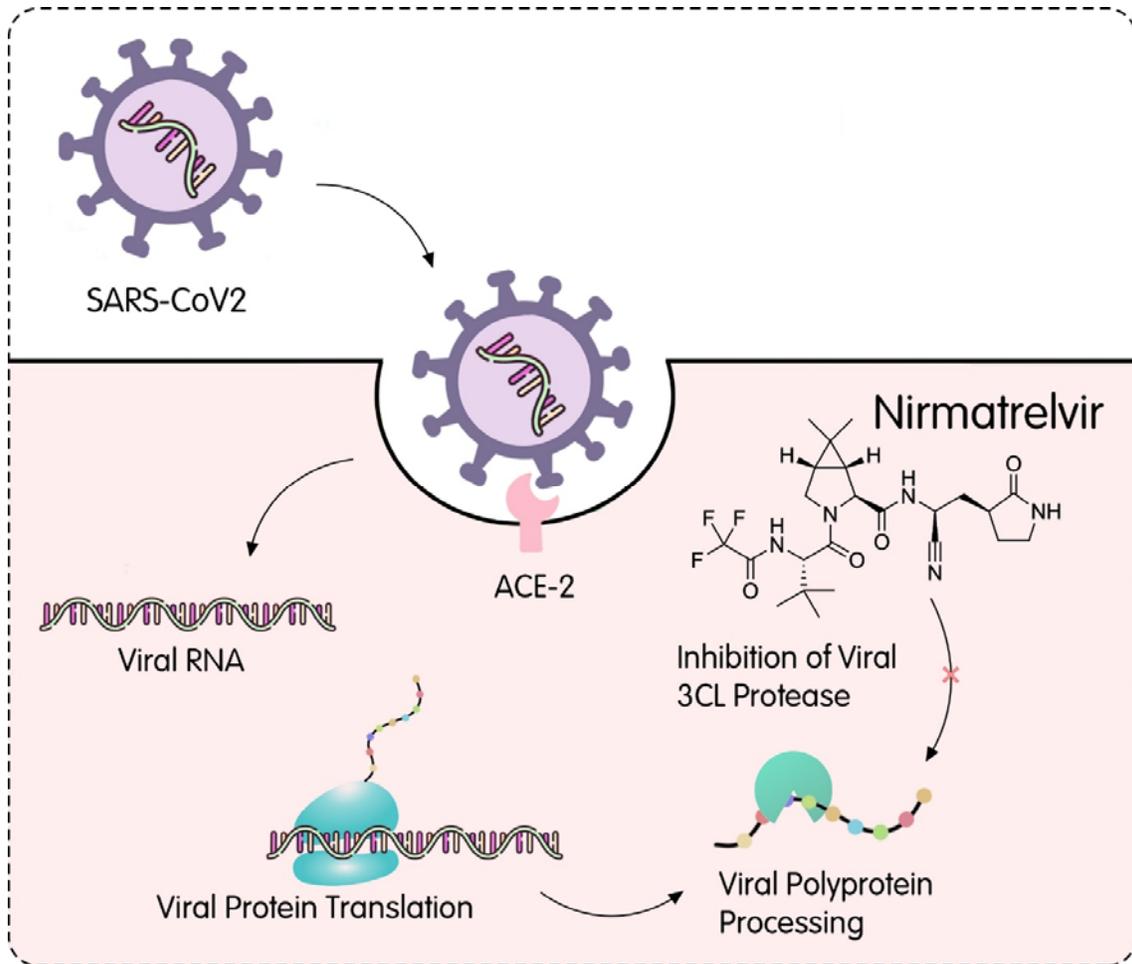


Figure 8: Nirmatrelvir

Nirmatrelvir targets the 3C-like protease enzyme required for COVID-19 viral replication and results in the prevention of virus replication.

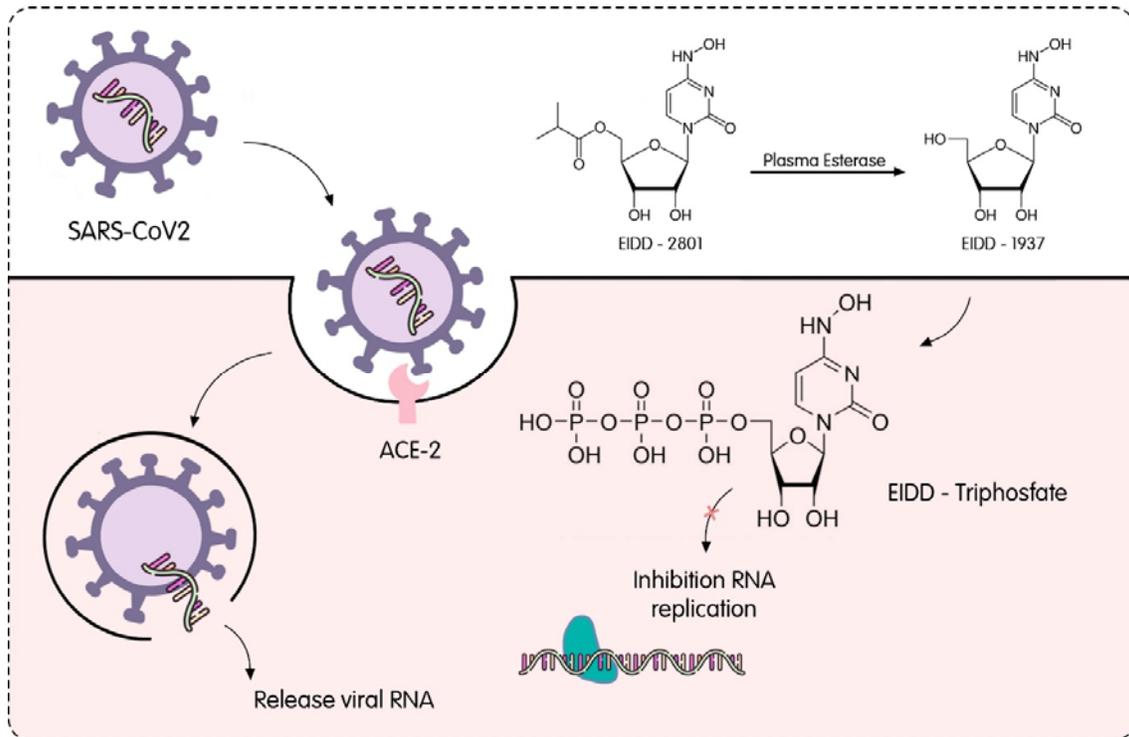


Figure 9: Molnupiravir

Molnupiravir is a prodrug that is a ribonucleoside small molecule derivative known as β -D-N4-hydroxy cytidine (NHC). NHC circulates in the body and undergoes intracellular phosphorylation to form NHC triphosphate. Subsequently, viral RNA polymerase combines with NHC triphosphate to misdirect the virus to bind to either guanosine or adenosine when replicating. This results in the accumulation of destructive mistakes in the viral genome, ultimately rendering the virus non-infectious and unable to replicate.

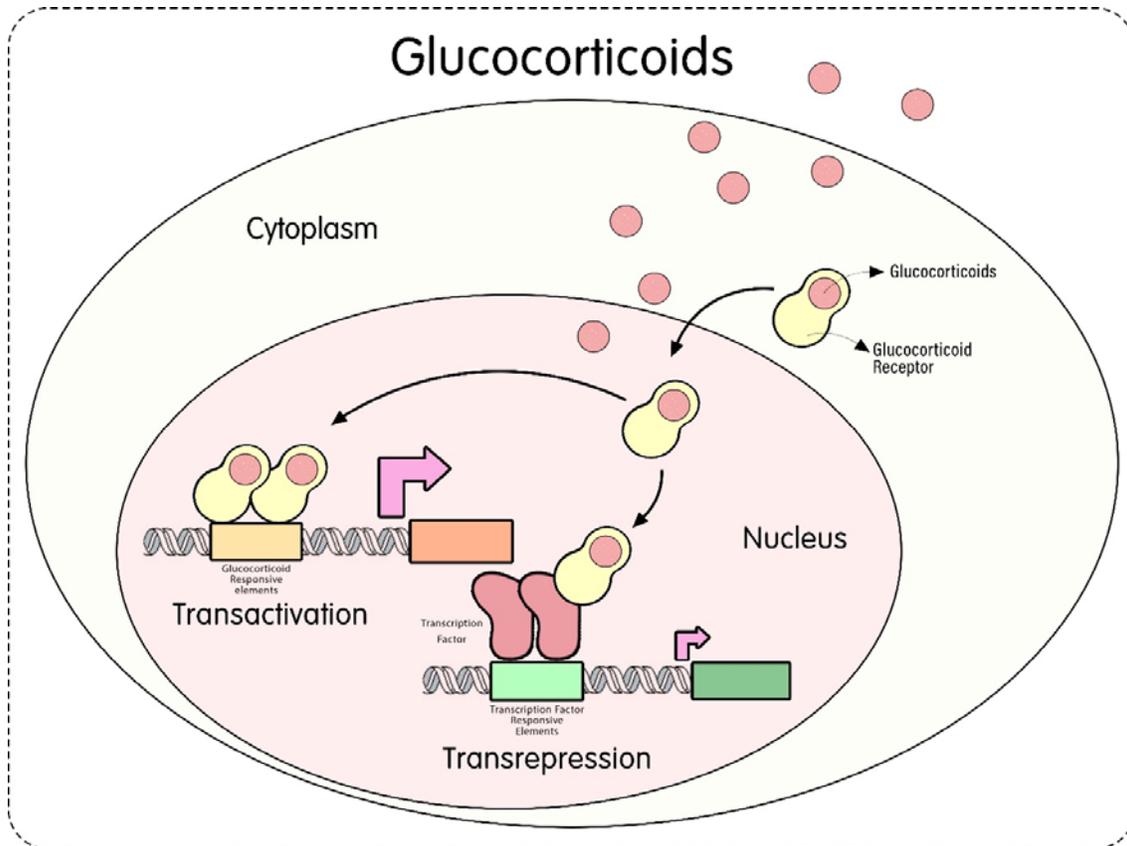


Figure 10: Corticosteroids

Corticosteroids bind with receptor proteins in the cytoplasm and make a steroid-receptor complex. This complex pass into the nucleus and can stimulate or inhibit the synthesis of specific proteins via DNA connection.

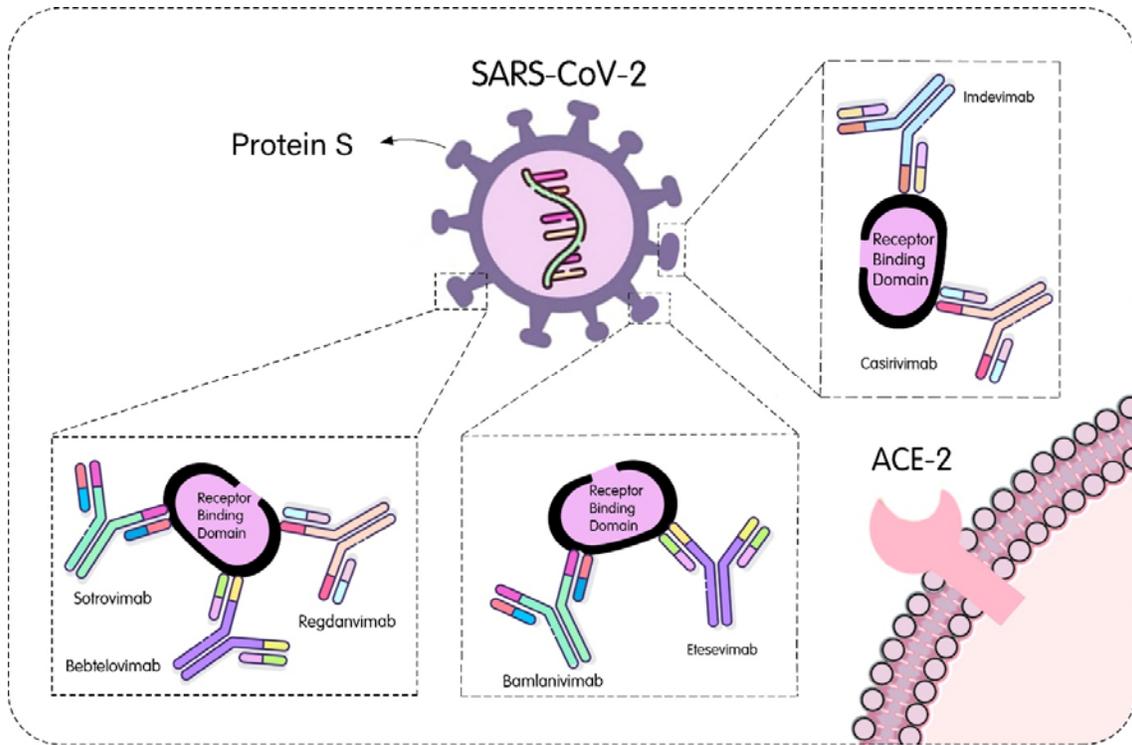


Figure 11: Monoclonal antibodies

Bamlanivimab plus etesevimab, sotrovimab and Casirivimab plus imdevimab target spike protein of SARS-CoV-2.

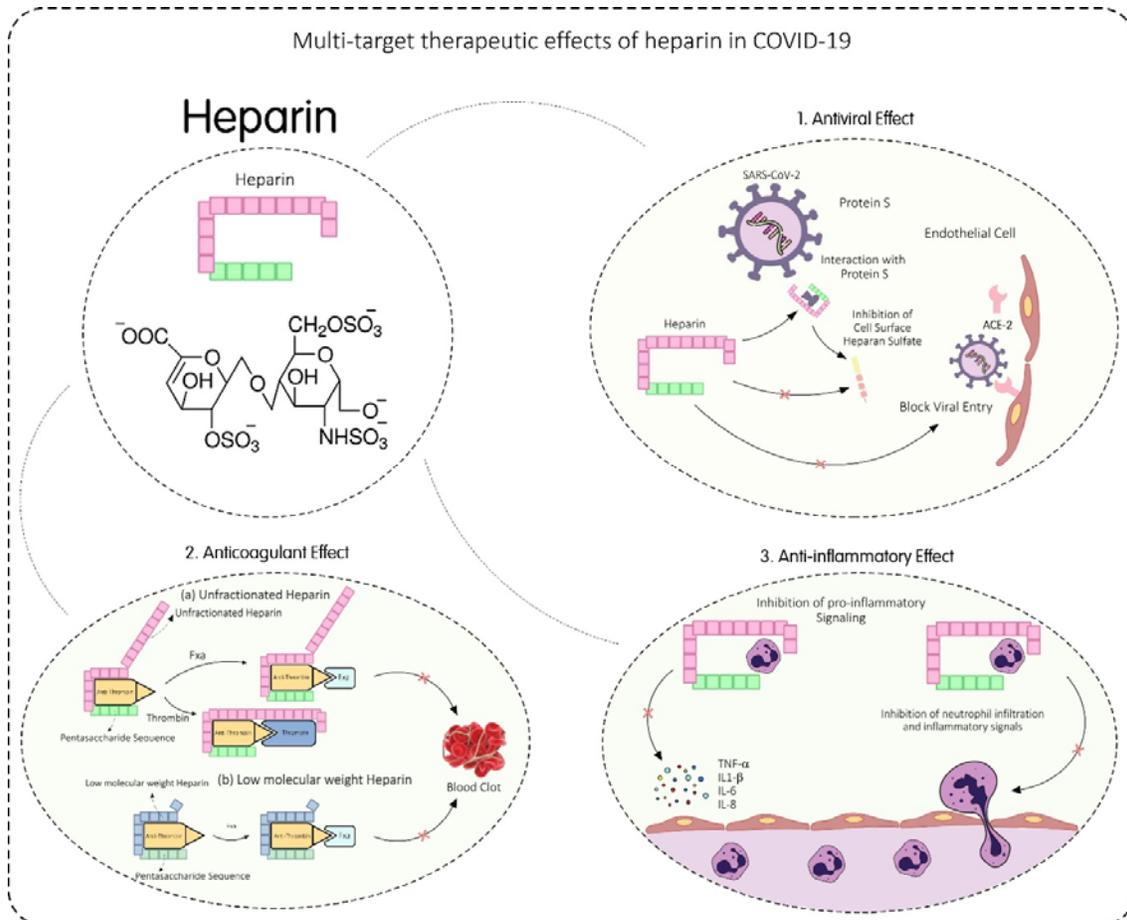


Figure 12: Heparin antiviral (1), anti-inflammatory (2), and anticoagulant (3) properties.

1. Heparin can also attach to the spike protein of SARS-CoV-2 and serve as a competitive suppressor for virus entry, thus reduce infectivity
2. it can reduce inflammation through impeding neutrophil infiltration and inhibiting the production of inflammatory factors, such as IL-8, IL-6, and TNF- α
3. Moreover, both unfractionated heparin (UFH) and low-molecular-weight heparin (LMWH) can bind to antithrombin III which inhibits the activation of FX. Unfractionated heparin (UFH) also can blocks thrombin by consisting heparin-antithrombin-thrombin complex and leading anticoagulation.

Table1: anti-inflammatory drugs used against COVID-19, their direct targets and SNPs, and SNPs association with other disease and drugs

Anti covid medicine	target	SNP	position	SNP association with other drugs	SNP association with diseases	reference
anakinra	IL-1R1	rs10490571	Intron	-	Rheumatoid Arthritis, osteonecrosis of the femoral head, lumbar disc herniation, breast cancer, IgA nephropathy	(75),(76),(77), (78), (79)
		rs12712127	Intron	-	-	-
		rs13019803	Intron	-	GVHD	(80)

	rs1302 0778	Intron	-	No evidence	-
	rs1558 641	Intron	-	atopic asthma	(81)
	rs2110 726	3 Prime UTR	-	Less Breast Pain in Women Prior to Breast Cancer Surgery	(82)
	rs2160 227	Intron	-	invasive pneumoc occal disease	(83)
	rs2192 752	Intron	-	location of papillary thyroid carcinom a	(84)
	rs2234 650	Intron	-	sudden sensorine ural hearing loss, risk of vertical HIV transmiss ion, systemic sclerosis	(85), (86), (87)
	rs2287 047	Intron	-	severe hand osteoarthr	(88), (89)

					itis, Invasive Pneumoc occal Disease	
		rs3917 225	Intron	-	osteoporosis predisposition, osteonecrosis of the femoral head, thyroid cancer, ankylosing spondylitis, IgA nephropathy, severe hand osteoarthritis	(90), (76), (91), (92), (79), (88)
		rs3917 254	Intron	-	invasive pneumoc occal disease	(83)
		rs3917 267	Intron	-	invasive pneumoc occal disease, Hepatitis B Virus Infection	(83), (93)

					of Children	
		rs949963	Intron	-	development of secondary lymphedema following breast cancer surgery, Childhood asthma	(94), (95)
tocilizumab	IL-6R	rs11265611	Intron	-	atrial fibrillation	(96)
		rs12083537	Intron	Tocilizumab(73), (97), (98), (99), (7(74)2)	ischemic stroke in patients with metabolic syndrome, severe heart failure, lung function in Chinese patients with asthma, coronary heart disease,	(100), (101), (102), (103)

		rs12133641		-	Abdominal aortic aneurysm	(104)
		rs12730935		-	No evidence	-
		rs2228144		-	No evidence	
		rs2228145	Misense	Tocilizumab(73), (72)	Multiple disorders such as: rheumatoid arthritis, asthma and dermatitis risk, COVID-19 risk, psoriasis, morbidity of preterm newborns, Alzheimer's Disease, obesity, Gulf War Illness, ischemic stroke, type 2 diabetes, bipolar disorder,	(105), (106), (107), (108), (109), (110), (111), (112), (113), (114), (115), (101)

					heart failure	
		rs2229 238	3 Prime UTR	-	Preeclampsia	(116)
		rs4129 267	Intron		Asthma, Ankylosing Spondylitis, cardiovascular diseases, Diabetic Ischemic Heart Disease, cardiovascular diseases and inflammatory disease	(117), (118), (106), (119), (106)
		rs4537 545	Intron	-	joint damage in rheumatoid arthritis,	(120)
		rs4845 617	5 Prime UTR	-	atherosclerotic lipid profiles, Neurological Status,	(121), (122), (108)

					psoriasis	
		rs4845 618	Intron	-	Melanoma, cardiovascular diseases and inflammatory disease, rheumatoid arthritis	(123), (106), (120)
		rs4845 622	Intron	-	Melanoma, severe heart failure	(123), (101)
		rs4845 625	Intron	-	atrial fibrillation, dyslipidemia, chronic kidney disease, cardiovascular disease, Atherosclerosis, cardiovascular diseases and inflammation	(96), (124), (125), (126), (127), (106), (119)

					tory disease, Diabetic Ischemic Heart Disease	
		rs6684439	Intron	-	Melanoma, multiple myeloma, severe heart failure	(123), (128), (101)
		rs6694817	Intron	-	No evidence provided	-
		rs7529229	Intron	-	Diabetic Ischemic Heart Disease, cardiovascular diseases and inflammatory disease, severe heart failure	(119), (106), (101)
secukimab	IL-17R	rs1057518744	Frame shift	-	-	-
		rs1057518745	Stop Gained	-	-	-

	rs1057 518746	Frame shift	-	-	-
	rs1057 518747	Frame shift	-	-	-
	rs1057 519079	Misse nse	-	-	-
	rs1215 9217	Intron	-	Cerebral Malaria	(129)
	rs2229 151	Synon ymous Vari ant	-	new- Onset Diabetes after Transplan tation	(130)
	rs2241 049	Intron	-	primary graft dysfuncti on after lung transplant ation, psoriatic arthritis	(131),(132)
	rs3879 06913	Stop Gaine d	-	-	-
	rs4139 6547	Intron	-	Cerebral Malaria	(129)
	rs4143 3045	2KB Upstre am Vari ant	-	Cerebral Malaria	(129)

		rs4819554	2KB Upstream Variant	anti-TNF drugs(133), etanercept(134)	Prognosis of Non-Small Cell Lung Cancer Patients Receiving Epidermal Growth Factor Receptor Tyrosine Kinase Inhibitors , Hashimoto's thyroiditis , Preeclampsia, Toxoplasma gondii infection, cardiovascular mortality in patients with congestive heart failure	(135), (136),(137),(138), (139)
		rs879574	Intron	-	chronic rejection after lung transplantation,	(140),(141)

					Crohn's disease	
		rs882643	3 Prime UTR	-	Crohn's disease	(141)
netaki mab	IL - 17 A	rs10484879	intron	etanercept(134), tolerance to FOLFOX chemotherapy in colorectal cancer treatment(142)	Psoriasis, Age-Related Macular Degeneration, Chronic Periodontitis, Risk of Vulvovaginal Candidiasis, chronic hepatitis B virus infection, colorectal cancer	(132), (143), (144), (145), (146), (147)
		rs1974226	3 Prime UTR	-	atopic asthma, Chronic Obstructive Pulmonary Disease, Gram-positive infection and mortality	(148), (149), (150)

					of severe sepsis	
		rs2275913	2KB Upstream	steroid - response(151), Antitumor Necrosis Factor Agents(152), etanercept(134), biologic and MTX treatment(153), anti-TNF therapy (154),	IgA nephropathy, polycystic ovary syndrome, cervical cancer, systemic lupus erythematosus, schizophrenia, colorectal cancer, Rheumatoid arthritis, hepatocellular carcinoma, gastric cancer, Chagas disease, appendicitis, Prostate Cancer, syphilis, influenza, acute respiratory distress syndrome, leprosy	(155), (156), (157), (158),(159), (160), (161),(162), (163), (164), (165), (166), (167), (168), (169), (170), (171),(145), (172), (173), (174), (175),(176), (177), (178),(179), (180), (181),(182), (183),(184), (185),(186), (187)

					<p> type 1 reaction, non- alcoholic fatty liver disease, Vulvovag inal Candidias is, recurrent aphthous ulcer, osteoarthr itis, bladder cancer, a sthma, ischemic stroke, ankylosin g spondyliti s, celiac disease, laryngeal cancer , Recurrent Pregnanc y Loss , severe viral bronchiol itis, lung cancer, Intestinal Behcet's disease, </p>	
--	--	--	--	--	--	--

					breast cancer, periodontitis, panuveitis, Oral squamous cell carcinoma	
		rs3748067	3 Prime UTR	FOLFOX tolerance (142),	breast cancer, gastric cancer risk, Graves' disease, tuberculosis, Pemphigus Foliaceus, asthma, brucellosis, gastric cancer, Cervical Cancer, obstructive sleep apnea syndrome	(184),(188), (189), (190), (191), (192), (193), (194), (195),(196)
		rs3804513	Intron	tolerance to FOLFOX(142)	ulcerative colitis	(197)
		rs3819024	2KB Upstre	-	gastric cardia	(198),(199), (200),

			am Variant		adenocarcinoma, metabolic syndrome, allergic bronchopulmonary aspergillosis in asthma, periodontitis and type 2 diabetes, brucellosis, colorectal cancer, syphilis, rheumatoid arthritis	(201),(193), (147), (147), (202),
		rs3819025	Intron	cytokine inhibitors(133)	breast cancer, Henoch-Schonlein purpura, brucellosis, rheumatoid arthritis	(184), (203),(193),(204)
		rs4711998	2KB Upstream Variant	response to blinatumomab (205), sustained responses to PEG-IFN α -2 α (206)	HBV related hepatocellular carcinoma, brucellosis	(207), (193),(208), (209)

					s, osteitis after BCG vaccination, hepatitis B virus-related liver cirrhosis	
rs7747909	3 Prime UTR	cytokine inhibitors(133)			colorectal cancer, age-related macular degeneration, giant cell arteritis	(147),(143), (210)
rs8193036	2KB Upstream Variant	Anti-Jo1 Antisynthetase Autoantibodies(211)			acute respiratory distress syndrome (ARDS), osteitis after Bacillus Calmette-Guerin vaccination, metabolic syndrome, allergic Rhinitis, restless legs syndrome, skin	(169), (208), (199),(199),(212),(213),(214), (183),(150),(215),(216)

					structure infections , Parkinson's disease with cognitive impairment, intestinal Behcet's disease, Gram-positive infection, breast cancer mortality, rheumatoid arthritis	
		rs8193037	2KB Upstream Variant	cytokine inhibitors.(133)	Graves' disease, Coronary Artery Disease, congestive heart failure, nonsmall cell lung cancer, ulcerative colitis	(189),(217), (139),(218),(197)
		rs8193038	Intron	-	ulcerative colitis, brucellosis	(197),(219)

barcit inib	JA K1	rs1057 519753	Misse nse	-		
		rs1078 9166	intron	-	Dengue hemorrhagic fever	(220, 221)
		rs1120 8534	Intron	-	-	-
		rs1510 47872	missen se	-	-	-
		rs1712 7114	Intron	-	-	-
		rs2254 002	Intron	-	-	-
		rs2780 815	Intron	-	Behcet's disease	(222)
		rs2780 831	Intron	-	Dengue hemorrhagic fever	(220)
		rs2780 885	Intron	-	-	-
		rs2780 889	3 Prime UTR	-	-	-
		rs2780 890	intron	-	-	-
		rs2780 895	Intron	-	-	-
		rs3102 30	Intron	-	Vogt- Koyanagi -Harada (VKH) syndrom	(223)

		rs3102 36	Intron	-	Vogt- Koyanagi -Harada (VKH) syndrome	(223)
		rs3102 41	Intron	-	Behcet's disease with ocular involvement, Vogt- Koyanagi -Harada (VKH) syndrome , allergic rhinitis, Psoriasis	(222), (223), (224), (225)
		rs3102 45	Intron	-	-	-
		rs3790 532	Intron	-	Behcet's disease with ocular involvement	(222)
		rs3806 277	Intron	-	-	-
		rs4786 65	Intron	-	-	-

		rs4916008	Intron	-	-	-
		rs578481	Intron	-	head and neck squamous cell carcinoma	(226)
		rs705509	Intron	-	head and neck squamous cell carcinoma	(226)
		rs869312953	Misense	-	-	-
tofctinib	JA K2	rs10119004	Intron	-	ankylosing spondylitis	(227)
		rs1057519721	Misense	-	-	-
		rs1057519722	missense	-	-	-
		rs1057519723	missense	-	-	-
		rs1057520016	missense	-	-	-
		rs10815144	Intron	-	-	-
		rs10815149	Intron	-	-	-
		rs10974944	Intron	-	myeloproliferative	(228), (229), (230),

					neoplasms, essential thromboc ythemia	
		rs1219 12472	missense	-	-	-
		rs1219 12473	missense	-	-	-
		rs1234 0895	Intron	-	myeloproliferative neoplasms	(231)
		rs1234 2421	Intron	-	polycythemia vera, myeloproliferative neoplasms	(232), (231)
		rs1234 3867	Intron	anti-TNF response(233)	Myeloproliferative Neoplasms, cytogenetically normal acute myeloid leukemia	(234), (235)
		rs1234 9785	Intron	-	myeloproliferative neoplasms, splanchnic venous	(231), (236)

					thrombosis	
		rs2230722	Synonym variant	-	-	-
		rs2230724	Synonymous Variant	-	recurrent spontaneous miscarriage, gastric cancer, acute myeloid leukemia	(237), (238), (235),
		rs2274471	Intron	13- <i>cis</i> -retinoic acid (13-cRA)(239)	Psoriasis, colon and rectal cancer	(240), (241),
		rs368927897	missense	-	-	-
		rs3780374	Intron	-	-	-
		rs3780378	Intron	-	myeloproliferative neoplasms	(231)
		rs4495487	Intron	-	myeloproliferative neoplasm	(242)
		rs56118985	Missense	-	breast cancer, acute leukemia	(243), (244)

		rs7737 5493	Mis se nse	-	myelopro liferative	(245),
		rs7849 191	Intron	-	breast cancer, myelopro liferative neoplasm s	(246), (231)
		rs7851 969	Intron	-	-	-
		rs7857 730	Intron	-	Algerian, myelopro liferative neoplasm s, ankylosin g spondyliti s	(247), (231), (227)

