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Review

Repurposing anticancer drugs for the management of COVID-19



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Abstract Since its outbreak in the last December, coronavirus disease 2019 (COVID-19) caused by SARS-CoV-2 has rapidly spread worldwide at a pandemic proportion and thus is regarded as a global public health emergency. The existing therapeutic options for COVID-19 beyond the intensive supportive care are limited, with an undefined or modest efficacy reported so far. Drug repurposing represents an enthusiastic mechanism to use approved drugs outside the scope of their original indication and accelerate the discovery of new therapeutic options. With the emergence of COVID-19, drug repurposing has been largely applied for early clinical testing. In this review, we discuss some repurposed anticancer drugs for the treatment of COVID-19, which are under investigation in clinical trials or proposed for the clinical testing.

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1. Introduction

Novel coronavirus disease 2019 (COVID-19) has recently emerged as a potential life-threatening disease of public health interest, worldwide. Few pharmacotherapeutic options are available at the present time, and patients are experiencing adverse outcomes in the most severe presentations. The treatments used in the clinical care are mainly based on supportive measures and intensive care management for complicated cases [138]. Antiviral, antibiotic and antimalarial molecules have also been used to avoid severe outcomes as well as plasma therapy from recovered patients [12,19,75]. Despite the lack of strong evidence from clinical trials, several pharmacological treatments have been used in clinical practice and included in possible protocols based on preliminary findings of promising efficacy and a rationale against COVID-19, as suggested by recent systematic reviews [37,217]. However, if available, clinical trials remain the option of choice for patients with COVID-19, to enhance the data and progress in medical research. In this context, drug repurposing can be an attractive and promising approach for patients with COVID-19 in order to benefit from the safety and effectiveness of the molecules with proposed antiviral and/or immunomodulating properties [80,142]. Repurposed pipelines may have a significant impact, particularly in this critical time [44,78,161]. In this review article, we discuss the potential of a number of compounds licensed for cancer conditions that seem to have a rationale to target some pathogenic key mechanisms of COVID-19, under clinical investigation or proposed for clinical trials.

2. Drug repurposing as a substitute solution for emerging viral diseases of public health interest

The development of novel antiviral drugs requires long-term investigations in clinical trials. The advantage to repurpose drugs to validate off-label use is related to the known safety profile, though it may vary across the diseases, and consolidated data of pharmacodynamics, pharmacokinetics and efficacy in phase I–IV trials. Of interest, some host cellular targets interfering with the viral growth cycle, such as kinases, are broadly shared in the mechanisms of several viral infections and other conditions such as cancer, suggesting the possibility to translate knowledge across medical disciplines and disease models [56,156]. Therefore, pharmacologic approaches for targeting key pathogenic mechanisms, such as signalling pathways, constitute a promising tactic during outbreaks of emerging pathogens. A cost-effective drug repurposing has been previously used to combat intracellular pathogens as an urgent alternative during the occurrence of rapidly spreading and deadly infectious diseases: in fact, repurposed medicines may

help reduce the investments in the earlier phases of drug development, with an immediate use in the specific setting of unmet need [66,118]. For instance, an accumulated experience on repurposed therapeutics during previous outbreaks caused by Ebola virus, dengue virus, severe acute respiratory syndrome coronavirus (SARS-CoV), Zika virus and Middle Eastern respiratory syndrome coronavirus (MERS-CoV) has led to the establishment of a rationale for synergistic drug combinations [9,20,32,47,213]. For example, the use of sofosbuvir plus ribavirin treatment, which is a combination of a nucleotide polymerase inhibitor and a synthetic antiviral nucleoside analogue approved for hepatitis C infection by the Food and Drug Administration (FDA) [187], has been repurposed for treating Zika virus infection. The two antiviral drugs were all tested *in vivo* for Zika disease and showed promising efficacy [22,57,160], thus prompting the clinical testing [16]. The antiprotozoal nitazoxanide developed by Romark Laboratories® is another important example [1,6]. Nitazoxanide is a thiazolide that triggers the host immune response by potentiating interferon release, interfering with post-translational stage of viral haemagglutinin and intracellular trafficking [157,191]. This drug showed potential properties with its broad-spectrum effects on other microbes encompassing *Mycobacterium leprae* [10], human norovirus [40] and Ebola virus [95]. Additionally, it has also been repurposed for influenza and is being tested in late-phase clinical trials ([79] (for details, see: NCT03336619, NCT02612922, NCT01227421 and NCT01610245). Meanwhile, nitazoxanide is presently under investigation in two randomised clinical trials (NCT04343248 and NCT04341493) for COVID-19. Romark® is a phase III, randomised, double-blind, placebo-controlled trial evaluating the efficacy and safety of nitazoxanide (twice daily for 6 weeks) for post-exposure prophylaxis of COVID-19 in 300 patients (NCT04343248). A phase IV trial is already currently enrolling patients with COVID-19 and will compare nitazoxanide-hydroxychloroquine to hydroxychloroquine alone (NCT04341493). Hydroxychloroquine is an antimalarial drug that has been shown to interfere with the late-stage entry process of viruses based on cellular endocytic pathways. It is currently being investigated worldwide in more than 70 urgent clinical trials for COVID-19 (<https://www.clinicaltrials.gov>; accessed 14-04-2020). However, emerging evidence from a randomised and controlled phase III trial comparing hydroxychloroquine alone or with azithromycin versus standard care treatments for hospitalised patients with mild to moderate disease did not show clinically significant improvements after 15 d of treatment [25]. In the real world, similar results were obtained in an observational study with a larger cohort [68]. Notably, the latest meta-analysis with Bayesian examination of 29 studies including three randomised trials on this topic found evidence against

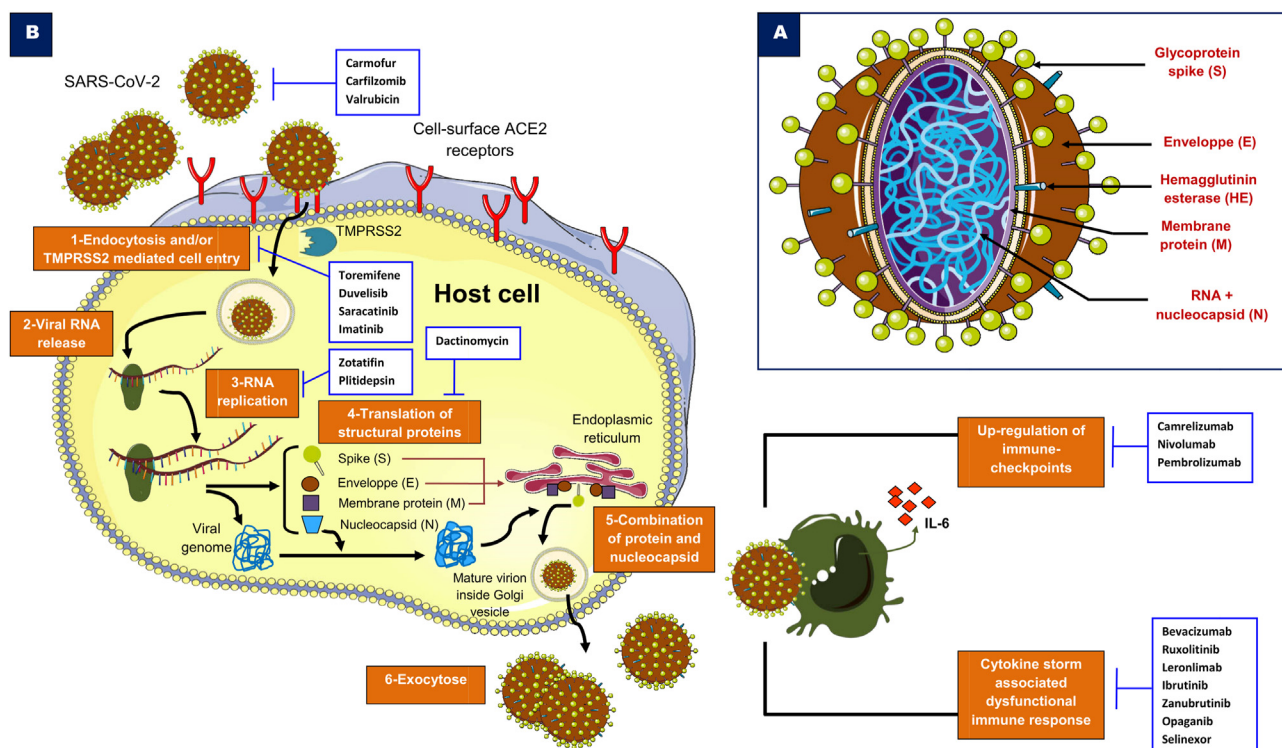


Fig. 1. A: SARS-CoV-2 structure. B: A simplified illustration of the targetable pathways by anticancer drugs in COVID-19. For details, see text. Abbreviations: ACE2: angiotensin-converting enzyme 2, IL-6: interleukin 6, RNA: ribonucleic acid, TMPRSS2: transmembrane protease serine 2.

the use of hydroxychloroquine alone or in combination with azithromycin in this setting [58].

3. Druggable targets in COVID-19-host signalling pathways

COVID-19 exhibits mild to moderate symptoms, which may progress into severe pneumonia, with lymphopenia, and an exhausted cytokine release syndrome [24]. The disease is caused by a novel strain of coronaviruses (SARS-CoV-2) that emerged in November–December 2019, probably naturally selected in an animal host before zoonotic transfer. However, SARS-CoV-2 has then crossed the animal species barriers and acquired the capacity of human-to-human infection, resulting in a rapid spread of COVID-19 at a pandemic proportion [83].

The SARS-CoV-2 is closely related to the previous strain SARS-CoV-1 that has caused the SARS epidemic in China during the period 2002–2004 [216]. SARS-CoV-2 is an enveloped, non-segmented and single-stranded RNA virus and belongs to the sarbecovirus subgenus of the *Coronaviridae* family [226]. The ultra-structure of SARS-CoV-2 is characterised by four key proteins, namely surface spike S glycoprotein, envelope protein E, membrane protein (M) and nucleocapsid protein (N) (Fig. 1A).

The mechanisms of infectivity and pathogenesis of this coronavirus are poorly known and seem to be

similar to those of SARS-CoV and MERS-CoV [106] (Fig. 1B). The host cell entry of SARS-CoV-2 depends on angiotensin-converting enzyme 2 (ACE2) and transmembrane protease serine 2 (TMPRSS2) [86,205,207]. Human cells with rich cell-surface receptor ACE2, an important modulator involved in the blood pressure regulation, are susceptible to SARS-CoV-2 infection. According to the recently emerged evidence, the elucidated crystal structure of SARS-CoV-2 showed that its receptor-binding domain (RBD) has an enhanced affinity to bind to the human ACE2 [167]. The receptor-binding motif of the RBD mediates the contact with ACE2. Two virus-binding hotspots at the RBD–ACE2 interface are stabilised by several residue changes of amino-acids in SARS-CoV-2 as compared to the previous SARS-CoV-1 [167]. The entry into cells requires the protein S of the viral spike that binds to the receptor ACE2 facilitated by some host cell proteases, such as TMPRSS2 [71,173]. This phenomenon, based on protein S priming, enables fusion of viral and cellular membranes and escapes from antiviral humoral immune response [71,173]. This interaction seems to be targetable by inhibiting host proteases [86,169].

The marketed orally bioavailable camostat mesylate approved in Japan for the treatment of chronic pancreatitis and reflux oesophagitis after gastrectomy [101,123] demonstrated a significant inhibition of TMPRSS2 [86], suggesting a potential role to treat COVID-19. These effects were also observed in prostate

cancer in which TMPRSS2 inhibition by camostat was associated with suppression of metastasis *in vivo* [109]. In addition, the use of soluble human ACE2 can inhibit early stage SARS-CoV-2 infections as revealed by a recent study including human organoids, by using the recombinant ACE2 as a decoy receptor [121]. Fig. 1B shows an overview about the potential targets for COVID-19 therapy.

After the fusion of viral and host cell membranes, the virus accesses by the endosomal pathway, facilitated by cathepsins (cysteine pH-dependent proteases) [18,144]. The virus disassembles and releases its nucleocapsid and RNA material into the host cell cytoplasm via caveolae- and clathrin-mediated endocytosis [215]. This pathway involves phosphatidylinositol binding clathrin assembly protein (PICALM), which is a cargo-selecting clathrin adaptor that drives membrane curvature during endocytosis [90,119]. The PICALM is a target of chloroquine that suppresses its expression [90,209], inhibiting in this way the clathrin-mediated endocytosis process. The genomic RNA of SARS-CoV-2 contains open reading frames, which are translated into two replicase polyproteins (pp1a and pp1b) [55]. Therefore, the replicase transcriptase complex (RTC) is generated by processing pp1a and pp1b to produce non-structural proteins (16 putative NSPs), which form RNA polymerase and helicase that have a central role in the transcription and replication of coronaviruses [26,195]. The RTC attaches to the endoplasmic reticulum (ER) membranes and produces double-membrane vesicles of the coronavirus. A part of the RTC undergoes other molecular changes for mRNA synthesis and splicing to generate the structural and accessory proteins of the virus. The glycoproteins of the envelope will reach the ER and Golgi apparatus in which they form with the nucleocapsid protein and the viral RNA the vesicle containing the virion [26]. After assembly via the ER–Golgi intermediate compartment, the release of the new virion-containing vesicles from the infected host cells is achieved by exocytosis [26,31]. An alternative cell surface non-endosomal route for coronavirus entry involves TMPRSS2 that cleaves the spike glycoprotein into S1 and S2 subunits [26]. This enables a fusion of the membranes of the coronavirus and target cell. Of note, TMPRSS2 is druggable by camostat that demonstrated efficacy as a single treatment in lung cell entry blockade of MERS-CoV [172].

The SARS-CoV-2 action on host immunity encompasses innate and adaptive immune response activation. Unfortunately, local and systemic uncontrolled inflammatory response as well as altered adaptive immune reactions can damage organ tissues [24]. Leukopenia and lymphopenia with reduced natural killer (NK) cells in addition to increased C-reactive protein can be seen in patients with severe COVID-19 [24,27]. A marked decrease of total CD8⁺ T cells and NK translates the early failure of antiviral immunity during SARS-CoV-2 infection [223]. Upregulation of the expression of CD94/NK group 2 member A (NKG2A) by these immune cells

suggested a functional exhaustion of cytotoxic lymphocytes in the development of COVID-19 [223]. These features can result in high mortality and more severity of the disease [27,91,92,146].

Elevated serum levels of pro-inflammatory cytokines (known as cytokine release syndrome or cytokine storm), such as tumour-necrosis factor alpha (TNF- α) and interleukin 6 (IL-6), seem to cause multiple organ failure by damaging tissues, contributing to the poor outcomes in COVID-19 patients [14,24,70,117,159,210,218,220]. Notably, immunosuppression obtained with drugs targeting IL-6 axis such as tocilizumab (Actemra®, Genentech) or siltuximab (Sylvant®, Janssen) has been proposed [61,117,134,221]. Based on these assumptions, several phase III trials are currently ongoing (NCT04320615, NCT04345445 and NCT04330638). However, at least for tocilizumab, the clinical use in hospitalised patients with severe COVID-19 has not met the expectations in randomised clinical trials (COVACTA/NCT04320615), and areas of implementation are now explored to identify possible predictive biomarkers of benefit for a subset of selected patients [124]. Finally, the pro-inflammatory microenvironment in COVID-19-associated lung failure induces massive infiltration of immune cells and damages the alveolar tissues. This mechanism increases vascular permeability and alveolar fluids accumulation, which is a hallmark of the deadly acute respiratory distress syndrome (ARDS) [115]. Both the mechanisms of immune-escape in COVID-19 might be potentially targetable and should be urgently delineated.

4. Potential anticancer drugs for COVID-19

4.1. Ruxolitinib

Ruxolitinib is a potent orally bioavailable Janus kinase (JAK) 1 and JAK2 inhibitor indicated for various myeloproliferative malignancies including myelofibrosis and polycythemia vera [3,198,200]. These neoplasms, alongside with the acute myeloid leukaemia, are known by their aberrant activation of the JAK-STAT signalling [199]. This pathway plays a pivotal role in cell proliferation and survival and seems to be targetable for a spectrum of conditions beyond malignancies.

Regarding COVID-19, ruxolitinib was recently repurposed to quell the immune-hyperactivation, thus dampening the cytokine storm and improving ARDS. An artificial intelligence-driven study has identified ruxolitinib as a promising drug for COVID-19 along with other molecules capable to disrupt the JAK-STAT signalling, used for myelofibrosis and rheumatoid arthritis, such as baricitinib and fedratinib [181]. The clinical trials using ruxolitinib for COVID-19 are summarised in Table 1. In particular, ruxolitinib is being investigated against placebo for COVID-19 patients with ARDS in two randomised phase III trials (NCT04362137 and NCT04377620). The

Table 1
Summary of potential anticancer drugs for repurposing in COVID-19.

Anticancer drug	Mechanism of action	Viral–host targets	Tested in clinical trials for COVID-19? (country)	NCT identifier	Eligible population	Primary end-point	Estimated completion date
Ruxolitinib	Reduction of hyperinflammation during cytokine storm	JAK-STAT pathway	Yes (USA, Germany, France, Mexico, Canada, Italy and Spain)	NCT04348071	Hospitalised patients	Cumulative incidence of Grade 3 and 4 AE; eight-point ordinal scale	October 2021
				NCT04359290	Severe lung disease	Overall survival	July 2021
				NCT04355793	Hospitalised patients	Description of the disease course	Not provided
				NCT04354714	Critical disease	Overall survival	Withdrawn
				NCT04362137	Hospitalised patients	Proportion of patients who die, develop respiratory failure or require ICU care	November 2020
				NCT04377620	COVID-19-associated ARDS	Overall survival	October 2020
				NCT04366232	Hypoxic pneumonia, ARDS or end-stage organ failure	Biological criteria of response, based on biomarkers change*	September 2020
				NCT04334044	Pneumonia with dyspnea	Recovery of pneumonia	June 2020
				NCT04374149	Critical disease	Overall response rate	April 2021
				NCT04338958	Severe lung disease	Overall response rate	August 2021
				NCT04337359	Critical disease	Description of the disease course	Not provided
				NCT04331665	Disease requiring O ₂ -support	Rate of clinical worsening	February 2021
				NCT04361903	Disease requiring O ₂ -support	Number of patients who avoid mechanical-assisted ventilation	June 2020
				NCT04348695	Hospitalised patients	Rate of clinical worsening	May 2020
				NCT04424056	• Eligible patients for resuscitation care in UCI	Ventilation-free days at day 28	November 2022
Bevacizumab	Vascular permeability inhibition	VEGF	Yes (France and China)	NCT04275414	Severe lung disease or critical disease	Change of PaO ₂ to FiO ₂ ratio	May 2020
				NCT04344782	Severe lung disease	Number of patients who avoid mechanical-assisted ventilation	December 2020
				NCT04305106	Disease requiring O ₂ -support	Time to clinical improvement	August 2020
Carmofur	Blockade of viral replication	SARS-CoV-2 main protease	No	—	—	—	—
Carfilzomib	Blockade of viral replication	SARS-CoV-2 main protease	No	—	—	—	—

Toremifene	Inhibition of viral membranes fusion with host cell endosomes	Interaction with coronavirus proteins	No	—	—	—	—
Zotatifin	Inhibition of protein biogenesis	Blockade of eIF4A	No	—	—	—	—
Plitidepsin	Interference with the viral cycle	Blockade of eEF1A	Yes (Spain)	NCT04382066	Hospitalised patients	Frequency of occurrence of Grade 3 or higher AEs	November 2020
Dactinomycin	Inhibition of viral cellular transcription	Inhibition of DNA topoisomerase	No	—	—	—	—
Valrubicin	Blockade of viral replication	SARS-CoV-2 main protease	No	—	—	—	—
Leronlimab	Immune homeostasis restoration	Disruption of the CCL5/RANTES-CCR5 pathway	Yes (USA)	NCT04343651 NCT04347239	Mild/moderate disease Severe lung disease or critical disease	Clinical improvement Overall survival	September 2020 April 2021
Camrelizumab	Immune homeostasis restoration	PD-1/PD-L1 pathway blockade	Yes (China)	ChiCTR2000029806	Severe lung disease	Proportion of patients with a lung injury score reduction	February 2021
Nivolumab	Immune homeostasis restoration	PD-1/PD-L1 pathway blockade	Yes (France and China)	NCT04343144	Disease requiring O ₂ -support	Time to clinical improvement	October 2020
				NCT04333914	Disease requiring O ₂ -support	28-day survival rate	Suspended
				NCT04356508	Clinically stable patients with mild or moderate disease and asymptomatic patients	Viral clearance kinetics	September 2021
Pembrolizumab	Immune homeostasis restoration	PD-1/PD-L1 pathway blockade	Yes (Spain)	NCT04335305	Severe lung disease or critical disease	Percentage of patients with normalisation of SpO ₂ ≥96% on room air	September 2020
Imatinib	Blockade of cell entry and endosomal trafficking	BCR/ABL kinase inhibition	Yes (France, Spain and USA)	NCT04357613	Hospitalised patients	Rate of prevented severe disease worsening	December 2021
				NCT04346147	Disease diagnosed <7 d	Time to clinical improvement	September 2020
				NCT04394416	Hospitalised patients diagnosed <7 d	All-cause mortality	June 2023
				NCT04356495	Outpatients, diagnosed <5 d	Hospitalisation rate, death rate	February 2021
Duvelisib	Immune homeostasis restoration and viral replication inhibition	PI3K inhibition	Yes (USA)	NCT04372602	Critical disease	Overall survival	April 2022
Zanubrutinib	Protection against immune, lethal and sepsis-induced pulmonary injuries	Inhibition of the Bruton tyrosine kinase	Yes (USA)	NCT04382586	Disease requiring O ₂ -support	Respiratory failure-free survival rate	April 2021
Ibrutinib	Protection against immune-induced lung injury	Inhibition of the Bruton tyrosine kinase	Yes (USA)	NCT04375397	Hospitalised patients with severe pneumonia	Respiratory failure-free survival rate, overall survival	April 2021

(continued on next page)

Table 1 (continued)

Anticancer drug	Mechanism of action	Viral–host targets	Tested in clinical trials for COVID-19? (country)	NCT identifier	Eligible population	Primary end-point	Estimated completion date
Opaganib	Anti-inflammatory and antiviral properties	Inhibition of sphingosine kinase-2	Yes (Israel)	NCT04414618	Disease requiring O ₂ -support	Measurement of the daily O ₂ requirements	January 2021
Saracatinib	Antiviral properties	Possible blockade of endocytic pathways and nucleocapsid protein	No	–	–	–	–
Selinexor	Antiviral and anti-inflammatory properties	Blockade of nucleocytoplasmic transport	Yes (USA, France, Austria, Spain and United Kingdom)	NCT04355676	Hospitalised patients with moderate or severe disease	Percentage of participants with at least a two-point improvement in the ordinal scale	September 2020
				NCT04349098	Hospitalised patients with severe disease	Idem	Idem

Abbreviations: AE: adverse event, FiO₂: fraction of inspiration O₂, ICU: intensive care unit, O₂: oxygen, PaO₂: partial arterial oxygen pressure, VEGF: vascular endothelial growth factor. * C-reactive protein, ferritinemia, creatininemia, transaminases, eosinophil count and lymphocyte count.

RUXCOVID study (n = 402, NCT04362137) is a phase III, multicentred, randomised (2:1) and placebo-controlled trial, which evaluates the safety and efficacy of ruxolitinib added to best supportive care in patients with cytokine storm. The RUXCOVID-DEVENT (NCT04377620) is a phase III, placebo-controlled trial, which is currently evaluating ruxolitinib for the treatment of COVID-19 patients with ARDS requiring mechanical ventilation. Both the phase III trials are designed for clinically critical patients, aiming to tackle the pathological immune response and control the hyper-inflammatory syndrome in more severe patients. In addition, another ongoing phase III trial (NCT04424056) combines multiple immune suppressors including anakinra (an anti-IL-1 inhibitor), tocilizumab, in association with ruxolitinib for more severe COVID-19 cases, including ARDS, associated with possible cytokine storm and other immune dysregulations.

4.2. Carmofur

Carmofur (1-hexylcarbamoil-5-fluorouracil) is an old pyrimidine analogue and a lipophilic masked compound of fluorouracil. Its mechanism of action is independent of 5-fluorouracil-antitumour activity, based on the inhibition of the lysosomal acid ceramidase [43,148]. The upregulation of acid ceramidase is linked to survival and growth of cancer cells as well as drug resistance [23]. This enzyme is an emerging target for several malignancies [74], such as prostate cancer [108] and glioblastoma [130] as well as paediatric brain tumours [45]. In addition, carmofur exerts its antiproliferative and antimetastatic effects through the blockade of the Wnt/β-catenin signalling pathway [107]. This molecule has been previously tested in clinical trials for the treatment of colon cancer as single or in combination with other chemotherapeutics [162] and in early breast cancer [122]. Recently, carmofur has been found to block SARS-CoV-2 replication by inhibiting the viral protease through the covalent binding between the carbonyl reactive group and the catalytic Cys145. In another molecular modelling report, Gao *et al.* indicated that carmofur strongly bound to SARS-CoV-2 M^{pro} [63]. This suggests that repurposing low-molecular weight drugs such as carmofur against the main protease of SARS-CoV-2 based on *in silico* studies may show a therapeutic potential. Importantly, the identification of the specific mechanisms of viral blockade provided the paradigm to design analogues of carmofur, with improved efficacy for COVID-19 and less neurotoxicity. This last aspect is particularly relevant because of its association with drug-related leukoencephalopathy.

4.3. Bevacizumab

Bevacizumab is a monoclonal antibody with anti-angiogenic properties, capable to target the vascular endothelial growth factor [64]. This biological agent has

been successfully implemented in the clinic management of various cancer types, including advanced ovarian cancer, renal cell carcinoma as well as colorectal cancer in combination with chemotherapy [49,64]. The key mechanism of bevacizumab is related to the disruption of the malignant neo-angiogenesis, ultimately reducing the abnormal vascular permeability and nutrients supply to cancer cells. Based on this mechanism [15,208], the use of bevacizumab has been proposed for the control of the abnormal angiogenesis and the related vascular permeability undermining the accumulation of intra-alveolar fibrin deposition, resulting in ARDS and occurring during COVID-19 disease. A recent study found that new vessel growth induced by intussusceptive angiogenesis in the lungs of COVID-19 patients is approximately three times higher than in the lungs of patients with influenza [2]. Therefore, bevacizumab is being investigated also in this field (NCT04344782, NCT04305106 and NCT04275414). More in detail, the BEST-CP clinical trial (NCT04275414) (n = 20) is a Chinese pilot study that is currently recruiting patients with COVID-19 to be treated with intravenous bevacizumab (500 mg flat dose) with the aim to control the pulmonary oedema. Another multicentred and randomised trial in China is investigating the safety and efficacy of bevacizumab (7.5 mg/kg) in COVID-19 patients with severe pneumonia (NCT04305106). Additionally, the French CORIMMUNO-BEVA phase II trial (NCT04344782) is randomising COVID-19 patients to receive bevacizumab (7.5 mg/kg) versus best of the standard supportive care (Table 1). Based on the proposed mechanism of action, the use of bevacizumab is now included in clinical trials for more severe patients, presenting with the clinical features of ARDS.

4.4. Carfilzomib

Carfilzomib is an approved antineoplastic drug for the treatment of relapsed or refractory multiple myeloma patients who have received ≤ 3 previous lines of therapies [170,183,214]. It has been investigated in several clinical trials for other malignancies such as Hodgkin lymphoma (NCT02867618) and non-Hodgkin lymphoma like in diffuse large B-cell lymphoma (NCT02073097) and mantle cell lymphoma (NCT03891355). In detail, carfilzomib is a second-generation antiproteasome with higher maximal percentage of proteasome inhibition at the maximum tolerated dose than bortezomib [103]. This mechanism leads to the irreversible inhibition of the 26S proteasome, which plays a key role in the hallmarks of myeloma, by binding to its chymotrypsin-like site of the proteolytic core [62,227]. The multiple targets of proteasome targeting encompass the NF- κ B pathway, ER stress, cell-cycle arrest, downregulation of growth factor receptors and adhesion molecules expression, angiogenesis inhibition as well as immunogenic cell death

[103]. Carfilzomib was recently repurposed for COVID-19 based on virtual docking screening, a structure-based virtual screening method for molecules capable of binding specific proteins in key binding sites or domains [206]. Among several other promising drugs, including eravacycline, elbasvir, lopinavir and valrubicin, carfilzomib was excellent in terms of binding free energy, which is an important parameter for a rational drug design targeting the SARS-CoV-2 main protease, anticipating a potent antiviral activity *in silico* [206].

4.5. Toremifene

Toremifene is a selective modulator of oestrogen receptors used in the endocrine therapy of breast cancer since 1997 [201]. Toremifene represents an alternative drug to tamoxifen, showing a similar efficacy in patients with advanced breast cancer [113]. Additionally, this antioestrogen drug was shown to destabilise the envelope glycoprotein of Ebola virus [222], leading to the inhibition of the fusion between viral and endosome membranes during intracellular trafficking [222]. The subsequent *in vivo* studies confirmed the anti-Ebola activity of toremifene regardless of its oestrogen-modulating activity, by inhibiting virus entry and internalisation [97].

Finally, a recent *in silico* study identified toremifene as a potential therapeutic approach against SARS-CoV-2, particularly when combined to emodin [225]. Emodin is a natural-derived compound with anticoronavirus pharmacological properties, belonging to the traditional Chinese medicine *Lianhua Qingwen* used for the prevention and treatment of viral influenza [84,165]. The authors suggested that toremifene interacts with several key host coronavirus proteins including ribosomal proteins like RPL19, HNRNPA1, NPM1, EIF3I, EIF3F and EIF3E [225]. However, the exact mechanism of its activity is still unclear. Based on the action on viral entry, the use of this agent can be proposed for prophylaxis of the infection and in patients with milder disease, in the earlier stages of infection.

4.6. Zotatfin

Zotatifin is a potent and selective small molecule inhibiting the eukaryotic initiation factor 4A (eIF4A) [190]. eIF4A is an RNA helicase activated by the B-cell receptor to stimulate several oncogenes linked to tumour growth, cell survival and metastasis [82,190]. The inhibition of this target by zotatifin and other synthetic molecules demonstrated promising antineoplastic preclinical efficacy [28,125,190]. For example, zotatifin is being studied in a phase I/II dose-escalation and cohort expansion clinical trial (n = 45) as monotherapy for pancreatic cancer and other malignancies (NCT04092673). Promisingly, a recent study published in *Nature* suggests a potential of zotatifin repurposing for COVID-19 [73]. Based on affinity-purification mass

spectrometry, the authors revealed 332 SARS-CoV-2-human protein–protein interactions after cloning and expressing the key SARS-CoV-2 proteins in human cells [73]. Importantly, 66 targetable human proteins were identified to be pharmacologically druggable by 69 compounds including 29 FDA-approved drugs [73]. Several of these compounds exhibited antiviral activities against mRNA translation and Sigma1 and Sigma2 virus receptors expressed on host cells [73]. In this study, zotatifin emerged as an effective drug in reducing viral infectivity by inhibiting protein biogenesis through eIF4A blockade along with other compounds such as ternatin-4 [73,135]. The proposed disruption of the translation of the Sigma proteins coupled to the direct inhibitory activity against them seems a promising strategy of drug targeting against SARS-CoV-2 infection and propagation, possibly indicated in the earlier stages of the infection.

4.7. Plitidepsin

Plitidepsin (also known as dehydrodidemnin B) is a natural compound discovered in a Caribbean marine organism (*Aplidium albicans*) [48]. Its antitumour effects encompass several cancer targets such as angiogenesis and cell survival [5,72]. Additionally, plitidepsin may also enhance radiotherapy response by a bystander effect [155]. The activation of c-Jun N-terminal kinase is a key mediator of plitidepsin sensitivity along with the eukaryotic elongation factor 1 alpha 2 that were recently suggested as potential predictive biomarkers of drug response [50,126]. So far, the drug has been investigated in several clinical trials for multiple myeloma, lymphoma, liposarcoma and prostate cancer [48]. Only one phase III randomised and controlled trial has evaluated the clinical activity of plitidepsin combined to dexamethasone in relapsed/refractory multiple myeloma (NCT01102426/ADMYRE), showing mild improvements in survival outcomes [180]. The development of this drug has been halted or terminated because of the arrival of competitive targeted agents as well as the poor enrolment in clinical trials.

On the other hand, plitidepsin has also antiviral properties, showing promising findings in the COVID-19 treatment on *in vitro* models (Press release 1, [136,137]). According to these models, plitidepsin blocks the eEF1A protein in human cells that is required for SARS-CoV-2 infection (Press release 2, [136,137]). Therefore, the drug is being tested in a phase I Spanish trial (NCT04382066/APLICOV) with the aim to assess the safety and early efficacy of three doses of plitidepsin (1.5, 2.0 and 2.5 mg/day) for three consecutive days in patients requiring hospitalisation for COVID-19 (n = 27). Based on the activity on a key mechanism of the viral pathogenesis, the use of this drug can be considered either in the earlier stages of infection, in a

multidrug regimen and for patients with more advanced disease, together with proper immunomodulators.

4.8. Dactinomycin

Historically, the anticancer drug dactinomycin was discovered in *Streptomyces parvulus* bacterium by Selman Waksman and colleagues in the 1940s [203] and launched after its initial FDA-approval in 1964. This intravenously administered cytotoxic antibiotic acts by intercalation to DNA and interruption of RNA transcription [38,87]. Recently, dactinomycin was shown to induce immunogenic cell death, suggesting a direct antineoplastic activity enhanced by immunomodulatory properties [93]. In fact, this drug is used in combination with other cytotoxic drugs in the management of a broad array of solid malignancies, such as gestational trophoblastic disease, Wilms' tumour and Ewing sarcoma [4,67,105,196].

The use of dactinomycin against coronaviruses was firstly described in 1979 by Kennedy *et al.* in the context of alpha-coronavirus 229E, involved in a spectrum of respiratory diseases from upper to lower airways infections [98]. Later on, Lewis *et al.* showed a potent inhibition of the feline enteric coronavirus strain 79-1683 in whole feline embryo cells [104]. Based on these and subsequent evidence, dactinomycin was widely used as an *in vitro* inhibitor of viral cellular transcription in infected cells [197]. Recently, a combination of dactinomycin and sirolimus showed a synergistic action against SARS-CoV-2-host proteins [225]. In fact, the association may inhibit DNA topoisomerase required for RNA synthesis as well as mammalian target of rapamycin (mTOR) signalling in human coronavirus infected cells [225]. Notably, the mTOR pathway is a target of sirolimus that has also shown a promising activity against MERS-CoV in addition to its known anticancer, immunosuppressive and antifungal properties [100]. However, these results for SARS-CoV-2 are limited by their *in silico* nature and warrant more clinical experiences beyond speculations. In addition, the use of this antibiotic is associated with an adverse safety profile when used at therapeutic doses as an antineoplastic agent that may be less suitable for severely ill patients with an acute infectious disease. In particular, the emetogenesis and myelosuppression can deserve a special consideration, and antiviral doses may be acceptable only if lower, as expecting a better safety profile.

4.9. Valrubicin

Valrubicin is a semisynthetic derivative of the anthracycline doxorubicin, used as intravesical topic treatment for the management of refractory non-muscle invasive bladder cancer [168]. The mechanism of action of valrubicin affects DNA synthesis and metabolism by inhibiting the incorporation of nucleosides and

topoisomerase II [13,168]. Based on computational hierarchical virtual screening, recently Wang showed that valrubicin alongside with other drugs, such as lopinavir and carfilzomib inhibits the main protease of SARS-CoV-2 [206]. Moreover, another report based on a machine-learning simulation has recently shown that valrubicin is a potential COVID-19 protease inhibitor [99]. However, no considerations on the effective doses have been modelled: the clinical use as an agent for systemic administration is not validated, being approved only as a topical solution, and any speculation for the design of clinical trials should account for the adverse safety profile of this chemotherapeutic.

4.10. Leronlimab

Leronlimab (PRO 140) is a humanised monoclonal antibody targeting the C–C chemokine receptor type 5 (CCR5) on T lymphocytes [96]. The role of CCR5 in the immunopathology of disease has been extensively studied since its discovery as a co-receptor for human immunodeficiency virus entry in 1996 [21] (NCT03902522 and NCT02859961). In addition, CCR5 is also expressed on a broad array of cancers and it is linked to various cancer hallmarks, particularly “avoiding immune destruction” [96]. It is under investigation in trials for few cancers, such as metastatic colon and breast cancers (NCT03838367), for which several other inhibitors of CCR5 are being developed (reviewed elsewhere: [96]). Previously, Chen *et al.* demonstrated that mRNA transcripts of CCR5 and other chemokine receptors were found upregulated in animals infected by SARS-CoV [30]. Additionally, enhanced production of these mediators was associated with recruitment of inflammatory cells and T lymphocytes influx into the lung parenchyma [30].

PRO 140 has shown a promising antiviral potency with a high barrier to resistance and encouraging short-term safety and tolerability [189]. A recently published *Preprint* suggested that leronlimab disrupts the CCL5/RANTES-CCR5 axis and restores immune homeostasis in COVID-19 patients [133]. The authors observed a marked CCR5 receptor occupancy on macrophages and T lymphocytes, rapid drop of circulating IL-6, CD4/CD8 ratio restoration and a significant decrease of patients’ viremia [133]. These encouraging findings boosted the rapid entrance of leronlimab into two clinical trials for mild to moderate and severe COVID-19 infection (NCT04343651 and NCT04347239). The largest trial using this drug is a randomised, double-blind and placebo-controlled, two-arm phase IIb/III trial investigating weekly leronlimab at a dose of 700 mg given subcutaneously (NCT04347239). A multicentred enrolment will include 390 patients with severe or critical disease with a mortality rate after four weeks of treatment as a primary end-point. Likewise, the other study is a phase II (n = 75) with a similar design that will enrol patients with mild to moderate symptoms

(NCT04343651). Patients will be randomised to receive leronlimab (700 mg) or placebo weekly in a 2:1 ratio with changes in total symptom score based on fever, myalgia, dyspnea and cough at day 14 as a primary end-point. Notably, the other outcome measures in this clinical trial include serum changes of cytokines and chemokines from baseline, CCR5 occupancy levels on macrophages and T^{reg} cells and CD3+, CD4+ and CD8+ T cell count from baseline. Based on the specific action in-between the viral pathogenesis and the immune-dysregulation, the use of this agent has been proposed for patients with mild clinical COVID-19 to prevent severe outcomes and reduce the related mortality.

4.11. Antineoplastic immune-checkpoint inhibitors

Based on the hypothesis suggesting the upregulation of immune-checkpoints such as PD-1/PD-L1, cytotoxic T lymphocyte antigen-4 (CTLA-4), lymphocyte activation-gene-3 (LAG-3) and T cell membrane protein-3 (TIM-3) pathways in excessive inflammatory response during sepsis [132], these types of drugs were repurposed beyond their use for the treatment of cancers [132,147]. Importantly, a recent meta-analysis including four studies showed PD-1 blockade improves patient outcome in sepsis [219].

Camrelizumab is a humanised IgG4-kappa anti-programmed cell death 1 (PD-1) monoclonal antibody used in cancer immunotherapy [114]. The blockade of PD-1 binding to its ligand PD-L1 enables reactivation of cell-mediated immune response against cancer cells [114]. Camrelizumab was tested in phases II/III clinical trials for advanced or metastatic oesophageal squamous cell carcinoma [91,92], osteosarcoma [212], relapsed or refractory classical Hodgkin lymphoma [177,178] and advanced hepatocellular carcinoma [145]. It received its first approval in China for treating RRHCL patients who received at least two lines of chemotherapy [114]. Additionally, it is currently being studied in two ongoing phase III trials for local advanced nasopharyngeal carcinoma (NCT03427827) and gastric and gastroesophageal junction adenocarcinoma (NCT04208347). The use of anti-PD-1 inhibitor camrelizumab in COVID-19 patients with pneumonia is being evaluated in a Chinese clinical trial (n = 120; ChiCTR2000029806) in comparison with the standard supportive care and the immunomodulatory agent thymosin.

Nivolumab and pembrolizumab are other immune-checkpoint inhibitors (ICIs) that were successfully introduced into the management of various solid cancers, particularly for melanoma [110,204]. These ICIs are both humanised and fully human anti-PD-1 IgG4 monoclonal antibodies. Their marked clinical activity depends on PD-L1 expression in some tumours, as well as on emerging predictive biomarkers, such as tumour-

mutational burden and microbiota that seem to mediate drug response [51,81,158].

Nivolumab is currently used in a multicentred phase II trial comparing its efficacy against tocilizumab, GNS561 (a chloroquine analogue), and standard supportive care in advanced or metastatic cancer patients with COVID-19 (NCT04333914). A total of 273 participants will be randomised to receive one of three treatments based on two cohorts. Patients that are asymptomatic or with mild disease will be prospectively assigned to the group comparing nivolumab versus GNS561 versus standard of care in a 1:1:1 ratio. Similarly, in patients with moderate to severe symptoms, nivolumab will be compared to GNS561 and tocilizumab. Survival after four weeks of randomised treatment is the primary end-point in this trial. Moreover, another ongoing French phase II trial ($n = 92$) is investigating the safety and efficacy of nivolumab in COVID-19 patients requiring hospitalisation (NCT04343144). Patients will be randomly allocated to receive nivolumab or standard of care (1:1 ratio). One pilot trial is also testing nivolumab in the earlier setting of COVID-19, in patients presenting with no or mild symptoms and not hospitalised – to examine the role of this ICI as an immune-enhancer, to accelerate the clearance of SARS-CoV-2 (NCT04356508).

Pembrolizumab was also repurposed for COVID-19. The COPENICO is a multicentred two-arm phase II trial (NCT04335305) that is currently enrolling patients with COVID-19 associated with pneumonia in Spain and who are non-responsive to frontline therapy. Twenty-four participants will be prospectively randomised to receive intravenous tocilizumab plus pembrolizumab or standard treatments (at physician's choice). The use of ICIs for COVID-19 is essentially intended as an immunomodulator, based on the proposed role of PD-L1 axis in the pathogenesis of ARDS and the expedited clearance of SARS-CoV-2 with anti-PD-L1 therapies [163]. Accordingly, the studies ongoing are testing these ICIs both for more severe cases, intending to attenuate the exuberant cytokine production [166], and for mild presentations, to prevent the onset of the hyper-immune pathological response and then dampen the viral replication [163]. Moreover, a recent report showed that PD-1 checkpoint inhibition in lung cancer patients who were diagnosed with COVID-19 does not appear to affect the prognostic outcomes [111]. Therefore, this provides a promising rationale for immunotherapy, which seems to have no unfavorable effects in this setting.

4.12. Kinase inhibitors

The blockade of kinase cascade is one of the successful targeted strategies in treating cancer in the modern era of oncology. Several kinase inhibitors including palbociclib, crizotinib, dabrafenib and imatinib, were introduced in practice for various cancers [69]. The rationale

for using kinase inhibitors in COVID-19 is principally based on their ability to inhibit host intracellular-viral events particularly during trafficking induced during the viral life cycle [140]. Viruses commonly use a number of host kinases during their invasion and thus, they are considered potential targets for kinase inhibitors repurposing as broad-spectrum antivirals [66,164]. Several previous reports demonstrated potent antiviral properties for kinase inhibitors in various virus-induced diseases, including SARS-CoV infections [8,17,35,46,100,128,129].

Imatinib is a marketed tyrosine kinase inhibitor for chronic myeloid leukaemia and gastrointestinal stromal tumours in the first-line setting [54,85]. The BCR-ABL1 oncoprotein with kinase activity was found to mediate virus entry in several reports encompassing hepatitis C and Ebola viruses [65,120]. In coronaviruses, recent studies indicated that imatinib blocks cell–cell and cell–virus membrane fusions [36,175]. Additionally, in *in vivo* and *ex-vivo* mouse models, the use of imatinib protected the lungs from ischemia/reperfusion injury [112,187], attenuated inflammation and vascular leak [154] and prevented lung injury and death [182].

Clinically, the IMAGE-19 trial is a single-arm phase II study in France that will enrol 90 elderly COVID-19 patients (NCT04357613). Early imatinib therapy at a dose of 800 mg daily for two weeks will be evaluated to prevent the progression to severe disease in patients >70 years with COVID-19. The efficacy of imatinib at a dose of 100 mg is also being examined in combination with hydroxychloroquine in the Spanish Covid-19HUF phase II trial ($n = 165$) (NCT04346147). Participants will be randomised to receive either oral lopinavir/ritonavir (protease inhibitors), imatinib or baricitinib (JAK inhibitor), all combined to hydroxychloroquine. The trial completion date is estimated in September 2020.

The COVERAGE study is a large French multiarm phase III trial (NCT04356495) that will investigate the efficacy of imatinib and several other experimental treatments. A total of 1057 symptomatic COVID-19 patients will be randomly allocated to receive imatinib, or telmisartan, or favipiravir, or hydroxychloroquine as compared to a control group with the objective to reduce the risk of hospitalisation or death in elderly patients. Another phase III trial using imatinib versus placebo for hospitalised COVID-19 patients is currently recruiting in the United States ($n = 204$, NCT04394416). Given the acceptable safety profile of imatinib and the several available generics, this seems to be an effective strategy for clinical repurposing. The drug repurposing of imatinib in COVID-19 has been embedded for the early phases of the disease, commonly referred to as the “viral response” phase [174]. Imatinib seems to tackle the mechanisms of viral entry and cell–cell spread, functioning as an antiviral agent. In this initial stage, the drug development is intended to reduce the duration of the symptoms, minimise the

contagiousness among people and prevent the progression to severe COVID-19.

Duvelisib is an orally bioavailable phosphatidylinositol 3-kinase (PI3K) selective inhibitor that is being studied for haematologic malignancies [94]. The PI3K–AKT–mTOR signalling is frequently dysregulated in cancer as well as during hyperinflammation and its therapeutic targeting is highly represented in ongoing clinical trials. Duvelisib was investigated in the DUO phase III trial and compared to ofatumumab for refractory and relapsed chronic lymphocytic leukaemia and small lymphocytic lymphoma [59]. It was approved by the FDA in 2018 for treating adult patients with relapsed or refractory chronic lymphocytic leukaemia or small lymphocytic leukaemia after at least two prior therapies [88]. There is currently one pilot phase II trial examining the efficacy of duvelisib twice daily at a dose of 25 mg for 10 d in COVID-19 patients as monotherapy (NCT04372602). This single-arm study will test the hypothesis that PI3K blockade could hamper the immune system hyperactivation and therefore reduce lung inflammation and interfere with the viral cycle. It is estimated to enrol 25 participants, with overall survival as a primary end-point. Notably, the PI3K inhibitors including the drug duvelisib can cause lung inflammation and increase the risk for infections, as consistently demonstrated for multiple compounds of this pharmacological class, and a special caution is warranted during the clinical trials using this class of molecules.

The aberrant activation of the Bruton tyrosine kinase has a key role in the tumorigenesis of B-cell lymphoma [77]. Its abrogation seems to protect lungs from acute immune-induced or lethal influenza-induced injuries [60,102] as well as sepsis-induced injury [224]. Zanubrutinib is an orally administered irreversible inhibitor of the Bruton tyrosine kinase [186,218,220]. This drug showed a durable and high overall response rate in patients with relapsed/refractory Mantle cell lymphoma [179]. Recently, zanubrutinib received accelerated FDA approval for treating adult patients with this rare and aggressive cancer who have received at least one line of therapy [186]. For COVID-19, a phase II trial in the United States is ongoing, aiming to dampen the disease-related immune dysregulation and hyper-inflammation (NCT04382586). This study is expected to randomise 52 participants to receive zanubrutinib combined to supportive care versus placebo and to test the hypothesis of increasing respiratory failure-free survival rates in patients with pulmonary distress. Ibrutinib is another inhibitor of Bruton kinase inhibitor, approved to treat indolent B-cell malignancies and chronic graft-versus-host disease, now being repurposed for COVID-19 [193]. An earlier experience with this drug in COVID-19 patients with cancer showed marked improvements and resolution or near resolution of COVID-19 symptoms during the follow-up

[193]. In the same way, the iNSPIRE (NCT04375397) is a phase II that will enrol 46 COVID-19 patients to randomly receive ibrutinib versus placebo along with standard supportive care. The trialists expect that ibrutinib will reduce respiratory failure and their findings are expected to be released in September 2020.

Opaganib (ABC294640) is a first-in-class selective inhibitor of sphingosine kinase-2 (sk2). This key enzyme in lipid signalling promotes tumour growth, cell proliferation and cytokine production during inflammation [139,143,176–178]. It was previously noticed that sk2 is the host factor that impacts the viral replication process of the chikungunya virus [149] and maintains viral latency and survival in Kaposi's sarcoma-associated herpes virus [39]. This drug has an orphan drug designation for cholangiocarcinoma [89] and is being tested in clinical trials for patients with metastatic castration-resistant prostate cancer (NCT04207255) and advanced cholangiocarcinoma (NCT03377179). It is also under development for COVID-19-associated pneumonia in a phase II study (NCT04414618). This is a randomised, double-blind, placebo-controlled trial in 40 adult patients with positive pneumonia, using daily opaganib for two weeks.

Saracatinib (AZD0530) is a tyrosine kinase inhibitor of the c-SRC family of proto-oncogenes. This pathway mediates signal transduction for several cell functions such as proliferation, cell-to-cell transmission, epithelial-to-mesenchymal transition, vascular permeability, invasion and metastasis [33,131]. Moreover, it interferes with additional cell signalling such as PI3K/mTOR and MAPK [151]. Saracatinib selectively abrogates this enzyme and was investigated in several clinical trials for recurrent osteosarcoma [11], platinum-resistant ovarian cancer [116], unresectable thymic malignancy [76] and cancer-induced bone pain [41]. SRC kinases have also a role in endocytic trafficking [150] and there is a growing body of evidence indicating the inhibitory properties of viral replication of dengue viruses and MERS-CoV by the anticancer agent saracatinib [42,171,228]. This makes this drug a possible candidate for further investigations on other members of the Coronaviridae family. In a recently published *Preprint*, molecular modelling showed that saracatinib binds to the nucleocapsid protein sites of SARS-CoV-2 along with other previously discussed drugs such as imatinib and camostat [188]. To the best of our knowledge, saracatinib has not entered into any COVID-19 clinical trial. The clinical use of molecular blockers of PI3K–AKT–mTOR, Bruton tyrosine kinase, sk2 and c-SRC is anticipated to exert immune-modulating rather than antiviral effect. To date, the drug development of this group of tyrosine kinase inhibitors is included in the protocols for hospitalised patients with severe COVID-19, to rescue the immune-dysregulation and improve the outcome.

4.13. Selinexor

Selinexor is a first-in-class orally bioavailable inhibitor of Exportin-1 (XPO1) [185]. XPO1 is an export receptor for various proteins and RNA molecules with oncogenic properties and is frequently altered in cancer [7]. Moreover, it is implicated in several cancer pathways including cell growth and survival, differentiation and drug resistance [127]. XPO1 suppression by selinexor in human clinical trials demonstrated promising results for patients with heavily treated relapsed or refractory multiple myeloma [29,152,202]. The drug received first an accelerated approval by the FDA in the United States of America (USA) in 2019 in combination with dexamethasone [152]. The antiviral activity of selinexor is based on its potential to interfere with host targets including XPO1 that interacts with SARS-CoV-2 proteins [73,194]. In fact, blockade of this nucleocytoplasmic transporter can sequester viral materials in the host cell nucleus and therefore reduce its replication mechanisms and viral load [194]. Moreover, selinexor has also selective and potent anti-inflammatory effects on sepsis pathophysiology [211]. Based on these concepts, two phase II randomised and controlled trials of low dose selinexor were initiated for COVID-19 patients with moderate to severe disease (NCT04355676 and NCT04349098). The largest trial that is currently recruiting patients is single-blind and placebo-controlled and is expected to enrol 230 hospitalised participants with severe COVID-19 (NCT04349098).

5. Perspectives

The COVID-19 pandemic continues to spread globally with 24,484,968 cases and 832,533 deaths registered worldwide, including 6,016,069 cases only in the USA (as of 27th August 2020). New effective therapeutics are urgently needed, as the results from the first clinical trials or experiences with drugs for COVID-19 are mostly disappointing. There is a massive demand for drug repurposing in this rapidly evolving setting, relying on the benefit of a shorter drug development and the capability to turn preclinical intuitions in clinical experiences. In fact, it is a potential successful drug discovery process, more efficient and resource sparing than the classic drug development from early phases for new molecules, taking generally a longer time (typically 10–15 years). While drug repurposing is not an excuse to design low-quality clinical trials or emphasise biased results from preliminary findings and non-controlled cohorts, the possibility to use molecules with an established safety profile certainly represents an advantage. The US [ClinicalTrials.gov](https://clinicaltrials.gov) database shows that ≥ 3100 clinical studies for COVID-19, including more than 1700 interventional trials (accessed as of 27-08-2020), are ongoing, portraying the commitment of the global

medical community to accelerate the COVID-19 research agenda – in a successful interplay between laboratory researchers and clinical trialists, ultimately to serve clinicians and improve patients' care. However, in the urgent need to provide a response to a burning unmet need, many isolated smaller clinical trials or off-label use of medicines outside the investigations have been observed, perhaps fragmenting the research agenda in uncoordinated silos – thus diluting the research efficiency. The large proportions of ongoing clinical trials are single-centred observational studies with high risk of bias and therefore providing low level of evidence when their findings are reported. The capacity to inform the decision-makers is inevitably weak when the clinical studies generate doubts over certainties. This was recently outlined in a critical appraisal of registered COVID-19 trials on [ClinicalTrials.gov](https://clinicaltrials.gov) database [141]. In fact, only 13.6% of these clinical trials are prospective cohorts that may provide level 2 evidence [141]. Therefore, the lack of effective international cooperation is suggested, along with the negligence of other key aspects to tackle the pandemic, including the disease prevention in high-risk populations. Another emerged critical issue is the tremendous heterogeneity in the rates of COVID-19 new cases across different geographic areas with a significant decline in some countries that may halt and decrease expected enrolment in ongoing clinical trials [192]. The important number of launched monocentric studies and the observed decline of incident COVID-19 cases in some countries should promote considering adaptive study designs to fulfil enrolment [192]. For this perspective, trialists are encouraged to follow master protocols when testing their hypotheses, as suggested by the World Health Organization, in the design of the international adaptive trial SOLIDARITY (e.g. NCT04330690). A significant experience with master protocols in oncology and haematology was achieved, which may accelerate drug development and reduce ineffectiveness in urgent clinical trials [184].

Among the dozens of coronavirus drugs that are being repurposed or in development, a number of anticancer compounds are also being studied as potential future drugs for COVID-19. At this time, the best option for patients is the enrolment in clinical trials. In the current scenario where no drug has provided a “parachute” effect on the prognosis of patients with COVID-19, especially with more severe disease, the use of off-label medicines based on over-emphasised interpretations of the preliminary data from small cohorts or anecdotes is strongly discouraged. With nearly one thousand clinical trials ongoing, there is virtually room to allocate much more patients in truly promising clinical trials, tailoring specific populations across the continuum of care – from the prevention to the treatments and palliative care. The notion is clear: treatment options in settings of high medical unmet need can be improved only in controlled clinical trials, both offering the most innovative

therapeutic approaches and prompting the disengagement from established clinical interventions with narrow added value, a common contingency observed when no treatment is available for a particular disease. Having an impact on health requires quality behind.

In response to the “high clinical unmet need” and to support clinicians, who are facing the daily challenges with the aim to manage severe health conditions associated with dismal prognosis, the implementation of clinical trials may serve to boost the access to treatments, making treasure of such precious information. In fact, the off-label use of some medicines, most strikingly hydroxychloroquine almost everywhere, outside a clinical trial or a registry-based accountability is the primordial error and the enhancer of confusions — generated by the disconnect between the anecdotal clinical perception of benefits and the objective data collected. In this regard, the regulatory agencies for medicines in many regions and countries have reformulated simplified procedures and supported more flexibility in the design, evaluation and adoption of clinical trials, to meet the patients’ needs. For example, the European Medicines Agency (EMA) has delineated regulatory flexibilities to help pharmaceutical companies cope with the consequences of the pandemic, emphasising the need to keep high-level of quality, safety and efficacy for medicinal products developed for COVID-19. The message is clear: patients’ safety and quality of science in clinical trial conduction is non-negotiable [52] (accessed 30-05-2020). EMA has established rapid procedures through accelerated scientific advice, rolling review and expedited treatment of the applications for marketing authorisations [53] (accessed 30-05-2020).

The harmonisation of regulatory processes and enhanced drug development must necessarily proceed in parallel, committing to accelerate the drug repurposing, ensuring safety thus prompting early interruptions of trials when new safety signals emerge, or major results of activity are collected. However, while endorsing the acceleration in the revisions, decisions, approval and authorisation, the regulators should commit to ensure the design of quality clinical trials, pushing for the definition of the benefits based on patients’ relevant end-points and dissecting the reasonability and opportunity to favour surrogate end-points in selected instances — when appropriate for the setting and highly predictive or correlated with the mortality outcomes — the truly important ones.

The high-throughput screening of approved molecules capable to dock to the key protein domains of SARS-CoV-2 pathogenesis or related to the immune-dysregulation will identify thousands of possible compounds, from *in silico* models. An important success for this approach was recently achieved based on a library profiling of approximately 12,000 clinical-stage or FDA-approved compounds [153]. The authors of this paper published in *Nature* identified various molecules such as the kinase inhibitor apilimod that antagonise SARS-

Box 1. Recommended reading of particular interest

	DOI
Larson C <i>et al.</i> COVID-19 and Cancer: A guide with suggested COVID-19 rule-out criteria to support clinical decision-making. <i>Biochimica et Biophysica Acta (BBA) - Reviews on Cancer</i> . 2020; 188412.	https://doi.org/10.1016/j.bbcan.2020.188412
Zeggini E <i>et al.</i> Biomedical Research Goes Viral: Dangers and Opportunities. <i>Cell</i> . 2020; S0092-8674(20)30576-6.	https://doi.org/10.1016/j.cell.2020.05.014
Sanders <i>et al.</i> Pharmacologic Treatments for Coronavirus Disease 2019 (COVID-19): A Review. <i>JAMA</i> . 2020; 10.1001/jama.2020.6019.	https://doi.org/10.1001/jama.2020.6019
Mercorelli B <i>et al.</i> Drug Repurposing for Viral Infectious Diseases: How Far Are We? <i>Trends Microbiol</i> . 2018; 26(10):865–876.	https://doi.org/10.1016/j.tim.2018.04.004
Farha MA, Brown ED. Drug repurposing for antimicrobial discovery. <i>Nat Microbiol</i> . 2019; 4(4):565–577.	https://doi.org/10.1038/s41564-019-0357-1
Boldeanu V <i>et al.</i> Broad-spectrum agents for flaviviral infections: dengue, Zika and beyond. <i>Nat Rev Drug Discov</i> . 2017; 16(8):565–586.	https://doi.org/10.1038/nrd.2017.33
Takizawa N, Yamasaki M. Current landscape and future prospects of antiviral drugs derived from microbial products. <i>J Antibiot (Tokyo)</i> . 2017; 71(1):45–52.	https://doi.org/10.1038/ja.2017.115
Kaufmann SHE <i>et al.</i> Host-directed therapies for bacterial and viral infections. <i>Nat Rev Drug Discov</i> . 2018; 17(1):35–56.	https://doi.org/10.1038/nrd.2017.162
Pfefferle S <i>et al.</i> The SARS-coronavirus-host interactome: identification of cyclophilins as target for pan-coronavirus inhibitors. <i>PLoS Pathog</i> . 2011; 7(10):e1002331.	https://doi.org/10.1371/journal.ppat.1002331
Zumla A <i>et al.</i> Coronaviruses - drug discovery and therapeutic options. <i>Nat Rev Drug Discov</i> . 2016; 15(5):327–347.	https://doi.org/10.1038/nrd.2015.37
de Wit E <i>et al.</i> SARS and MERS: recent insights into emerging coronaviruses. <i>Nat Rev Microbiol</i> . 2016; 14(8):523–534.	https://doi.org/10.1038/nrmicro.2016.81

CoV-2 replication *in vitro* and in primary human lung explant model [153]. While it is essential to boost the drug-discovery based on approved medicines, to accelerate the drug development overall, the construct of preclinical models should be *in vitro* and *ex vivo* strictly

Box 2. Useful resources and database

	Website
The US Clinical Trials Database	https://www.clinicaltrials.gov/
COVID-19 clinical trials	https://www.clinicaltrials.gov/ct2/results?cond=COVID-19
Coronavirus Disease 2019 (COVID-19) Treatment Guidelines	https://www.covid19treatmentguidelines.nih.gov/
COVID-19 Studies from the World Health Organization Database	https://www.clinicaltrials.gov/ct2/who_table
Elsevier Coronavirus Resource Directory	https://www.elsevier.com/novel-coronavirus-covid-19
COVID-19 Databases and Journals	https://www.cdc.gov/library/researchguides/2019novelcoronavirus/databasesjournals.html
EULAR COVID-19 Database	https://www.eular.org/eular_covid19_database.cfm
COVID-Evidence Database	https://covid-evidence.org/database
Cell Press Coronavirus Resource Hub	https://www.cell.com/COVID-19
COVID-19 SARS-CoV-2 preprints from medRxiv and bioRxiv	https://connect.biorxiv.org/relate/content/181

implemented in collaborating research facilities, committed for the research on COVID-19. This is an effort of everyone. In the fight against COVID-19, the entire chain of medical researches has rapidly become multidisciplinary, truly intersected with specialists of several branches, transversally positioned. Offering technology and knowledge to develop the best science at the service of a public health global emergency recalls the humanitarian mandate of medical science – to improve population health and wellbeing for all.

The hope is that drug repurposing to boost the discovery of new treatments for human diseases will find the right place and that this innovative and transversal operational model of scientific research will get common, to result in timely impact on populations at the most effective and efficient utilisation of resources and knowledge. Time for wasteful duplications based on competitive conflicts should be over; the COVID-19 will enlighten the solidarity in science, sublimating narcissism for the ultimate goals of serving people, especially when facing the hardest moments of their lives, for deadly diseases.

6. Conclusion

Controlling the SARS-CoV-2 pandemic warrants multidisciplinary collaboration to find relevant treatment options. From the evidence reviewed here, a number of anticancer drugs seem to retain a promising activity to treat patients with COVID-19. The high level

of evidence should still be the primary objective of the clinical research, despite the critical circumstances during a pandemic. Hence, testing a battery of known drugs in clinical cohorts will certainly provide an additional value to these molecules (for more information, see Boxes 1 and 2). Although the ideas of using repurposed anticancer drugs might be criticised because of the toxicity profile of some of them, we believe that this review article may provide inputs and guide the future directions for clinical research – envisioning an approach beyond the definition of separate medical subjects by vaporising the common barriers between medical disciplines. This might help address innovative operational models and regulatory flexibilities in this unique *momentum* of the international research, united to control a global pandemic, committed to serve the people in all over the world.

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Conflicts of interest statement

Khalid El Bairi, Dario Trapani, Angelica Petrillo, Cécile Le Page, Hanaa Zbakh, Bruno Daniele, Prof. Rhizlane Belbaraka and Prof. Said Afqir have no COI to declare. Prof. Giuseppe Curigliano has received honoraria from Pfizer, Novartis, Lilly, Roche; fees for expert testimony and medical education from Pfizer and has participated in advisory boards for Pfizer, Roche, Lilly, Novartis, Seattle Genetics, Celltrion.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.ejca.2020.09.014>.

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