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COVID-19: From Cough to Coffin

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Abstract

The primary determinants of Covid severity are ACE genotype, the falciparum antigen (CD147), and vitamin D status. Specifically the I (insertion) and D (deletion) alleles, ACE polymorphisms determine the balance between the RAS (Renin Angiotensin System) and the KKS (Kallikrein Kinin System) in the response to SARS CoV2 (SARS2). CD147, the falciparum antigen, mediates the damage. Vitamin D modulates the immune response. The RAS and KKS connect Covid-19 to Kawasaki's Disease (KD) and Toxic Shock Syndrome (TSS). Covid-19 pathogenesis is embroiled in a nature versus nurture debate, as it seems to target people of color, unless you live in sub Saharan Africa. There are only three plausible explanations for the latter and they have all been selectively ignored/suppressed by mainstream medicine. This article speaks to the genotypic nature of Covid-19. Angiotensin II, bradykinin, ACE2, ACE and its two polymorphic alleles play vital roles. They predict disease severity. They portend the ARDS variants. They portend extra pulmonary disease or not. The heavily glycosylated CD147 epitope on the spike protein S is key. It has been dismissed as non-existent by flawed studies. Yet its interaction with CD147 receptors on erythrocytes and T lymphocytes cannot be denied and is at the heart of the myocarditis conundrum. Using this key, multiple dots are connected and a red alert issued, whether Covid-19 or vaccine related. These include thrombosis, immune deficit, cancer progression, autoimmune disease, and ADE (Antibody Dependent Enhancement) for those at risk. In susceptible vaccinees its deleterious effects are accelerated. Assessment of this and preventative approaches are explored.

Subject Areas

Pathology

Keywords

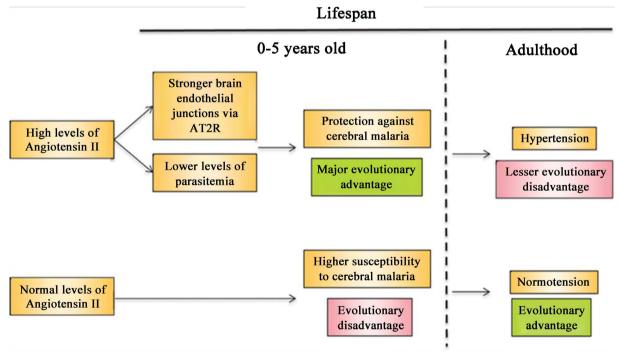
CD147, CD8, Lectin, Glycan, Epitope, Angioedema

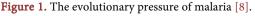
1. Introduction

In early 2020 pulmonologists found that many of their patients did not fare well after initiating mechanical ventilation [1]. COVID-19 ARDS had a worse outcome than ARDS with mortality rates ranging from 65.7% to 94% in patients who required mechanical ventilation. When mechanically ventilated patients were anti-coagulated mortality dropped from 62.7% to 29.1% [2]. Then in April 2020 a research group in the Netherlands discovered the critical KKS contribution to Covid-19 pathogenesis [3]. This was quickly followed in July by a similar article from the Oak Ridge National Laboratory linking the RAS and KKS systems [4]. Recent 2021 articles have highlighted differences between Covid ARDS and typical ARDS [5] [6]. Once perspective is broadened from just the RAS to include the KKS clarity which begins to emerge. The impact of ACE polymorphisms is glimpsed. Correlating the physiology with the clinical findings becomes possible. There appears to be two types of Covid-19 ARDS that reflect the predominance of either the ACE I allele or the D allele. One tends toward angioedema (typical ARDS), the other toward microthrombosis (Covid ARDS). The dual angiotensin II/bradykinin nature to Covid-19 drives this. The evolutionary connection between these ACE polymorphisms and falciparum malaria is well known [7] (see Figure 1), as is the parasite's dependence on erythrocyte CD147 receptors for entry.

2. Discussion

The ACE DD genotype seems to have evolved in Africa over many millennia as protection against malaria. ACE is almost 70% higher in ACE DD v ACE II [9]





[10]. This seems to work via angiotensin II and AT2Rs, enhancing endothelial tight junctions and preserving the blood brain barrier. Surviving cerebral malaria as a child was worth the risk of death due to hypertension as an adult. Given the overarching evolutionary pressure falciparum malaria has exerted on the ACE genotype and given the close clinical and lab parallels between malaria and Covid-19, more intense scrutiny of ARDS through the recently described KKS lens seems warranted.

2.1. ARDS

Covid-19 ARDS seems to swing between two forms: the traditional angioedema type as seen in SARS1/other viruses and something resembling a microthrombotic type of ARDS, kind of a "malarial pneumonia." The latter is driven by the RAS and angiotensin II (CD147, ACE D allele) and the former by the KKS and bradykinin (ACE2, ACE I allele). ACE2 receptors dominate via alveolar type 2 (AT2) cells before the blood-gas barrier, while CD147 receptors, up-regulated in the elderly, the obese, and the comorbid [11], are the primary determinants after the breach primarily via their receptors on lymphocytes, erythrocytes, platelets and endothelial cells. ACE2 receptors are not present on lymphocytes or erythrocytes and their presence on platelets [12] or endothelial cells [13] [14] is putative.

2.2. Covid Symptoms

Most exposed to Covid-19 have minimal symptoms (ACE I allele). Dry cough is one of its earliest signs and can indicate mild pulmonary edema [15]. In Covid-19 it is mediated by bradykinin [15] [16] [17] [18]. Anosmia, ageusia occur late and are good prognosticators [19] (KKS, ACE I allele). Dry cough and headache [19] are features of both early Covid (shorter clinical course) and long-haul Covid [20] (KKS, ACE I allele). It is another manifestation of bradykinin and the KKS. Estrogen down regulates ACE [21]. Brain fog and myalgias favors post Covid, non comorbid, non hospitalized Caucasian females [22] [23]. Indeed long-haul Covid may represent the new fibromyalgia/chronic fatigue syndrome [24]. Covid toes are also probably a skin manifestation of bradykinin. Bradykinin via B1R/B2R facilitates vascular permeability. ACE blocks BK access to B2Rs. Instead it is shunted to des-Arg9-bradykinin, which is metabolized by ACE2 (see Figure 2).

2.3. MIS-C, MIS-A, KD, TSS

ACE/ACE2 increases with age, accelerated by comorbidities [25] [26]. Without a comorbidity pediatric ARDS is rare [27]. AT2Rs (facilitate tight endothelial junctions) and seemingly B2Rs (increase solute permeability) decrease with age [28] [29] and seem to cancel each other [30] Amongst 104 11 - 17 year olds bradykinin was negatively and des-Arg9-bradykinin positively correlated with body mass index (BMI) (increased ACE). Des-Arg9-BK was also positively correlated

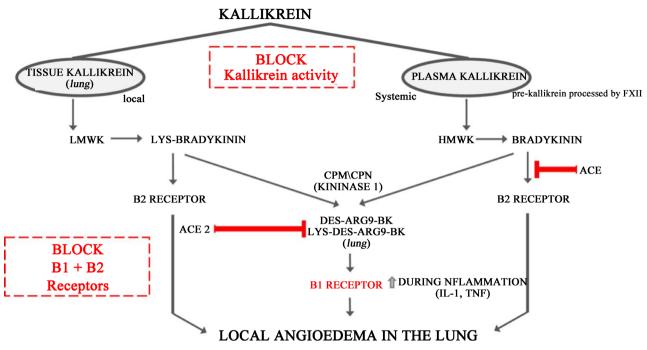


Figure 2. RAS and KKS are linked [3].

with systolic blood pressure (BP) [31]. Increasing pediatric BMI, diabetes, BP are burgeoning problems, especially amongst African Americans and Hispanics [32]. By correlating these age and gender related changes in children the KKS connection can be extrapolated to explain MIS-C [33] and MIS-A [34]. Children do not appear to express typical or Covid type ARDS, unless they have a comorbidity. But if their Th1/Th2 becomes imbalanced as seen in asthma/allergic atopy [35] or vitamin D deficiency [36], the balance can tip toward the KKS and higher risk [37]. The mucocutaneous erythema, edema, hypotension, and rash of MIS-C/MIS-A suggests the work of bradykinin and appealingly connects both KD and TSS [38] [39] to MIS-C. In MIS-A the slightly older (median age 21, predominantly male, noncomorbid, slightly overweight [40] status also seems symptomatically to represent a KKS response. CD147 and its glycan (sugar) shield interact with MBLs (Mannose Binding Lectins) and trigger the LCP (Lectin Complement Pathway) [41]. This ties the coronary arteritis and myocarditis of MIS-C/MIS-A to KD [42] [43] [44]. The staphylococcal and streptococcal toxins, although possessing no glycan shield, also invoke the LCP. Although myocarditis has been described in TSS, coronary arteritis has not. MIS-C/MIS-A, KD, and long haul Covid are probably best classified as post viral inflammatory/autoimmune states.

2.4. CD147

SARS2 in numbers removes ACE2 and increases ACE/ACE2 and Angiotensin II. Angiotensin II via AT1Rs activates TACE (Tumor necrosis factor Alpha Converting Enzyme) aka ADAM17 (A Disintegrin and Metalloproteinase 17) [45], which up-regulates TNF- α and IL6. TNF- α and IL-6 up-regulate CD 147 receptors [46] [47]. TNF-*a* up-regulates high mannose glycans [48] due to TNF-*a* inhibition of mannosidase in the ER (endoplasmic reticulum) [48]. N-glycosylation in the ER of the heavily glycosylated spike S protein CD147 epitope (and the host CD147 receptors) then favors the high mannose N-glycan (sugar) for their glycan shield [49]. This then brings us to the CD147 epitope present on the SARS2 spike protein S [50]. Two recent widely referenced articles [51] [52] claimed not present, but have been discredited [53]. The CD147 epitope, whether on the emerging virus or on the manufactured spike protein S, is heavily N-glycosylated. MBLs are a prominent component of innate immunity especially in the young. They bind the high mannose glycans that are shielding the CD147 receptors are also involved in the pathogenesis of coronary artery disease [54].

2.5. ACE D ALLELE

ACE D allele is more frequently encountered in African-Americans and others, e.g., Italians, Iranians, differentially targeted by Covid-19 [55] [56]. The ACE D allele is more frequently encountered in elite athletes specializing in power sports [57]. The ACE I allele frequency correlates with stamina but only in male athletes [58]. CD147 receptors on erythrocytes are up regulated in quarter horses (power) versus Arabian horses (stamina) [59]. If quarter horses are likened to elite power athletes with the DD genotype, might this up-regulation also apply? The myocarditis appears to be mediated by CD147 receptors on cardiac pericytes [60] and CD147 epitopes on spike S. MBLs remove "immune complexes". After all CD147 is a member of the immunoglobulin super family. MASPs (MBL Associated Serine Proteases) increase endothelial inflammation, permeability and angioedema (up-regulate bradykinin) [61]. This would explain the myocarditis and pericardial effusion complication post vaccination in power athletes.

Age, gender, race, weight have significant impact on Covid phenotype. Furthermore other less easily quantified inputs, e.g., inoculum dose, vitamin D status, other polymorphisms, ..., cannot be overlooked. The Covid-19 phenotype/ACE genotype linkage is strong but not absolute and not always direct. For example, elevated RDW (red cell distribution width) reflects spleen size and a poor prognosis. Perhaps pre-existing spleen size is also a negative determinant. The previous reference to increased CD147 receptors on erythrocytes in quarter horses indicated a close relationship between CD147 and MCT1 (monocarboxylate transporter). This relationship enables more efficient removal of lactic acid by erythrocytes. These erythrocytes can be stored in the spleen, as a reserve, when needed. Enlarged spleens due to chronic hypoxia are native to high altitude Sherpas and the free diving Bajau people (sea nomads) of Malaysia. ACE genotype does not dictate this, but the D allele does lead to higher ACE levels and more angiotensin II. This creates faster twitch, red fibers and gives its owner an advantage in power sports, which, if exploited with interval training, leads to more CD147-MCT1 receptors on erythrocytes stored in the spleen, as in quarter horses. Since the Covid phenotype seems unduly harsh for the anaerobically trained athlete, perhaps the same is true for Sherpas and the Bajau. Covid 19 has devastated Nepal at the top of the world. The free diving Bajau peoples comprise barely 10% of the Malaysian population yet over 50% of the Covid deaths. This is even more striking, given their sea nomadic lifestyle and probably excellent vitamin D status in contrast to city dwelling Malaysians and the D allele is rare. Splenic release of these super sensitized CD147 laden erythrocytes could result in myocarditis if encountering increased CD147 antigens (epitopes) on the spike protein S (recent viral or vaccine exposure). In addition splenomegaly offers another connection to KD and coronary artery changes [62].

3. T Cells and Glycans

Covid-19 related lymphopenia is primarily of T cells, especially cytotoxic CD147+CD8+ T cells [63]. CD4/CD8 is increased in Covid-19 v AIDS, where it is decreased. This has important implications for complications, discussed below. Th17, a CD4+ T cell, is a marker for autoimmune disease (see Figure 3, Figure 4) and elevated Th17/Th1 is strongly associated with COVID mortality [37]. It is activated by IL-1,6,23 and produces IL-17. It is elevated in KD [64], Macrophage Activated Syndrome (MAS), and associated auto immune diseases, e.g., Stills Disease [65] and Juvenile Idiopathic Arthritis. Th17 in particular is associated with coronary ectasia [66], seen in almost all of these auto immune diseases. Coronary arteritis in MIS-C, MIS-A, KD involves CD147 receptor antibodies and appears to involve pericytes [60]. Although KD exhibits a Th17

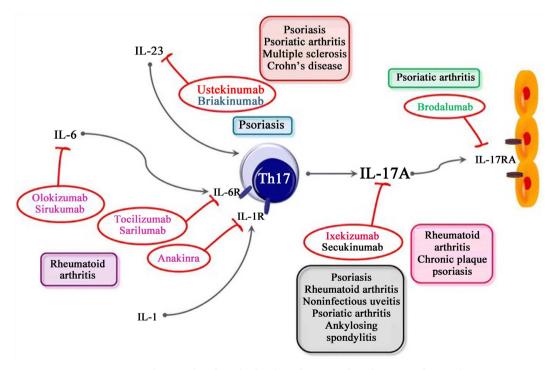


Figure 3. Many commercial monoclonal antibodies have been employed against Th17 and its autoimmune proclivities.

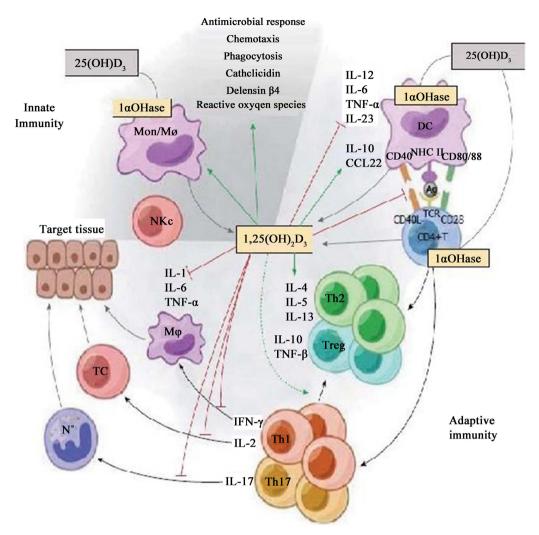


Figure 4. Vitamin D covers the action of all these Th17 related commercial monoclonal antibodies.

auto-immune type response, all KD patients with coronary aneurysms exhibited eosinophilia, a Th2 allergic type response [67]. So, TNF-alpha and IL-6, the two most critical cytokines to COVID-19 severity [68] [69], mediate both the MBL response and the Th-17 response. MASPs up-regulate bradykinin and cause cardiac angioedema [61]. This might explain the pericardial effusion that often accompanies the myocarditis in the young and otherwise healthy. These autoimmune and/or allergic effects are blocked by vitamin D (see Figure 4), which balances Th1/Th2 and suppresses Th17.

4. Lectins

Unfortunately this spike S/CD147 (BSG) interaction has dire consequences for those with comorbidities. Destruction of ACE2 receptor bearing cells by SARS2 increases ACE/ACE2. Although SARS2 may also enter cells after attaching to CD147 receptors [50], their high mannose glycan shield may be more problematic [49]. The RBD (Receptor Binding Domain) is extensively shielded from antibody recognition by its glycan cover, with the notable exception of the ACE2

receptor binding domain [70] [71]. This translates to plenty of ACE2 antibodies, but few CD147 antibodies. The poorly shielded ACE2 epitope provides easy access for antibody production. CD147 antibodies but not mRNA induced antibodies are protective against all the variants [72]. Vaccines trigger antibodies to the predominant strains, clearing the field and encouraging the rise of variants simple Darwinian survival of the most resistant, not unlike disruption of the normal balance of gut flora by antibiotics, thereby providing a gap for pathogenic microorganisms. Natural immunity is non-discriminating and attacks all strains old and new as they appear. Mutations are not granted special status and usually get diluted. Many recent studies have reported overwhelming superiority of natural immunity over vaccine immunity (Johns Hopkins and Israeli reports). This spike protein S, whether viral or vaccine related, presents plenty of high mannose glycans for the MBLs [73]. The efficacy of meplazumab (antibody against CD147) [74] and an LCP inhibitor [75] [76] lends additional mechanistic support. The glycan shield on the RBD of SARS2 triggers the LCP via circulating MBLs [77] [78]. Complement triggered by CD147 is also a prominent player in malaria [79]. Complement fixing MBLs are strongly correlated to plasma D-dimer levels, a marker of COVID19 coagulopathy [80] [81] [82]. IVM and other lectins (red algae [83]) bind and saturate not only these high mannose glycan shields but also the high mannose glycans on CD147 receptors on erythrocytes, platelets, lymphocytes, and endothelial cells, intermediating the complement and clotting cascades (microthrombotic and cardiovascular pathology). Lending credence to this hypothesis is an ongoing clinical trial involving Rhodophyta, a red algae. Rhodophyta (Gigartina) contain mannose specific lectins that bind spike glycoprotein specific to SARS-CoV2 to inhibit viral entry. This leaves MBLs inactivated and dormant. This would seem protective whether facing Covid-19 or its vaccines [84].

5. Vaccines (Pfizer and Moderna)

The blood-gas barrier seems to separate typical ARDS from microthrombotic type. Once the blood gas barrier is bypassed by the vaccine, the introduction of large numbers of CD147 bearing S spike-proteins accelerates the process in the susceptible. Vaccines bypass this barrier in a manner analogous to the bite of an Anopheles mosquito injecting P falciparum. Both the protozoan and the S protein attach to the CD147 receptors on erythrocytes, platelets, and endothelial cells, causing thrombosis and vasculitis. No replication required. Knowing Avogadro's number, the 30 microgram Pfizer/100 microgram Moderna load of mRNA with 150 kD molecular weight, one can calculate the number S units produced (12 trillion for Pfizer and 40 trillion for Moderna). S antigen was detected as early as day 1 post-vaccination, and peak levels were detected on average 5 days after the first injection. S in all participants declined and became undetectable by day 14 [85]. According to analysis of a Japanese biodistribution study of the Pfizer mRNA vaccine, the S1 subunit can be found in spleen, bone

marrow, the liver, adrenal glands, ovaries, heart and brain [86]. Each new administration represents a CD147/MBL booster and is the reason D-dimers are acutely slightly elevated post vaccination. Richard Hoffe, MD reported July 2021 mildly elevated D-dimers in 62% after 4-7 days in the 900 tested [87]. Vaccines create a mismatch between circulating spike protein S with their high mannose glycan shielded CD147 epitopes and host CD147 receptors on T cells, erythrocytes, and endothelial cells. Many otherwise healthy with enlarged spleens, e.g., athletes, Sherpas, Bajau people, may face dire consequences upon release of these CD147 receptor upregulated erythrocytes that are sequestered in the spleen. Vitamin D deficiency would compound this.

6. Covid-19 and Vaccine Complications

We know the short term risks of Covid-19, but those of its vaccines are less clear. What might be the intermediate and long term consequences?

6.1. Thrombosis

Clearly Covid-19 causes TMA (Thrombotic Microangiopathy) in those susceptible, i.e., the elderly, the obese, and the comorbid (RAS dominant). However, outside this group those with the ACE DD genotype and those with anaerobically induced splenomegaly are also at risk. The large number of vaccinees has highlighted the problem of thrombosis amongst the otherwise healthy. The AZ and JJ [88] vaccine adverse reactions appear to be more platelet activating and thrombotic and less cytotoxic, whereas the mRNA vaccine adverse reactions seem to favor the MBL pathway. The thrombotic reactions can be either HIT [89] or VITT. Heparin antibodies are present in about 5.4% of the population [90]. Thrombosis could also be triggered without heparin. After the first AZ/JJ dose CD147 epitopes on the inactivated virus could activate platelets. These activated platelets then release platelet factor 4. PF4 combines with endothelial heparan sulfate (a heparin analogue) to form complexes [91] on the endothelial cells (no exogenous heparin needed). These immunogenic complexes have a clearance time of about 50 days [89]. They could trigger thrombosis if Covid-19 exposure was recent or this could theoretically follow the second dose, depending on the timing. Instead of HIT, it's called vaccine-induced immune thrombotic thrombocytopenia (VITT) and involves primarily 20 to 50 year old Caucasian females [88]. This may be unrelated to the RAS/KKS balance. However, one can speculate about the role of birth control pills in this thrombotic process. The increase in PF4 seems to be mediated by the S spike protein [88] [90]. VITT post AZ/JJ vaccines is probably due to an antibody to the immunogenic PF4/heparan complex (not just PF4 [92] [93]). See Figure 5.

6.2. Immune Compromise

CD8+ T cells are selectively but not solely reduced by SARS CoV2, increasing CD4+/CD8+ [63]. In one study of 88 children with chicken pox (varicella) and

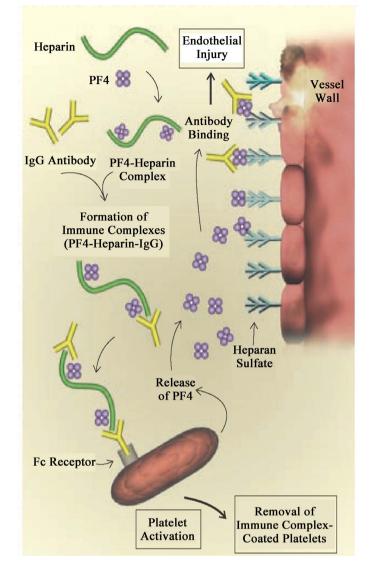


Figure 5. Endothelial heparan sulfate, like heparin, is an anticoagulant, but under certain circumstances it can become a pro-coagulant.

60 without, a significant decrease in the level of CD8 positive cells was found in those with viral DNA. CD4 differed insignificantly [94]. Reactivation of herpes zoster is often seen in the immunocompromised, e.g., those undergoing chemotherapy, the elderly, the stressed, ... Reactivation of herpes zoster (HZ) post Covid-19 has been described [95]. European EudraVigilance database reported 4103 cases of HZ after receiving the Pfizer vaccine [96]. This has also been seen after both the Moderna and inactivated viral vaccines as well [97]. CD4+ and cytolytic CD8+ T cell responses play an important role in controlling HZ replication [98] [99].

6.3. Cancer Acceleration

In addition to erythrocytes and platelets SARS2 and its vaccines impact T lymphocytes. T cell lymphopenia is well described in malaria and a poor prognosticator. CD147 receptors are expressed on CD4+ (T helper) and CD8+ (T cytotoxic) cells. CD8+ T cells are selectively but not solely reduced by SARS CoV2, increasing CD4+/CD8+ [63]. This is the inverse of HIV where the ratio decreases [100]. Both viruses utilize highly glycosylated CD147. Loss of CD8+ T cells translates to loss of control over progression of CA (growth, metastasis,...) [101] [102] [103] and CD147 is specifically involved. This is because CD4+ T cells monitor premalignant cells, *i.e.*, dysplasia and carcinoma in situ. CD8+ T cells suppress those that actually invade (loss of p53 function) [101]. In short loss of CD4+ T cells renders an individual susceptible to opportunistic infections. Loss of CD8+ T cells renders an individual susceptible to cancer recurrence. Presence of cytotoxic CD8+ expressing CD147 receptors limits cancers expressing CD147 antigens [102] [103] [104] [105]. This is why cancer is spiking post Covid 19 vaccine [106].

6.4. Autoimmune Disease and Antibody Dependent Enhancement (ADE)

The cause of autoimmune disease is multifactorial but appears to involve Th17 [107]. Vitamin D deficiency is clearly contributory [45]. Some consider autoimmune disease to be the longterm sequelae of a viral infection. Others have shown a distinct link with CD8+ T cell deficiency. So the low CD8+ T cell count (CD4/CD8 is increased) in SARS2 [108] is worrisome.

ADE was first seen with the Dengue virus about 40 years ago. It has also been described with Yellow fever, Zika, West-Nile fever, respiratory syncytial virus, influenza, measles, HIV, SARS, and MERS [109] [110]. CD8+ T cells can block Dengue [111] and Zika virus ADE [109]. The ADE potential for SARS2 has already been documented in the lab, using the Wuhan strain and the delta strain [112]. IVM [113] [114] [115], and Vitamin D [4] have much to offer.

7. Early Diagnosis

D-dimers, RDW, platelet, lymphocyte and eosinophil counts are good proxies for early diagnosis of the more lethal microthrombotic ARDS (angiotensin II dominant) or extra-pulmonary TMA. The CD147 epitope on the spike protein S interacts with CD147 receptor bearing erythrocytes. These erythrocytes, which now lack available CD147 receptors, are entrapped in the spleen [116], which itself enlarges. RDW parallels splenic size [117] [118] [119]. This same interaction between CD147 receptors on platelets (releasing PF4) and endothelial cells triggers the LCP with angioedema and thrombosis, producing the D-dimers and thrombocytopenia. The same process drives cerebral malaria [120]. ACE2 receptors are probably not involved. But the answer is probably moot [121]. All four cell types, erythrocytes, platelets, lymphocytes, and endothelial cells, conspire for a poor prognosis [119] [122] [123] [124] [125] [126]. All four are rich in CD147 with poor/no ACE2 receptor component. Covid-19 has had little impact in sub-Saharan Africa. There are only three plausible explanations for this and all three have been denigrated to some extent by mainstream medicine. 1) falciparum antibodies [127] [128]; 2) IVM cross coverage for malaria/river blindness [129]; 3) adequate vitamin D.

IVM is a horse dewormer! Many parasitic helminths, e.g., Onchocerca volvulus (cause of a filariasis known as river blindness), horse worms like Ascaris, ... employ glycan shields against bacteria as their larval or filarial forms migrate through the bloodstream. Ascaris employs an N-glycan shield [130]. P. falciparum's RH5 epitope is also heavily glycosylated. Given its efficacy for river blindness (Nobel Prize) and as a horse dewormer (not to mention anti-malarial), IVM may also bind these glycans, pre-empting MBLs without the inflammation and thrombosis.

Vitamin D levels were lower in north and south Africa compared with sub-Saharan Africa, in urban areas compared with rural areas, in women compared with men, and in newborn babies compared with their mothers [131]. Amongst Brazilians over 50 years 25% were vitamin D deficient and 62% were vitamin D insufficient [132]. In one unpublished study 88% of Filipinos were vitamin D deficient (<20 ng/ml) or insufficient (<30 ng/ml). In India 80% - 90% were deficient [133]. Adequate vitamin D counters the development of either phenotype [4]. IL6 and severity/mortality are inversely and independently related to Vitamin D levels. Adequate vitamin D counters MIS-C and MIS-A. IL-6 and TNF- α are the prime cytokines of COVID-19. TNF- α inhibits mannosidase, increasing high mannose glycans on the RBD of the spike protein S [48] and IL-6 up-regulates CD147 [47]. These oligomeric glycans involve not only the RBD on circulating S protein of viral or vaccine origin but also the native CD147 receptors. They are upregulated by cytokines, especially IL-6, interact with MBLs, which initiate the LCP and thrombosis, and accelerate atherosclerosis [134] [135].

8. Therapy

There have been many studies evaluating the efficacy of vitamin D for prevention and therapy of COVID19. The vast majority have been positive [136] [137] [138] [139]. Those that have been less favorable, when reviewed, have obvious errors in structure, e.g., low levels accepted as sufficient, insufficient time for adequate levels to develop, ... Vitamin D is helpful in the vaccinated and the unvaccinated. Vitamin D also addresses the influenza risk. Using 2017-18 data from the CDC website [140] one can calculate the NNT for the flu shot to be about 40. The NNT for vitamin D during the winter (less than 25 nmol/L or 10 ng/ml) is just above 4 [141] (see **Figure 6**). NNT is the number needed to treat to eliminate one case of the flu. This translates to a vitamin D flu efficacy nearly an order of magnitude greater than that of the flu shot.

Vitamin D deficiency rickets was first described in the 1920s. Vitamin D resistant magnesium deficient rickets wasn't described until 50 years later in the 1970s (see Figure 7). Vitamin D deficiency can also impact allergy and autoimmune

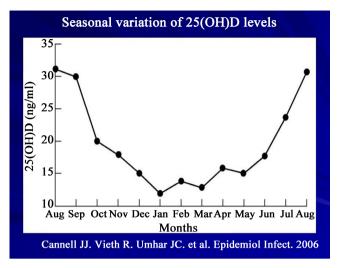


Figure 6. During the flu season in the northern tier of the US vitamin D levels are less than 15 ng/ml.

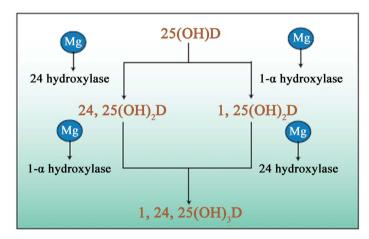


Figure 7. Magnesium is critical in producing the active form of vitamin D [142].

risks [143] [144] [145].

9. Conclusion

The ACE I/D polymorphisms seem to predict which Covid-19 ARDS variant will manifest in the few that do develop ARDS. A preliminary CD4/CD8 might provide some insight into susceptibility. Those without comorbidities that are KKS inclined generally escape severe ARDS, while those with comorbidities that are RAS mediated upregulate TNF- α and IL6. CD147 then takes center stage. D-dimers, RDW, lymphocyte, platelet and eosinophil counts should help in early diagnosis. There are numerous considerations, e.g., vitamin D status, polymorphisms, age, gender, inoculum dose, ..., that impact the clinical course. Vitamin D, IVM, red algae might assist in prevention and treatment of symptoms, whether vaccinated or not. In summary circulating MBLs attack the glycan shield on the CD147 epitope. This supercharges CD147 induced damage and triggers microthrombosis. In the susceptible consequences can be devastating.

Long-term impact of Covid-19 is unknown. But the commonality of CD147 between the virus and the vaccine raises some disturbing possibilities, including compromised immune function, autoimmune disease, accelerated progression of cancer, and ADE. There are many trillions of spike protein S units created by each mRNA dose/booster. CD147 epitopes are on each S unit, despite flawed articles claiming otherwise. This article constitutes a red alert on the lurking dangers that are becoming more apparent. Warning: Cough not required!

Conflicts of Interest

The author declares no conflicts of interest.

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