

Potential combination endothelial progenitor cell and antiviral therapies for reversal of impaired angiogenesis induced by SARS-CoV-2 infection

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Abstract

Late stage detection of COVID-19 disease pathogenesis induced by SARS-CoV-2 infection may prove to be fatal to the infected patients with the severity increasing, if inhibition of differentiation of hematopoietic endothelial progenitor cells, or their sustained deficiency, is not reversed. Antiviral drug treatments of such individuals need not necessarily resuscitate from a severe impairment of normal angiogenesis of the vascular endothelium. The virus targets the endothelial progenitor cells which co-express the hematopoietic stem cell marker CD34 and the angiotensin converting enzyme 2 (ACE2), latter being the host receptor for this pathogen. There is not an apparent segregation of CD34 and ACE2 antigens, strongly implicating inhibition of differentiation of virus infected progenitor cells. The embattled clinical condition of SARS-CoV-2 infected patients may well require a dual-mode endothelial progenitor cellular infusion and antiviral drug molecular therapies to stage a rapid clinical recovery. Umbilical cord blood derived CD34+ progenitor cells are the most optimal for rapid availability, harvesting and infusion into the severely ill infected patients, who are most likely to be non-responsive to solely antiviral drug treatments. This is about rapidly prepared allogeneic cell infusion therapy and not autologous cell transplantation which is impractical for these relatively acute and short-term conditions and treatments of SARS-CoV-2 infections. A combination progenitor cell and antiviral drug treatment is suggested for recovery and maintenance of normal angiogenesis of infected patients who may otherwise not survive. Hence the concept and its basis discussed needs to be urgently advanced to translation stage.

Keywords: SARS-CoV-2, CD34, CD133, endothelial progenitor cells, angiogenesis inhibition, transient treatment, progenitor cell therapy, neovascularization, thrombocytopenia, microRNA, iPSC, organoids

Hypotheses

Variations caused by SARS-CoV-2 in levels of disease pathogenesis in infected patients from asymptomatic or minor symptoms, to severe damage to vascular cells, raises questions on the return to normal functioning of the vasculature and recuperation of the severely affected physiology of angiogenesis pathways.

The level or stage of the COVID-19 disease at which the patients' infection by SARS-CoV-2 is tested to be positive by PCR for existing infection, is critical to determine the therapeutic strategy to be adopted. If SARS-CoV-2 infection has advanced the virus

induced disease to injury of the vasculature and its endothelium, eliciting an abnormally potent cytokine storm due to the severe but acute surge of immune responses, then recovery may need the dual administration of exogenous cellular in addition to the molecular drug therapies.

Umbilical cord blood (UCB) derived endothelial progenitor cells (EPCs) despite their allogeneic nature are evidenced as the most resourceful and generally safest for infusion into the patients [1], in this instance battling the severity of SARS-CoV-2 infection. However, it is important to distinguish between our

suggested intravenous (IV) cell infusion for recovery and homeostasis of angiogenesis than undertake the impractical cell transplantation, in such a critical clinical scenario. Infusion can be performed following rapid cell preparation including any pre-IV HLA matching on a few UCB derived cells [1].

Herein a combination cellular and molecular therapy that has all the prerequisites for rapid clinical administration to the virus infected patients struggling between life and death is proposed. **Figure 1** depicts the desirable therapeutic approach if the infected patient is already at the disease stage where she needs ventilator support to remain stable and for recovery.

Lung microvascular endothelial progenitor cells (EPCs) play a pivotal role in maintaining the life of an individual [2]. In our **Figure 1**, UCB derived endothelial progenitor cells expressing the characteristic CD34 antigen of the hematopoietic lineage (HEPC) [1,3], may come to the rescue of replenishing the severely depleted endothelial cells of the vasculature of the critically ill and even non-responsive patients, arising from SARS-CoV-2 infection. However, such human umbilical cord hematopoietic endothelial progenitor cell (hUCHEPC) infusion still would need an anticipated antiviral drug treatment for containment of the virus replication, to aid in the HEPC/EPC homeostasis and stabilization for reversal of the otherwise aggravated patients' physiological condition.

Inhibitors of this viral RNA synthesis (from its +ssRNA

strand), such as remdesivir and ED-1931/ED-2801, are very much in clinical trials with much anticipated efficacy thru' aborted viral replication, due to the incorporated modified nucleotides [4,5]. Viral protease inhibitors of SARS-CoV-2 serine protease are also being developed [6-9]. Therefore, for the infected individuals who become severely ill from damaged vasculature, in order to re-attain stable vascularization (neo-vascularization), hUCHEPC may be the most plausible therapy to be adopted [10-12], aided by the antiviral drugs [4-9], to achieve and sustain the normal angiogenesis recovery.

Binding of the SARS virus spike (S) protein to the endothelial cell angiotensin converting enzyme-2 (ACE2) [13], as it also happens with strong affinity to SARS-CoV-2 in particular, is physiologically deleterious to the host HEPC/EPC self-renewal and differentiation into mature cells of different types of the human vasculature [14-16]. The steady state disruption between the HEPC/EPC in effect during the virus infection also necessitates the antiviral drug induced containment and clinically eventual cessation of virus replication to maintain functioning levels of endothelial progenitor cells. Else, resurgence and function of the HEPC/EPC prevail during decreasing viral loads not requiring the cellular therapy component, presumably in the virus infected asymptomatic individuals, or patients that are responsive to antiviral drugs without clinical advancement to pneumonia and severe lung-vascular cell injury.

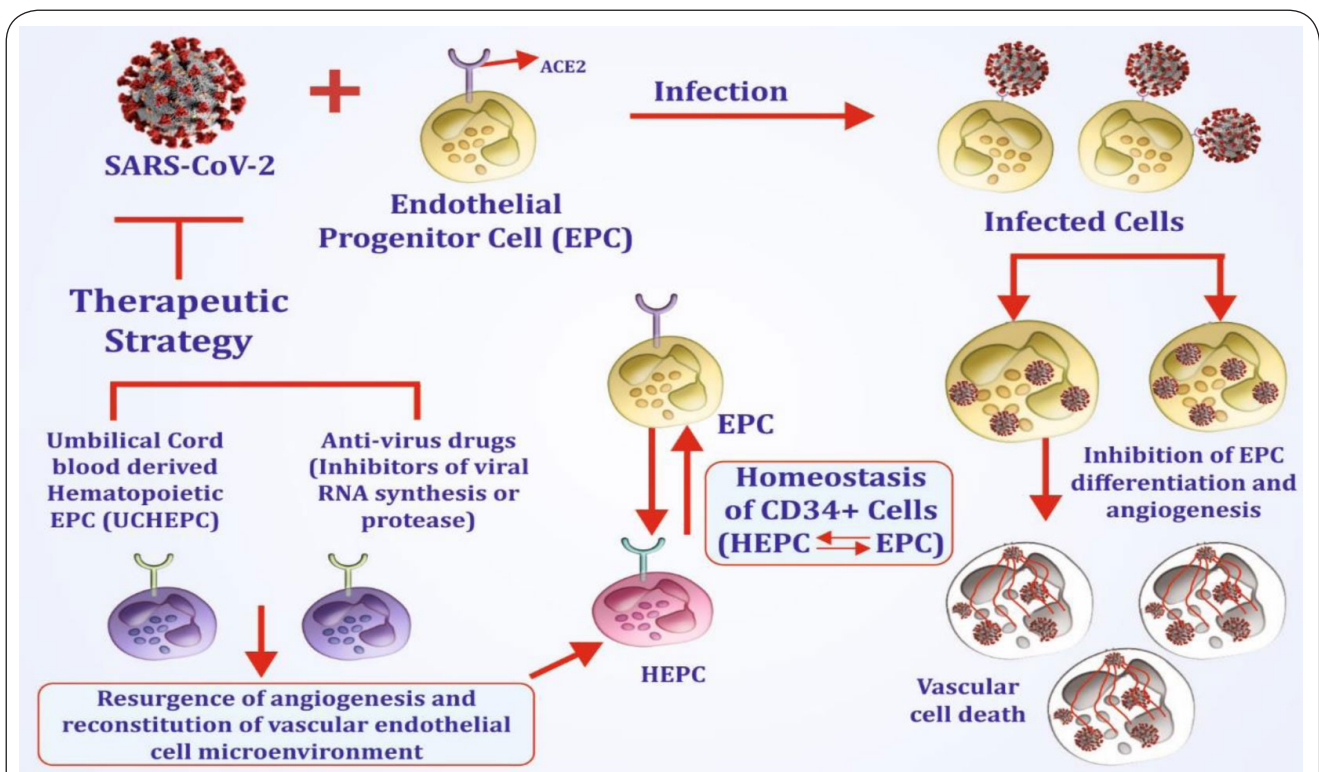


Figure 1. Schematic showing the proposed endothelial progenitor cell and antiviral drug dual approach therapies to simultaneously replenish the lost cells and contain the virus replication, for rejuvenation of the vasculature, in late stage pathogenesis of SARS-CoV-2 infected patients. Note: The virus and cell are taken from Google online search.

Despite the common haematopoietic lineage association of the endothelial progenitor cells that also express the characteristic hematopoietic cell surface CD34 antigen marker [1,17], onset of stage specific divergent lineage commitment of HEPC and EPC (CD34+) lead them to differentiate separately into immune and vascular cells respectively. HEPC are in greater proximity to being stromal than EPC. Previously, we have performed engraftments and reconstitution of CD34+ progenitor cells, isolated rapidly by AutoMacs, in human stromal microenvironment in SCID-hu animals [18,19]. In the context of EPC, CD34 expression lends credence to the stromal proximity and coexistence between HEPC and EPC populations which are discriminated only through their lineage differentiation into cell types categorized by their functions designated prior to HEPC/EPC "split". Lung vascular EPC microenvironment [2] preservation is essential for the circulation of vascular cells which are the primary target of this virus and interruption or failure of this function is potentially lethal to the infected patients. UCB derived CD34+ H/EPCs [19] for the cellular therapy can be used for regeneration of the severely depleted vascular progenitor cells in the SARS-CoV-2 infected patients.

Moreover, purified CD34+ cells, even from peripheral blood are significantly more preventative of graft-versus-host disease (GVHD) in allogeneic transplantations [20]. Further, UCB derived CD34+ cells carry the potential to confer greater survival incidence and ward off GVHD related mortality in transplantation recipients [21]. Thus, when GVHD due to CD34+ cell transplantation is at a minimal in causing patient mortality, a transient readily preparative and administrable CD34+ cell infusion therapy that we have proposed is not expected to confer a burden on the immune system of the COVID-19 patients. Rather any such presumably mild reaction [20,21] could be even synergistically beneficial in repressing or lowering any prevailing clinical cytokine storm condition in these SARS-CoV-2 infected patients, in conjunction with some of the other prescribed and adopted treatments.

Interestingly, regions of the S/pike protein of SARS-CoV-2 cause cytokine storm induced thrombocytopenia, is similar to that effected by HIV-1 V3 loop determinants [22,23]. The commonality of virus-induced depletion of the megakaryocyte lineage CD41+ cells [24], may well strengthen the involvement of microRNA in the inhibition of differentiation of progenitor cells [25], H/EPC, causing insufficient supply of cells required to be maintained in normal blood circulation and in the case of both these viruses, for platelet formation and levels.

Our recent report wherein differential regulation between microRNAs, miR-15a and miR-24, is implicated in HIV-1 induced inhibition of differentiation of CD34+ hematopoietic progenitor cells occurs, purportedly via HOX, and thus ensue different lineage cytopenias [25]. Similarly, miRNA implicated phenomena are involved in EPC differentiation and these cells' regulatory signals on miR-141-3p (via Hsr1) and miR-126 (via VEGFR2) in destabilization of proteins controlling prevention of lung injury [26], and type 2 diabetes [27], respectively. Such findings

compel us to anticipate microRNAs to become the focus of consideration for drug therapies to address depletion of progenitor cells. Further, microRNAs may lead to cell-free therapies possibly encapsulated in bio-scaffolds. Targeting or using microRNAs may further delineate strategies to tackle the virus pathogen or its deleterious clinical conditions.

Sustained maintenance of stromal microenvironments in the human vasculature [28] post-therapy, as depicted in **Figure 1**, pertaining to the homeostasis between HEPC and EPC balance of the cell surface antigens, such as the inevitable CD34 [26] marker expression in this regard, is of utmost underlying physiological necessity. The homeostasis between HEPC and EPC as we depicted in **Figure 1**, seems to be driven by the co-expression of CD133 and CD34 on the EPC, with retention of the hematopoietic origins, for a sustained replenishment of the double-positive (CD34+CD133+) EPC that participates in the angiogenesis or neovascularization [1]. Whether a selection enriched for the multi-potent ALDH+CD133+Lin-phenotype cells' repopulating potential as in *in vivo* [29], of these CD34+ subset cells when derived from the UCB, would be more efficacious for the H/EPC+ therapy for severe acute SARS-CoV-2 infections, needs to be determined. The vascular endothelium may have been depleted of the repopulation supporting stromal microenvironment in the SARS-CoV-2 infected patients but then again, this ALDH+CD133+Lin-/CD34+ cell population enrichment [29] may very well be the cofactor in the midst of the whole CD34+ cells together providing the cellular microenvironment needed for rapid and irreversible neovascularization. These co-populating progenitor stem cell subset phenotypic requirements but led by the focus and ease to collect large numbers of the CD34+ cells from different UCB samples, will be expected to counter and recover the SARS-CoV-2 induced damage to the vasculature. This will address the well-being of the affected patients in general, and SARS-CoV-2 infected patients in particular, for attainment of the elimination of the entire and last vestiges of infection and the virus induced pathogenesis.

Thus we hope that basic science and clinical researchers may follow up on these suggested lines of therapeutic treatment to evaluate the feasibility and efficacy in severely ill COVID-19 patients to stage a recovery, not without a sense of urgency.

Furthermore, since SARS-CoV-2 infections also cause dysfunction of disparate human organs including the lungs, heart, brain, kidney, intestine, etc., disease modeling by employing the human induced pluripotent stem cells (hiPSC) derived organoid model systems [30-40], lead the path towards the mechanisms underlying organ failures due to SARS-CoV-2 infections. Such experimental investigations using hiPSC derived organoid models could well generate molecular drug therapies to treat preemptive tissue damage by SARS-CoV-2 infections.

Ethical considerations

As this article deals only with presentation of perspectives on

the subject without any experimental, scientific and clinical investigations, including non-use or non-involvement of biological materials, or there were no recruitment of human subjects, therefore informed consent, ethics committee, or institutional review board (IRB) approvals and clinical trials registration are all not required.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

Authors' contributions	PSK	BR	STR
Research concept and design	✓	✓	--
Collection and/or assembly of data	✓	✓	--
Data analysis and interpretation	✓	--	--
Writing the article	✓	--	--
Critical revision of the article	✓	--	✓
Final approval of article	✓	--	✓
Statistical analysis	--	--	--

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