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Research Article

A Cohort of Cancer Patients with No Reported Cases of SARS-CoV-2 Infection: the Possible Preventive Role of Methylene Blue

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Abstract. We report the case of a cohort of 2500 French patients treated among others with methylene blue for cancer care. During the COVID-19 epidemics none of them developed influenza-like illness. Albeit this lack of infection might be by chance alone, it is possible that methylene blue might have a preventive effect for COVID-19 infection. This is in line with the antiviral activity of Chloroquine, a Methylene blue derivative. Both Chloroquine and Methylene blue have strong antiviral and anti-inflammatory properties probably linked to the change in intracellular pH and redox state.

Keywords. COVID-19, cancer, methylene blue, metabolic treatment.

Introduction

Europe has been recently hit by an epidemics of COVID-19. We report a cohort of patients treated for cancer in France. This cohort is managed by an association (Espoir Métabolique) and is a cancer support group. There are 2500 patients all at high risk for sepsis because of concomitant chemotherapy. One of us has interviewed (by telephone and by e mail) these patients to register the cases of COVID-19. As of March 27th 2020, there were no cases of registered COVID-19 or of flu-like syndroms. These patients were treated by a combination of standard therapy and α -lipoic acid (800 mg twice a day), hydroxycitrate (500 mg three times a day) and methylene blue (75 mg three times a day) as well as a low carb diet.

There were 52% women against 48% men. The most prevalent cancer type were breast cancer (40 %), lung (20%), prostate (10%), uterine (10%), colon (8%), liver (6%) miscellaneous (6%). Albeit this lack of influenza-like illness might be by chance alone, it is possible that one of these molecules might have prevented viral infection. Herein, we present a short scientific survey of what is currently known about the SARS-Cov-2 virus, hydroxychloroquine treatment as well as biochemical properties of methylene blue that was called once upon a time a “magic bullet” for healing a wide range of diseases.

Background

Coronaviruses (CoVs) are quite common viruses that are generally related in humans to the upper and lower respiratory tract family of disorders. They may trigger asthma in children and adults and severe respiratory disease in the elderly. They could also be responsible of pneumonia and bronchiolitis infections in the infant and child population. The first human coronaviruses was discovered in 1965 and named B814.¹ Shortly after this discovery, other coronaviruses were described that caused disease in multiple animal species, including, rats, mice, chickens, turkeys, calves, dogs, cats, rabbits and pigs.² In the late 1960s, two major human strains were studied HCoV-OC43 (“OC” meaning that they could be grown in organ cultures such as mouse brain) and HCoV-229E (“E” meaning that such viruses were ether-sensitive suggesting that they required a lipid-containing coat for infectivity), a strain that could be grown in tissue culture directly from clinical samples. Epidemiologic studies found that coronaviruses were endemic in humans, being responsible for 5-10% of all upper and lower respiratory tract infections associated to a quite low pathogenicity. But the situation changed in 2002-2003 after the discovery in Southern Asia of a new respiratory illness, termed Severe Acute Respiratory Syndrome (SARS), which was able to spread quickly in 29 countries of America, Asia and Europe causing 774 fatalities and resulting in a mortality rate of 9%.³ This SARS-CoV was responsible for a loss of \$40 billions in economic activity. Phylogenetic studies seemed suggesting a bat origin for this new SARS-CoV virus with an entry in the human population probably relayed by Himalayan palm civets. The strange thing was that 40% of wild animal traders and 20% of individuals, who slaughter animals were seropositive for SARS, without manifestation of any symptoms, meaning that the real causes of the disease remain still nowadays quite unclear. Infection control policies were able to halt the epidemics in 2004.

But the same year, a new strain named HCoV-NL63 was identified in the Netherlands from an infant with bronchiolitis, followed by a new strain HCoV-HKU1 in

2005 from a patient with pneumonia in Hong Kong. These new strains were however not able to trigger new epidemics in the human population. This was however not the case of the MERS-CoV (Middle East Respiratory Syndrome Coronavirus) isolated in 2012 from a patient with pneumonia in Saudi-Arabia that was able to spread in 21 countries with almost 600 related deaths and a mortality rate of \approx 40%. Again, bats were suspected of being at the origin of the virus, but with a new intermediate that were dromedary camels as natural hosts. It is worth noticing that since 1965, HCoV-229E, HCoV-OC43, HCoV-NL63, HCoV-HKU1 and MERS-CoV are commonly circulating in the human population causing general respiratory illness and cold symptoms during winter and spring months in healthy individuals and more severe disorders in the immuno-compromised and the elderly. We are now facing a new terrible challenge, with the emergence in late November 2019 of a new strain name SARS-CoV-2 in Wuhan, Hubei province, China. Concerning the origin of the virus, genomic studies have shown that Malayan pangolins (*Manis javanica*) illegally imported into Guangdong province contain coronaviruses similar to SARS-CoV-2 with as usual bats serving a reservoir hosts.⁴ Another possibility could be natural selection in humans following zoonotic transfer. But genetic data irrefutably show that SARS-CoV-2 is not derived from any previously known virus backbone, meaning that it is improbable that it emerged after laboratory manipulation of a related SARS-CoV-like coronavirus. Nowadays, CoVs belong to the *Nidovirales* order split into two subfamilies (*Coronavirinae* et *Torovirinae*) and further subdivided into four groups: α -CoVs (HCoV-229E and HCoV-NL63), β -CoVs (OC43, HKU1, SARS, MERS), γ -CoVs and δ -CoVs.

Genome and structure

SARS coronaviruses are medium-sized RNA viruses, holding inside an oily shell a 30 kilobases long single stranded RNA-genome, positive in sense, with a 5' cap structure ending with a 3' poly-adenylated tail. The virions appear to spherical in shape with a diameter of about 125 nm and several club-shape spike projections emanating from the oily surface, conferring to the virus its corona-like aspect (figure 1). The overall packing of the coronavirus genome is 5' – leader – UTR - Replicase (ORF1ab) - S (Spike) - E (Envelope) - M (Membrane) - N (Nucleocapsid) – 3'UTR – poly (A) tail with accessory genes interspersed within the structural genes at the 3' end. UTR corresponds to untranslated regions and ORF to open reading frames. The replicase gene occupying two-thirds of the genome encodes for nonstructural proteins (Nsps) through two polyproteins (pps) 1a (coding for nsps1-11) and 1ab (coding for nsps1-16) with the following identified functions:⁴

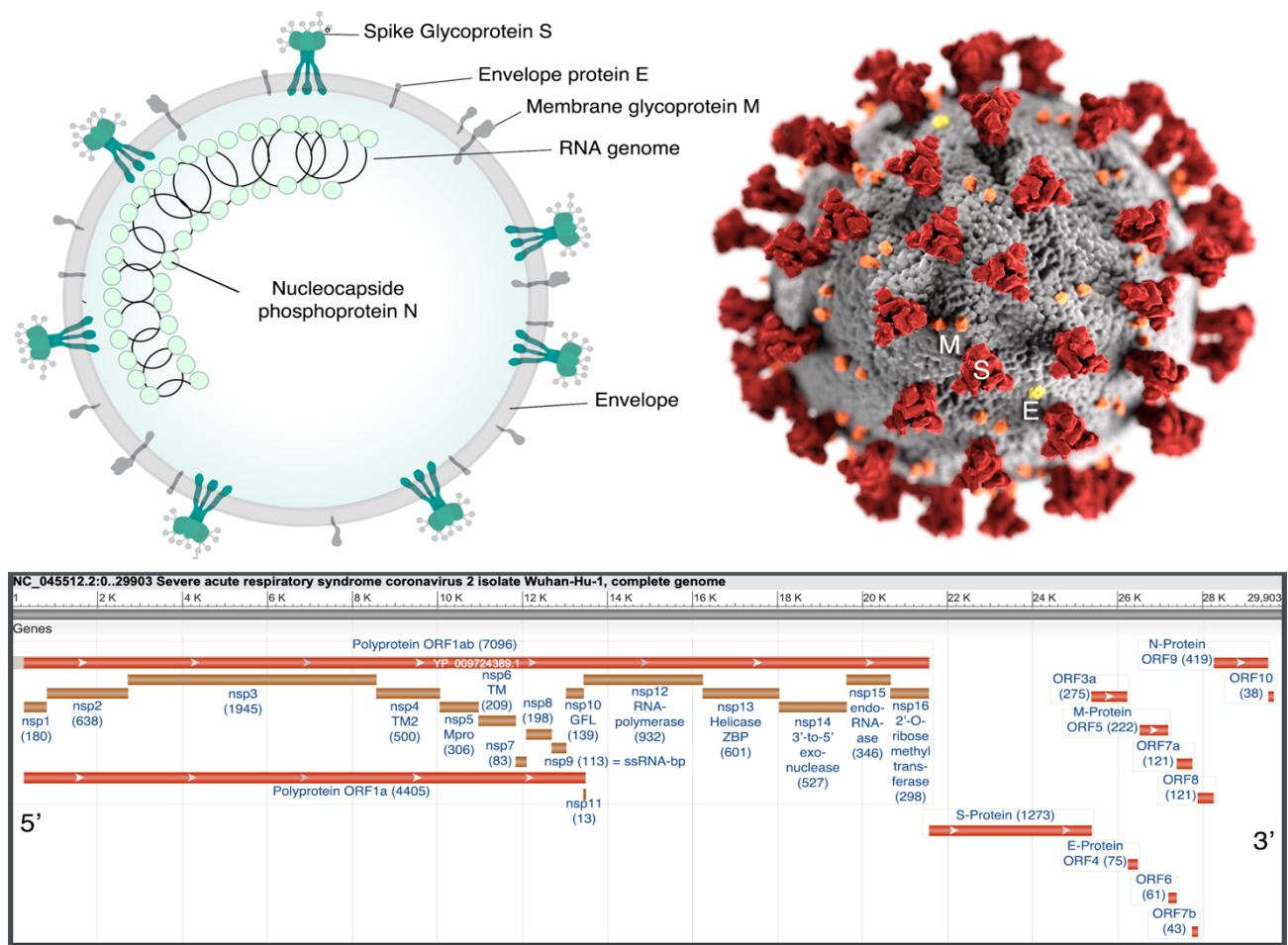


Figure 1. Structure and genome of SARS-CoV-2 responsible for the Covid-19 epidemics. Number in brackets gives the number of amino acids of each encoded protein.

- Gene nsp1 promotes cellular mRNA degradation and blocks host cell translation resulting in blockage of innate immune response.
- Gene nsp2 have currently no known function.
- Gene nsp3 encodes for a large multi-domain trans-membrane protein with ubiquitin-like and acidic domains that interacts with N-protein, ADP-ribose-1''-phosphatase (ADRP) activity that promotes cytokine expression, papain-like protease (PLPro) with deubiquitinase domain, responsible for cleavage at nsp1/2, nsp2/3 and nsp3/4 boundaries in order to block host innate immune response.
- A potential trans-membrane scaffold protein playing an important role for proper structure of double-membrane vesicles (DMVs) in encoded in gene nsp4.
- A serine type main protease (Mpro) in nsp5 gene is responsible for all cleavage events not mediated by PLPro.
- Genes (nsp7, nsp8) encodes for two hexadecameric complexes that may act as processivity clamp for RNA polymerase.
- A RNA binding protein is encoded in gene nsp9.
- Gene nsp10 encodes for a cofactor that forms heterodimer with nsp16 and nsp14 for stimulating activity of the corresponding proteins.
- Gene nsp11 from pp1a extended into pp1b becomes nsp12, encoding the RNA-dependent RNA polymerase (RdRp) that duplicates viral RNA. The recombination ability of coronaviruses during viral evolution is tied to the switching ability of RdRp during replication.
- A RNA helicase with RNA 5'-triphosphatase activity insures unpackage of the viral genome (gene nsp13).
- Gene nsp14 encodes an Exoribonuclease (ExoN) that insures replication fidelity (proofreading of the viral

genome) with N7-methyltransferase (N7-MTase) activity for adding 5' cap to viral RNAs.

- A viral endoribonuclease (NendoU) with unclear function is encoded in gene nsp15 and is a genetic marker (together with nsp14-ExoN) for the order *Nidovirales*.

- Finally gene nsp16 encodes for a 2'-O-methyltransferase (2'-O-MT) that shields viral RNA from MDA5 (melanoma differentiation associated protein 5) recognition.

Once cleaved, genes behave as mRNA for the ribosomal units of the infected cell in order to produce a set of Nsps that are able to self-assemble into a replicase-transcriptase complex (RTC) providing a suitable environment for producing both genomic and sub-genomic RNAs. The role of sub-genomic RNAs is to serve as mRNAs for the structural and accessory genes which resides downstream of the replicase polyproteins. Following replication and sub-genomic RNA synthesis, the viral structural proteins S, E and M are translated and inserted into the endoplasmic reticulum (ER) for further processing by the endoplasmic reticulum-Golgi intermediate compartment (ERGIC). After encapsidation of the viral genome by the N-proteins in the ERGIC, budding with viral structural proteins (first with M and E, then with S later on) leads to mature virions. Following assembly, virions are transported to the cell surface in vesicles for further release by exocytosis.

If some S-proteins remain outside the virions, they may also transit towards the surface for mediating cell-cell fusion between infected cells and adjacent uninfected cells, leading to giant multinucleated cells, responsible for virus spreading without detection or neutralization by virus-specific antibodies. Accordingly, the primary determinant for infection of a cell by coronaviruses is the attachment of the virion through non-covalent interactions of protein-S with a suitable receptor. For human coronaviruses it is known that HCoV-OC43 binds to N-acetyl-9-O-acetylneuraminic acid, HCoV-HKU1 binds to O-acetylated sialic acids, while HCoV-NL63 and SARS-CoV bind to heparin sulfate proteoglycans.⁵ After binding to the cell, coronaviruses use a broad variety of fusion receptors: aminopeptidase N (APN) for HCoV-229E, human leucocyte antigen molecule (HLA class I) or sialic acids for HCoV-OC43, angiotensin-converting enzyme 2 (ACE2) for HCoV-NL63 and SARS-CoVs and dipeptidyl-peptidase (DPP4) for MERS-CoV. The binding receptor of HCoV-HKU1 remains unknown. ACE2 receptors are expressed by epithelial cells of the lungs, intestines, kidneys and blood vessels, with a substantial up-regulation in patients with type 1 or 2 diabetes or hypertension, who are treated by ACE inhibitors and angiotensin II type-I receptor blockers (ARBs).⁶

Increased expression of ACE2 is also observed by

thiazolidiniones and ibuprofen. It has thus been inferred that people using ACE2-stimulating drugs may have a higher risk of developing severe and fatal COVID-19.

Chloroquine and hydroxychloroquine

Quite recently, a French team led by Pr. Didier Raoult in Marseille, has reported that it was possible healing in less than a week COVID-19 patients after administration of an anti-malaria drug, hydroxychloroquine (HCQ) and an antibiotic, azithromycin (figure 2).⁷ It is worth noticing that antibiotics are generally of no use against viruses, but could be nevertheless useful in order to prevent severe respiratory tract infections in patients suffering from viral infection. Such results are obviously very promising as the mean duration of viral shedding in patients suffering from COVID-19 in China was at least 20 days and up to 37 days.⁸ In vitro studies has suggested that a possible mechanism of action of chloroquine (CQ) and HCQ could be the augmentation of the intracellular pH in acidic organelles such as endosomes and lysosomes, owing to the fact that both molecules are weak bases.⁹ Accordingly, it is known that acidic media (pH < 5) are mandatory for endosome maturation and function. CQ was thus reported to elevate the pH of lysosome from about 4.5 to 6.5 at 100 μ M. Consequently, it could be surmised that endosome maturation might be blocked at intermediate stages of endocytosis, resulting in failure of further import of virions into the cytosol. Another possibility could be inhibition of SARS-CoV entry by CQ or HCQ through their ability of changing the glycosylation state of ACE2 receptors and S-proteins. Accordingly, it has been checked using immunofluorescence analysis (IFA) and confocal microscopy that the transport of SARS-CoV-2 from early endosomes (EEs) to endolysosomes (ELs) required for the release of the viral genome, was blocked (*in vitro*) by CQ and HCQ.⁹ Another point concerns the high concentration of cytokines (IL1B, IFN γ , IP10, MCP1, MIP1A, TNF α) in the plasma of critically ill patients infected by SARS-CoV-2. As HCQ is a successful anti-inflammatory agent that has been extensively used in autoimmune diseases, one may anticipate its ability to decrease the production of cytokines and pro-inflammatory factors. It is also worth noting that if HCQ is less toxic than CQ, prolonged and/or overdose usage of both molecules may lead to poisoning with cardiovascular and renal complications. There is thus an obvious need of finding other much less toxic molecules. Another constraint should be the low price and the large-scale availability of these molecules as on Saturday march 28, the COVID-19 has affected 177 countries, with more than 600'000 confirmed cases and about 28'000 deaths all around the world.¹⁰

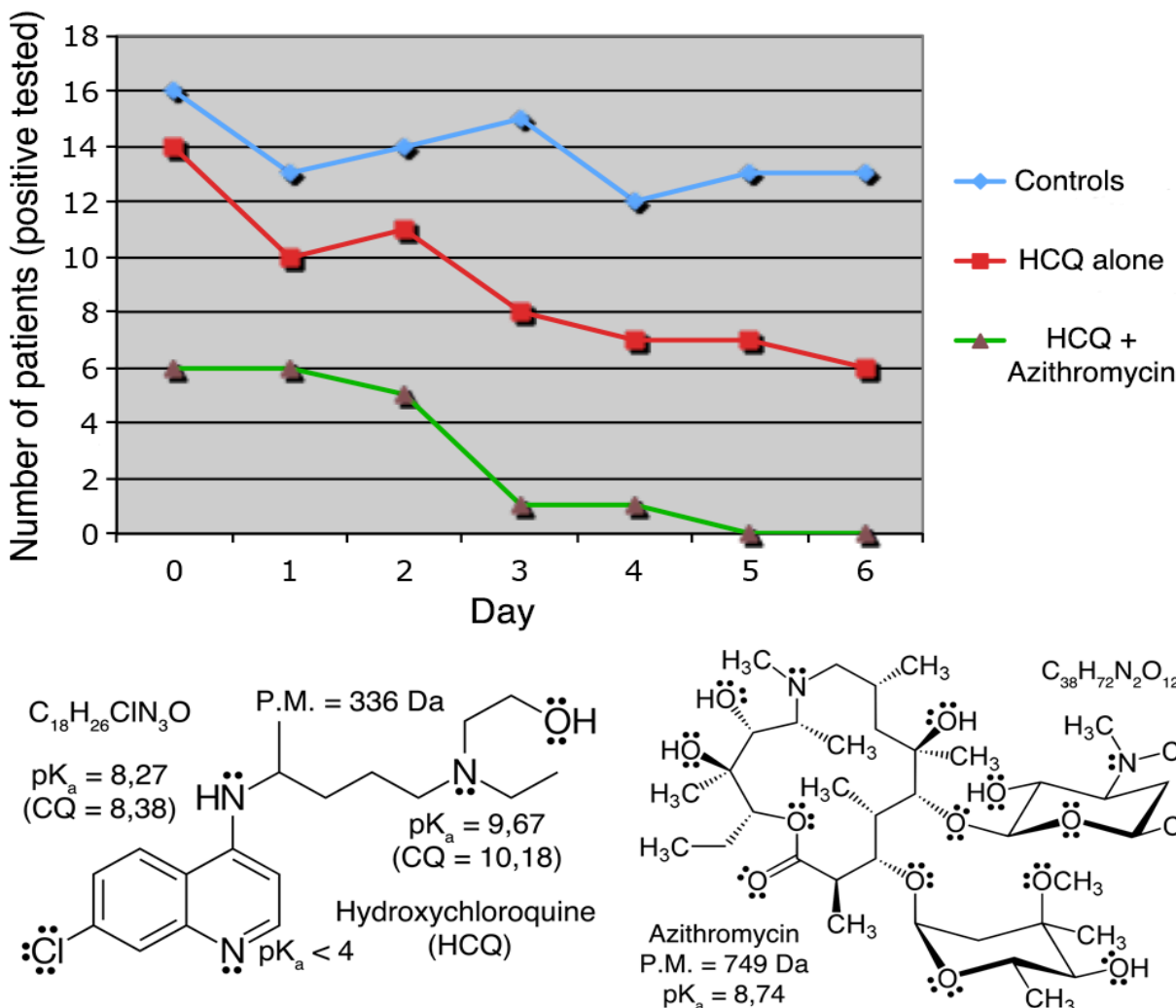
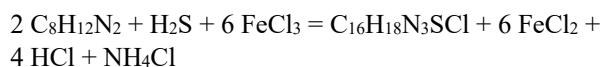


Figure 2. Effect of a treatment against COVID-19 based on daily administration of 3×200 mg of hydroxychloroquine sulfate with addition of azithromycin if necessary. Study made on 20 hospitalized patients (age > 12 years) with PCR documented SARS-CoV-2 carriage in nasopharyngeal samples at admission. Adapted from reference 7.

Methylene blue (MB⁺)

Cancer is another kind of disease able to lead to cytokines storms and one may expect a large number of deaths from COVID-19 in such patients. However, a survey among our database of patients treated with a combination of α -lipoic acid, hydroxycitrate and methylene blue suggests that this treatment prevents from severe infection by COVID-19. It may thus be anticipated, but yet not proved, that MB⁺ could be of considerable help for fighting against the COVID-19 epidemics. Here, we give a survey of the very interesting properties of this molecule with emphasis on chemistry, clinical trials being currently under investigation. Moreover, it is worth noting that methylene blue is the ancestor of modern anti-malaria drugs such as chloroquine and is associated to a lesser toxicity, the only drawback being a green-blue coloring of urine.

Methylene blue chloride is an old compound synthesized in 1876 by the German chemist Heinrich Caro, through oxidation of a mixture of dimethyl-4-phenylene-diamine $Me_2N-Ph-NH_2$ and hydrogen sulfide by ferric chloride:¹¹



The sulfur atom bridges two molecules of the *p*-phenylene-diamine backbone (figure 3), forming a phenothiazine heterocyclic molecule displaying a formally positive thionium ion in one mesomeric form. Through aromatic resonance among the three fused rings, this formal positive charge may be delocalized over the two nitrogen atoms of the right and left dimethyl-amino groups leading a

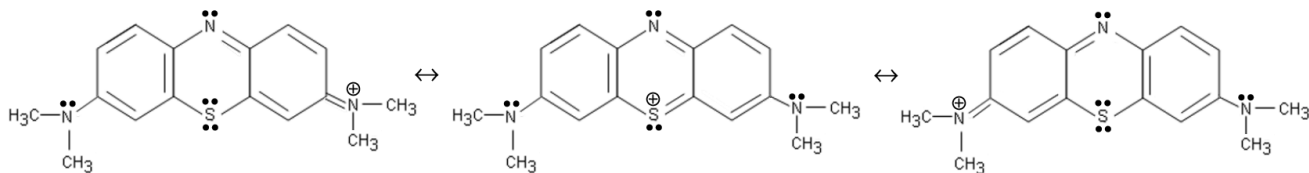


Figure 3. Methylene blue MB⁺ and its three mesomeric forms.

characteristic absorption at $\lambda = 663 \text{ nm}$ ($\epsilon = 75 \text{ mM}^{-1} \cdot \text{cm}^{-1}$) for the un-protonated cation, undergoing a red-shift at $\lambda = 740 \text{ nm}$ in HMB²⁺ after protonation.¹² Owing to strong absorption of the red part of the visible spectrum, cationic forms of methylene blue are deep-blue colored, a property used in 1882 by Robert Koch for staining the tubercle bacilli and extensively used by Paul Ehrlich (1854-1915) for differentiation between the different types of white blood cells.^{13,14} The chemical structure of methylene blue was established in 1884 and it was used in 1887 in combination with eosin by the Polish pathologist Czeslaw Checinski for evidencing presence of daisy-like (*Plasmodium malariae*) and sickle-shape (*Plasmodium falciparum*) parasites in blood smears. In 1891, Ehrlich discovered that methylene blue fell in the category of “magic bullets” drugs for its ability to target malarial organisms. Replacement of quinine, a natural substance derived from the *cinchona* tree of South America available in very limited supply, by methylene blue, a costless synthetic dye has allowed large-scale production of antimalarial drugs. Subsequent developments of what has been called at that time “chemotherapy” has led to the synthesis in 1931 of the very successful drug quinacrine, marketed by Bayer under the names of mecaprine and atrabine. Atrabine was further modified in 1934 by the German chemist Hans Andersag by replacing the acridine ring with a quinolone ring, giving access to a product named “resochoin” upon reaction of oxaloacetic acid diethylester with *m*-chloroaniline. But, the product was found to be too toxic for practical use in humans and in order to minimize toxicity, the compound 3-methylresochoin, named “sontochin” was synthesized and patented in November 1939, after testing over 1’100 patients with malaria. In November 1945, E. K. Marshall rediscovered resochoin giving to the compound its definitive name “chloroquin”, becoming the first-line antimalarial therapy for about 20 years, saving countless lives. Later on, hydroxychloroquine was developed and reported to be half as toxic as chloroquine.¹⁵ It follows that methylene blue may be considered as a template for the synthesis of substitutes of quinine in the cure of malaria. As shown in Table 1, methylene blue is in fact in the top-five drugs against 18 stains of *Plasmodium falciparum*.¹⁶ Moreover, a recent paper has evidenced that parasites responsible for malaria may also be hosts of yet unidentified RNA viruses,¹⁷ reinforcing the idea that it may exist a strong link between RNA viruses and anti-malaria drugs.

There is also recent evidence that methylene blue activated

with visible light effectively reduce Ebola Virus (EBOV), MERS-CoV, SARS-CoV,

Crimean–Congo hemorrhagic fever virus (CCHFV) and Nipah virus (NiV) infectivity in platelets and plasma, respectively.^{18,19} It follows that methylene blue used in conjunction with light may open a quite novel route of antiviral therapy by mixing biochemistry with photo-physics. The need for large-scale testing of methylene blue against COVID-19 is again reinforced.

Methylene blue in non-viral pathologies

Methylene blue is in fact what may be called a BONARIA drug.²⁰ Here the three letters “BON” means that it is a safe and efficacious remedy while the last letters indicates that it is also affordable (A), registered (R) and internationally accessible (IA). This comes from the quite peculiar chemical properties of this substance. As shown in figure 4, the behavior of methylene blue as a function of pH, redox potential and irradiation is quite diversified and fascinating.^{12,21} Upon assimilation, it is not a single molecule that enters blood circulation, but a full set of molecules, explaining the extreme versatility in a large number of pathologies ranging from microbiology to psychiatry. Concerning viral infections, it is worth noticing that when methylene blue undergoes a one-electron reduction, it becomes a neutral lipophilic MB[•] radical with good stability insured by its delocalization over several mesomeric forms. Moreover, such a radical acts as a weak base ($\text{pK}_a \approx 9$) that could favor, as with chloroquine, transient alkalization of cytosolic spaces.

Methylene blue also possesses antibacterial activity and is secreted in urine, explaining why it was heavily used for treating infections and painful disorder of the urinary tract in multi-ingredient prescriptions (polypharmacy).²² A quite useful combination was 5.4 mg of MB, 0.03 mg of atropine sulfate and 0.03 mg hyoscyamine (for pain relief of smooth muscle spasms), 40.8 mg of methamine (condensation product of formaldehyde with ammonia, breaking down in acidic urines), 5.4 mg of benzoic acid and 18.1 mg of phenyl salicylate (salol, an antiseptic). The mechanism of action of methylene blue against parasites has been partially elucidated.

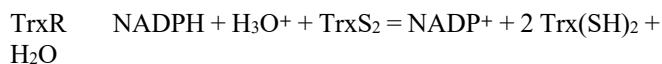
Drug	Abbreviation	<IC ₅₀ >/ nM	Min / nM	Max / nM
Dihydroartemisinin	DHA	0,72	0,1	5,08
Artesunate	AS	1,35	0,12	5,53
Lumefantrine	LMF	4,1	0,61	82,9
Proveblue	BM	5,3	0,9	40,2
Pyronardine	PND	5,8	0,4	19,7
Monodesethylamodiaquine	MDAQ	9,8	1,9	119,1
Mefloquine	MQ	30,1	7,1	63,4
Piperaquine	PPQ	32,2	2,5	168,0
Chloroquine	CQ	52,2	6,1	346,4
Quinine	QN	63,1	6,2	1430

Table 1. Ex vivo susceptibility of 18 Plasmodium falciparum isolates from Dakar.¹⁶ Proveblue is a methylene blue preparation that complies with the European Pharmacopoeia and contains limited organic impurities and heavy metals of recognized toxicity.

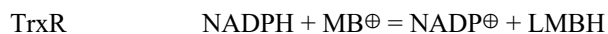
It involves homodimeric flavoenzymes of the glutathione (GR) reductase family that are present both in malarial parasite and the mammalian host cell.²⁰ The first affected enzyme is glutathione reductase (GR) allowing reducing glutathione disulfide (GSSG) to the sulfhydryl form glutathione (GSH):



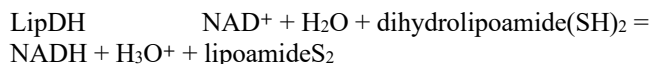
The second enzyme is thioredoxin reductase (TrxR) allowing reduction of thioredoxins (Trx) that are proteins facilitating the reduction of other proteins by cysteine thiol-disulfide exchange:



The role of GR and TrxR is to keep GSH and redoxins in the reduced state in order to maintain cytosolic spaces under reduced conditions. The key point is that methylene blue could be a substrate for TrxR with production of the reduced neutral and colorless form (leuco-methylene blue or LMBH) absorbing in the UV-part of the electromagnetic spectrum ($\lambda = 340 \text{ nm}$, $\epsilon = 3.3 \text{ mM}^{-1} \cdot \text{cm}^{-1}$ and $\lambda = 258 \text{ nm}$, $\epsilon = 17.4 \text{ mM}^{-1} \cdot \text{cm}^{-1}$):



It follows that methylene acts as an inhibitor of the natural reactions of TrxR. It is worth noticing that methylene blue is unable to inhibit the enzyme dihydrolipoamide dehydrogenase (LipDH), another flavoprotein enzyme that oxidizes dihydrolipoamide to lipoamide (functional form of α -lipoic acid):



However, the reduced form LMBH is unstable at micromolar concentrations of molecular oxygen and auto-oxidizes readily under such conditions as shown in figure 5. The antioxidant thiol-producing enzymes guarding the reducing milieu of cytosolic spaces are thus turned in contact with methylene blue into pro-oxidant H₂O₂-producing enzymes challenging the reducing milieu that they are meant to protect.

It follows that methylene blue is both an inhibitor and a subversive substrate allowing concomitant production of reactive oxygen species (ROS). Under such conditions, NADP(H) and molecular oxygen that are needed for the pathogen's metabolism are irreversibly consumed by methylene blue. In addition, there is less GSH available in the parasite as a substrate of GSH S-transferase for the detoxification of hemes and other lipophilic compounds. The anti-bacterial effect of methylene blue then lies in the fact that human GR reacts more slowly with methylene blue than parasites GR and that human TrxR is not present in erythrocytes. Accordingly, when parasitized red blood cells and normal erythrocytes are incubated together in MB-containing solution, the drug becomes concentrated selectively in the parasitized erythrocytes. A possible reason is that as LMBH bears no electrical charge, it easily permeates the membrane of digestive vesicles and is then auto-oxidized back to MB⁺, thus remaining trapped in the vesicles. It is finally worth noticing that synergistic effects have been found against *P. falciparum* in culture when using methylene blue in combination with artemisinin derivatives.²⁰ In the 1920s methylene blue proved also to be a dramatic antidote for carbon monoxide or cyanide poisoning.²³ Consequently, methylene blue could be very efficient since 1940 for treatment of methemoglobinemia, a pathology where the ferrous ion of hemoglobin becomes oxidized into ferric ion, impairing attachment of dioxygen and thus reducing the oxygen-carrying capacity of the blood.²² Upon IV-injection, methylene blue (1-2 mg·kg⁻¹ or 1% sterile solution) transforms in contact with reductases in the erythrocytes into the colorless *leuco*-methylene blue (LMBH) able to reduce methemoglobin back to normal hemoglobin.

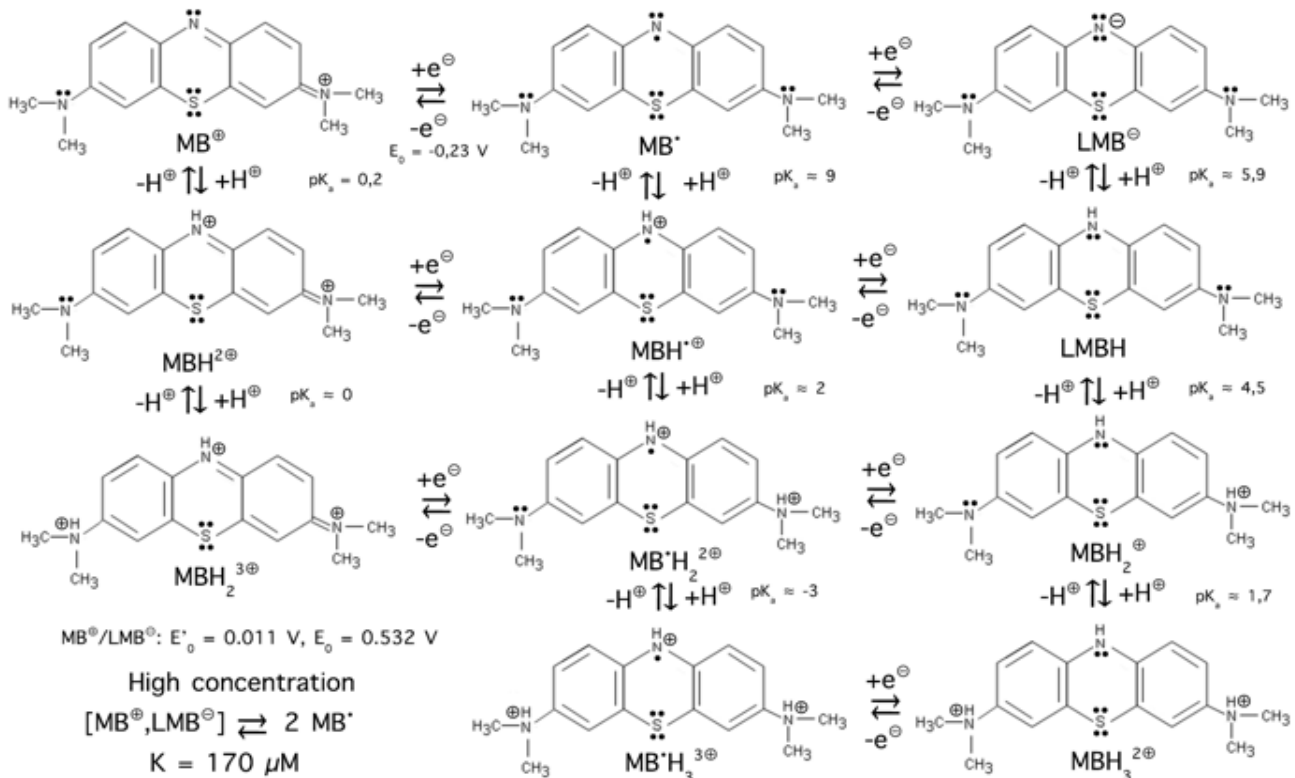
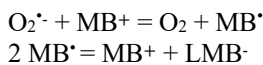


Figure 4. Methylene blue speciation as a function of pH and redox potential.^{12,21}

It follows that if used as such rather low concentrations, methylene blue functions as an alternative electron carrier in mitochondria, which accepts electrons from NADH or FADH₂ and transfers them to CoQ or cytochrome c and bypassing any complex I/III blockage (figure 6).²⁴ It is worth noting that in this case, a harmless product, water, is produced instead of hydrogen peroxide (cf. figure 5).

As a side effect, methylene blue is also able to scavenge any leakage in superoxide anion O₂^{•-} from the ETC according to the reaction:



Methylene blue is thus a potent antioxidant able to block the oxidative cascade at its very beginning. It may thus also be considered as pyromaniac firefighter able to increase oxidative stress by accelerating ATP production and also able to eliminate the same oxidative stress thus generated. Associated to these well-known antioxidant properties is the ability of methylene blue for attenuating any ischemia/reperfusion injury through inhibition of superoxide generation by xanthine oxidase.²⁵ The ability of methylene blue for minimization of free radical production in the mitochondrial ETC explains why it has positive effects under metabolically stressed conditions, such as ischemic brain injury, where excess free radicals may lead to cellular damage and cell death. Accordingly, *in vivo* studies have shown that methylene blue reaches its maximum concentration in blood by 5 min after

intravenous administration in humans.²⁶ The half-life of MB in blood after intravenous administration is 5.25 h in humans. No significant effects on vascular reactivity were observed using functional magnetic resonance imagery (fMRI) but a preferential potentiation of regions with a higher metabolic demand has been evidenced. Consequently, methylene blue concentrates in cortical regions with the largest metabolic energy demands in an activation task such as forelimb stimulation, boosting intellectual tasks. Using methylene blue in patients with septicemia, leads to positive results owing to a direct inhibition of nitric oxide synthases (NOS), both constitutive and inducible. It inhibits guanylyl cyclase (GC) by binding to the heme group of the enzyme and blocks the catalytic functions of NO synthase by oxidation of the enzyme-bound ferrous iron.²⁷

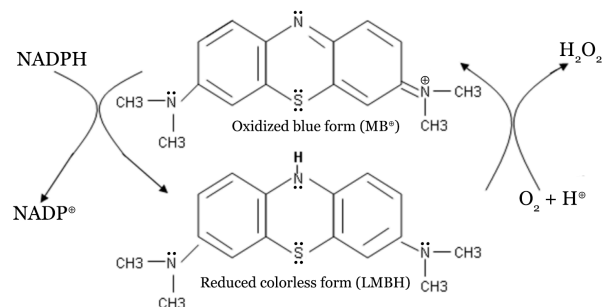


Figure 5. Methylene blue as a scavenger of NADPH and O₂ with production of hydrogen peroxide H₂O₂.

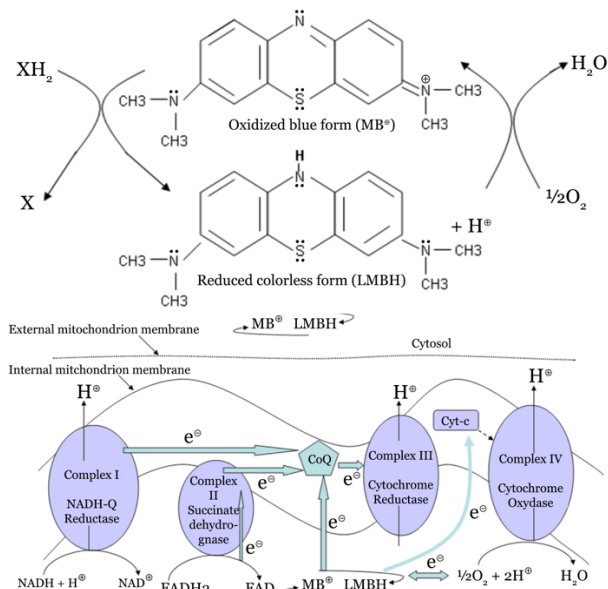


Figure 6. Methylene blue as an electron carrier within the electron transport chain (ETC) present in mitochondria.

An additional effect on soluble guanylyl cyclase, which normally leads to the formation of cyclic guanosine monophosphate (cGMP) as the second messenger of NO[•], adds to the inhibition of the NO-cGMP pathway. Methylene blue may be a more specific and potent inhibitor of NO synthase than guanylyl cyclase, because direct NO-donating compounds in the presence of methylene blue can still partially activate cGMP-signaling pathways. Methylene blue also inhibits platelet activation, adhesion, and aggregation synergistically with an inhibition of platelet thromboxane A2 and endothelial prostacyclin I2 production.²⁷

A very important point is that if methylene blue is able to increase complex-IV activity by about 30%, it also has a marked hormetic action by capturing electrons of the ETC at high dose (> 10 mg/kg) and through interaction with nitrogen oxide synthase (NOS) may produces cardiovascular effects.²⁶ Studies performed *in vitro* has evidenced that complex-IV achieves a maximum activity for a 0.5 μM concentration. Above 5 μM, complex-IV activity is inhibited, and the larger the concentration, the stronger the inhibition.²⁸ On the other hand, studies performed *in vivo* on rats has shown that a maximal locomotion activity was reached at a dose of 4 mg·kg⁻¹, no effects being observed below 1 mg·kg⁻¹ or above 10 mg·kg⁻¹. Finally, above 50 mg·kg⁻¹, locomotion activity becomes to be reduced.

Summing together all these properties explains why methylene blue could be of considerable use in neurodegenerative diseases²⁹, ageing³⁰, cancer³¹ or for healing psychic disorders.³² Figure 7 shows for instance how methylene blue may act on key enzymes involved in Alzheimer disease (AD). Beneficial effects of methylene blue against AD has been demonstrated in clinical studies and comes from the non-polar character of the reduced

LMBH form allowing it to cross easily the blood-brain barrier, for hitting multiple molecular targets. Concerning the inhibition of Tau protein aggregation by methylene blue, an *in vitro* target seems to be the microtubule affinity-regulating kinase (MARK4) at Ser262, through stabilization of its dimeric form following cysteine oxidation. But, as shown in figure 7, this is just one of the mode of action, the other ones being the down-regulation of cholinesterase activity for preventing acetylcholine (ACh) degradation and the ability of scavenging superoxide. But research in this field is progressing quite rapidly. For instance, it was recently shown that methylene blue reverses Caspase-6-induced cognitive deficits by inhibiting Caspase-6, and Caspase-6-mediated neurodegeneration (inhibition of the cleavage of the amyloid precursor protein APP) and neuroinflammation.³² As Caspase-6-mediated damage seems to be reversible months after the onset of cognitive deficits suggesting that methylene blue could benefit Alzheimer disease patients by reversing Caspase-6-mediated cognitive decline. As shown in figure 7, methylene blue is also an inhibitor of monoamine oxidases (MAOs), a biochemical fact that has been known for many decades.³² Figure 8 gives more details about the antipsychotic effects of methylene blue that have been exploited for more than a century. Accordingly, many studies have confirmed that methylene blue influences neuronal communication by altering cholinergic, monoaminergic, and glutamatergic synaptic neurotransmission both in the central and the peripheral nervous systems. First, methylene blue is able to induce neuronal membrane depolarization with inhibition of Ca²⁺-activated K⁺ channels, activation of Ca²⁺ channels, and facilitation of Na⁺ channel inactivation.³² It also modulates the functions of various integral membrane proteins involved in transports of solutes such as glucose and ions such as Na⁺, K⁺, and H⁺. It was demonstrated that the cGMP pathway does not mediate the actions of methylene, reported in the majority of these studies. Both glutamate and dopamine have been implicated in the pathogenesis of psychoses and methylene blue has been the lead compound for the development of classical antipsychotics. Thus, promethazine was made in the 1940s by a team of scientists from Rhône-Poulenc laboratories and shares with LMBH the same phenothiazine backbone and was the first-generation urgeon Henri Laborit in the hope of discovering a more effective anesthetic. While the drug did not cause his patients to lose consciousness, it did induce a remarkable calmness. This discovery was the very beginning of the “chemical lobotomy” revolution for the treatment of both acute and chronic psychoses, including schizophrenia and the manic phase of bipolar disorder, as well as amphetamine-induced psychosis. Considering the similarities in the chemical structures of methylene blue and antipsychotics, it is likely that methylene blue modulates the activity of dopamine receptors.

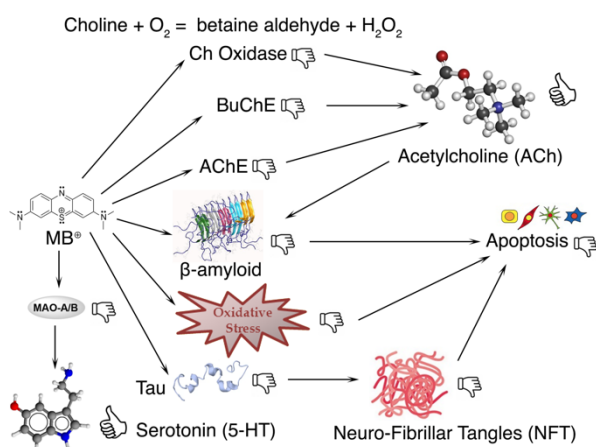


Figure 7. Methylene blue as an inhibitor of several enzymes involved in Alzheimer's disease.

And owing to its antioxidant properties, it could also be of considerable help in the prevention of Li-toxicity in bipolar disorder patients.

Among other interesting properties, methylene blue has been shown to modulate the physiological actions of hormones involved in the hypothalamo-pituitary-peripheral axis, increasing thyroid peroxidase activity and enhancing the iodination of thyronines, with subsequent increases in the synthesis of thyroxine.³² Moreover, when exposed to light, MB becomes photosensitized, leading to the release of cytotoxic, highly active, and short-lived oxygen-derived species such as singlet oxygen ¹O₂. It has also been shown that photons in the red-to-near-infrared frequency range of approximately 620–1150 nm penetrate to the brain and intersect with the absorption spectrum of cytochrome oxidase.³⁴ While low-dose methylene blue and low-level near-infrared light may produce different pleiotropic cellular effects, both interventions cause a similar up-regulation of mitochondrial respiration with similar benefits to protect nerve cells against degeneration. These human studies suggest that low-dose methylene blue may have potential therapeutic applications in neurology as a neuroprotective agent, and in psychiatry and clinical psychology to facilitate psychotherapeutic interventions. Similarly, low-level near-infrared light improved human neurological outcome after ischemic stroke,³⁵ and could help in conjunction with methylene blue to enhance emotional and neurocognitive functions such as sustained attention and working memory in humans.

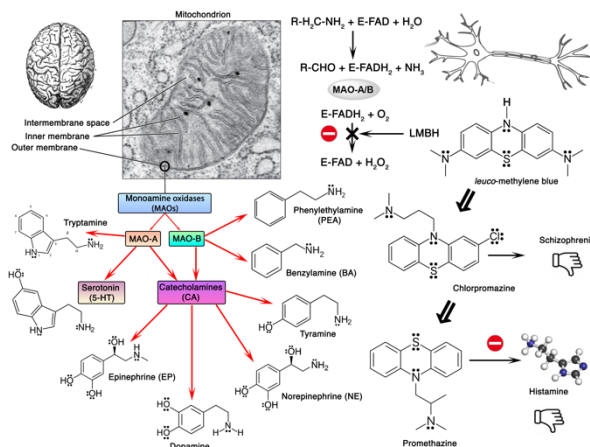


Figure 8. Leuco-methylene blue (LMBH) quickly crosses the blood-brain barrier and at daily dose of 15 mg is a MAO-A inhibitor boosting serotonin (5-HT) and increasing levels of catecholamines in the brain with thus reduced anxiety and depression and concomitant improvement of memory (increased levels of ACh) and mood

Conclusions

With the current COVID-19 outbreak, the world is facing a challenge and every possibility of helping people should be considered. Our preliminary data suggest but do not prove that Methylene blue might be a good treatment for influenza-like illnesses. Both Methylene blue and its derivatives such as chloroquine may share similar mechanism of action. Time is ripe for a prospective randomized clinical trial for the treatment of this dreadful disease.

Old drugs that have been tested for other indications (such as methylene blue or chloroquine) have a well-defined safety profile. They are often more effective than new drugs from the High Tech. The COVID-19 epidemics like cancer have a solution. Repurposing of known molecules will help cure these deadly disease

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