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Review of Phytochemical Compounds as Antiviral Agents against Arboviruses from the Genera Flavivirus and Alphavirus

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Abstract

Arboviruses are a diverse group of viruses that are among the major causes of emerging infectious diseases. Arboviruses from the genera flavivirus and alphavirus are the most important human arboviruses from a public health perspective. During recent decades, these viruses have been responsible for millions of infections and deaths around the world. Over the past few years, several investigations have been carried out to identify antiviral agents to treat these arbovirus infections. The use of synthetic antiviral compounds is often unsatisfactory since they may raise the risk of viral mutation, they are costly and possess either side effects or toxicity. One attractive strategy is the use of plants as promising sources of novel antiviral compounds that present significant inhibitory effects on these viruses. In this review, we describe advances in the exploitation of compounds and extracts from natural sources that target the vital proteins and enzymes involved in arbovirus replication.

Keywords: Arbovirus; Flavivirus; Alphavirus; Antiviral; Natural compounds; Plant products.

Introduction

Arbovirus (Arthropod-borne virus) is an epidemiological term used to describe diverse viral families that are transmitted to vertebrate hosts by hematophagous arthropod vectors [1]. Based on isolation of these viruses from vectors in nature, mosquitoes are the main vector in the transmission of arboviruses, although other biting flies, ticks, midges, and sand flies may also transmit the viruses [2, 3]. Most arboviral diseases are zoonotic and are infections which occur in vertebrates (monkeys, bats, rodents, birds, domestic animals, reptiles, and amphibians) and humans are an incidental host [3, 4]. Arboviruses have re-emerged as a serious public health problem worldwide. Today, this array of viruses is one of the major causes of emerging epidemic infections [5]. At present, there are over 534 viruses listed in the International Catalogue of Arboviruses, 134 of which are pathogenic for humans. However, only a few cause extensive epidemics with high mortality [6, 7]. According to reports on the ICTV (International Committee on Taxonomy of Viruses), most human pathogenic alphaviruses belong to one of three viral families: *Flaviviridae*, *Togaviridae* and *Bunyaviridae*. Most of these arthropod-borne viruses belong to the family *Bunyaviridae* but many important human arboviruses from a public health perspective belong to the families *Togaviridae* and *Flaviviridae*. However, there are many other human arboviruses belonging to the *Bunyaviridae* family [2, 4, 8]. About 30 alphaviruses and 53 flaviviruses have been identified, nearly one-third of which are clinically significant human pathogens [9, 10]. The viruses that cause substantial infections in humans are arboviruses. The wide variety of these viruses limit a review of the complete range of examples however, Table 1 lists the arboviruses that produce substantial infections in humans. Arboviral infections in humans and livestock can eventuate in a broad spectrum of clinical syndromes ranging from mild acute febrile infections with fever, to severe haemorrhagic fever, neuroinvasive diseases such as encephalitis and encephalomyelitis, and fatal disease [3, 8, 11]. The large number of diseases caused by arboviruses, such as dengue fever, tend to be epidemic with about 390 million cases occurring per year [12]. Chikungunya virus has recently spread throughout all regions of America and resulted in more than 2.9 million confirmed and suspected cases of infection, with approximately 300 deaths in the summer of 2016 [13].

The recent pandemic of Zika virus is therefore not an exception, and shows the risk potential of arboviruses that are implicated in human disease while the persistent challenges of dengue, Japanese encephalitis, chikungunya, and tick-borne encephalitis emphasize that people are at risk of these emerging and re-emerging viral diseases, although the mortality risk of many arboviral diseases is relatively low [11, 14]. Furthermore, the fast spread of arboviruses, especially in tropical and less developed regions can be considered as an important factor. This rapid spread results from a variety of factors, such as an increased human population, climate change, urbanization, globalization, and increased intercontinental travel which impose negative effects on the control of vectors [5, 15]. According to the literature, development of new diagnostic tools, antiviral therapies and efficient vaccines, is urgently required. Despite all the above factors, there are currently no effective antiviral therapies against the majority of these viruses, although approved vaccines exist for some flaviviruses such as yellow fever, Japanese encephalitis, and tick-borne encephalitis [7, 16, 17]. In the case of dengue, after several decades of endeavor, there is no effective vaccine or therapy for dengue virus, but several vaccines are currently in clinical trials. The first vaccine was recently authorized for use, but it bestowed only limited protection against four serotypes of dengue virus. Lifelong immunity against one serotype was induced, but immunity affecting the other DENG serotypes lasts only a few months. Moreover, there is no known efficacious therapy or vaccine for chikungunya virus [18, 19, 20]. In addition, despite extensive studies from the scientific community, currently there is no applicable arboviral therapy, though nonsteroidal anti-inflammatory drugs are recognized to suppress the replication of some arboviruses *in vitro*. However, *in vivo* research has been limited to a few rodent models, consequently, finding an effective treatment against these viruses represents a milestone in this field [5, 21, 22].

Plant-derived compounds can be used to contribute towards the design of structure-based drugs and plants are the main source of diverse natural compounds such as saponins, flavonoids, alkaloids, sterols and many other components. The antiviral properties of these biologically active compounds have been validated on the basis of their antioxidant-, antimicrobial- as well as immune-stimulatory properties, plus inhibition of viral DNA and RNA synthesis, inhibition of viral entry and other therapeutic activities [23, 24]. These agents are generally of low toxicity and in most cases, do not cause adverse effects. The success of natural compounds in the treatment of several diseases such as malaria, cancer and HIV has attracted research attention in the context of these agents against arboviruses [25, 26]. Clearer identification of natural antiviral compound modes of action is crucial to driving future research into antiviral treatment and to more effective viral control. Several compounds isolated from natural sources have been studied in relation to their inhibitory effects on different stages of the viral life cycle [24]. Accordingly, this review will focus on studies concerning phytochemical compounds possessing therapeutic potential as antiviral agents against arboviruses.

Table 1 – Clinically significant arboviruses, vectors, reservoir hosts, symptomology and geographic distribution

Genus	Virus	Vector	Reservoir host	Characteristic symptoms (in human clinical cases)	Geographic Distribution	References
Flavivirus	Dengue 1,2,3,4	In enzootic cycle: arboreal <i>Aedes spp.</i> In epidemic urban cycle: <i>A. aegypti</i> and <i>A. albopictus</i>	humans, primates	Fever, arthralgia, myalgia, rash	worldwide in tropics	[27, 28, 29,]
	Yellow Fever	<i>Aedes</i> and <i>Haemogogus spp.</i> (in urban cycle: <i>A. aegypti</i>)	humans, primates	Fever, hemorrhage, Jaundice	Africa, South America	[15, 30]
	Zika	<i>Aedes spp</i>	Primates	Fever, arthralgia, myalgia, rash. Infant microcephaly following infection in pregnancy	SE Asia, Africa, S and N America, Pacific Islands	[15, 27, 30, 31]
	Japanese encephalitis	<i>Culex spp</i> (especially <i>C. tritaeniorhynchus</i>)	Birds (Swine as secondary amplification host epizootic cycle)	Fever, encephalitis	Asia, Pacific	[1, 29, 31, 32]
	West Nile Fever	<i>Culex</i> species (especially <i>C. pipiens</i>)	Birds	Fever, arthralgia, myalgia, rash	Africa, Asia, Europe, North America	[29, 30, 32]
	Tick-borne encephalitis	<i>Ixodes</i> ticks	birds, rodents, domestic animals	Encephalitis	Europe, Asia, North America	[1, 15, 30]
	Murray Valley encephalitis; Kunjin	<i>Culex annulirostris</i>	Birds	Encephalitis	Australia, New Guinea	[1, 29, 31]
Alphavirus	Chikungunya	<i>Aedes spp</i> (in epidemic urban cycle: <i>A. aegypti</i>)	humans, primates	Fever, arthralgia, myalgia, rash	Africa, South Asia, Philippines, Indian Ocean islands, Caribbean, North and South America	[1, 27, 33]
	Eastern equine encephalitis	<i>Culiseta melanura</i> , <i>Coquilletidea perturbans</i> , <i>Aedes spp</i>	Birds	Encephalitis	Eastern and Gulf coasts of USA, Caribbean islands, Central America, North-East South America	[15, 31, 33]

Western equine encephalitis	<i>Culex tarsalis</i> <i>Aedes melanimon</i> <i>Aedes dorsalis</i> <i>Culex. (Mel.) spp.</i>	Birds	Encephalitis	Mid-West and Western USA, Canada	[1, 15, 31, 32]
Venezuelan equine encephalitis	<i>Aedes, Mansonia,</i> <i>Culex spp.</i> (epizootic subtypes)	Horses	Encephalitis	Venezuela, Colombia, Peru, Ecuador	[1, 31, 32]
O'nyong-nyong	<i>Anopheles funestus</i> <i>Anopheles gambiae</i>	Unknown	Fever, arthralgia, myalgia,	Africa	[1, 30, 31, 32]
Roos River	<i>Culex annulirostris,</i> <i>Aedes polynesiensis</i> <i>Aedes vigilax</i>	humans, marsupials	arthritic joint pain, fatigue and rash fever	Australia, South Pacific	[1, 15, 34]
Sindbis	<i>Culex. Spp,</i> <i>Culiseta sp.,</i> <i>Aedes sp.</i>	Birds	Fever, arthralgia, myalgia, rash	Asia, Africa, Australia, Europe,	[15, 30, 33]

Flaviviruses

The genus *Flavivirus* (here denoted as flaviviruses) is a group of arthropod-borne viruses that belongs to the *Flaviviridae* family and comprises nearly 75 viruses (53 species); of these, about 50% are associated with medically important infections that cause significant morbidity and mortality [1, 35]. This diverse group of arboviruses include: Dengue virus (DENV), West Nile virus (WNV), Japanese encephalitis virus (JEV), yellow fever virus (YFV), Zika virus (ZIKV), tick-borne encephalitis virus (TBEV), Murray Valley encephalitis virus (MVLEV) and many other viruses. These viruses are the most prevalent arboviruses worldwide and infect a significant number of people each year around the world. DENV, JEV, YFV and TBEV are the most dangerous pathogens from the genus *flavivirus* giving rise to mortality rates up to 30% [6, 36, 37]. The diseases caused by flaviviruses tend to fall into one of three categories (1) mild illness accompanied by fever, rash, and arthralgia, (2) encephalitis/meningitis, and (3) life-threatening hemorrhagic fever. Possible overlap may occur between these categories - for example, WNV generally induces a febrile illness with rash, although the minority of infections result in encephalitis or other neurological complications [1, 38].

Flavivirus molecular biology and replication

The mature flavivirus virion is small, icosahedral, enveloped, spherical and 500 Å in diameter [35, 39]. From a pathogenesis and therapy perspective, recognition of FLVs life cycle and replication in host cells with precision and resolution is of great importance. The first step in the life cycle of flaviviruses in the host cell is attachment of the virus E glycoprotein to the host cell surface receptor and cell entry via clathrin-mediated endocytosis [1, 40, 41]. Then the endocytic vesicle carries and transmits the flavivirus to endosomes. The low-pH environment within endosomes causes fusion of the viral membrane with the endosomal membrane mediated by molecular events within the E glycoprotein which result in release of the nucleocapsid into the cell cytosol and commencement of viral replication in the cytoplasm [41, 42]. Following the release of the viral genome into the cytoplasm, Polyprotein processing into component proteins takes place through a unique process at the surface of the endoplasmic reticulum (ER) where the created micro-environment allows the viral life cycle to continue while preventing interferon response signaling [41, 43, 54]. The replication complex (RC) is organized on ER membranes where negative-sense RNA is copied from the genomic RNA template. The RNA genome then interacts with C protein, which buds via the glycoprotein-containing ER membranes as immature viral particles into the lumen. Finally, these immature viruses in the process of maturation driven by the ER and Golgi complex, are released as mature virions from the cell [1, 54]. The flavivirus genomic RNA is single-stranded with positive polarity and the viral genome is approximately 11 kb in length, varying depending on the species. The RNA genome contains the 5' untranslated region (UTR), a single long open reading frame (ORF), and a 3' UTR [44, 45, 46]. The 5' UTR is typically capped, but at the 3' end the genome lacks a poly-A tail - instead there is a 3'-hairpin loop. The flanking UTRs are known as functionally active structural RNA elements that possess an important function in viral

replication, viral protein translation, virion assembly and immune modulation [1, 46, 47]. In infected cells, replication starts with the translation of ORF by the host cell machinery that produces a polyprotein precursor that needs post-translational processing by virus-encoded serine protease embedded in the N-terminal domain of non-structural protein 3 (NS3Pro) and host signals and furin proteases cleave the polyprotein into ten proteins including three structural proteins encoded toward the 5'-end of the genome (capsid, pre-membrane or membrane, and envelope) whereas seven non-structural (NS) proteins are encoded toward the 3'-end (NS1, NS2A, NS2B, NS3, NS4A, NS4B, and NS5) [44, 48, 49]. The three structural proteins are combined into the virion, whereas the non-structural proteins NS1-NS5 are used to coordinate the intracellular aspects of viral replication, virion assembly and modulation of host defense responses. The main role of NS proteins is viral replication, but several other functions have been identified for these proteins [48]. Among the NS proteins, NS1 is a highly conserved protein essential for flavivirus replication and inhibition of the complement-mediated immune response. This protein is recognized to activate the TLRs and is probably involved in negative-strand synthesis by an unknown mechanism [50]. NS3 is the second largest viral protein and has multi-enzymatic activity in the viral life cycle, including serine protease together with the cofactor NS2B, RNA triphosphatase necessary for capping nascent viral RNA, Nucleoside 5' triphosphatase (NTPase), and RNA helicase activities [51, 52]. NS5 has RNA methyl-transferase and RNA-dependent RNA polymerase (RdRp) domains necessary for viral replication and capping of nascent RNA. NS3 and NS5, together with the viral genome and cofactors plus host cell cofactors, assemble the virus replication complex (RC) on the intracellular membrane [46, 53]. NS2A, NS4B and NS4A are non-enzymatic membrane-associated proteins. NS2A, a small hydrophobic transmembrane protein, is essential for replication of the virus and is involved in the generation of membranes induced by the virus during virus assembly, whereas the viral NS2B protein acts as cofactor for the NS3 protease function. NS4A induces the rearrangement of the membrane and autophagy to amplify viral replication, while NS4B restrains the host immune response by stopping α/β interferon signaling and the helicase activity of NS3 [1, 48, 51, 54].

Alphaviruses

Alphaviruses are small, spherical, lipid-enveloped, and vector-borne viruses of the *Togaviridae* family. These virions are 70 nm in diameter and constitute more than 30 viruses with a broad host range, one-third of them being clinically important to humans [55, 56, 57]. They are universally divided into "Old World" and "New World" viruses according to the geographical region where these viruses were initially isolated. "Old World" alphaviruses, such as chikungunya (CHIK), Semliki Forest virus (SFV), O'nyongnyong viruses (ONNV), Sindbis virus (SINV), Mayaro virus (MAYV), and Ross River (RRV) are found in Africa, Asia, and Europe. Infection by these alphaviruses is commonly associated with febrile illness and polyarthralgia or polyarthritis [58, 59]. It has been estimated that millions of people are infected with these viruses annually worldwide. Recently, Chikungunya virus spread to South, North, and Central America resulting in more than 2.9 million suspected and confirmed cases which were responsible for about 300 deaths in the summer of 2016 [60]. "New World" alphaviruses include Venezuelan equine encephalitis virus (VEEV), Eastern equine encephalitis virus (EEEV) and Western equine encephalitis virus (WEEV), which are found in North and South America. These virulent pathogens are strongly related to potentially lethal encephalitis in humans and other mammals [58, 59].

Alphavirus genome and replication

These viruses consist of a single-stranded, positive-sense RNA ((+)RNA) genome. The alphavirus genome is approximately 11.8 kb long and contains two open reading frames (ORFs) [55, 61]: a 7 kb open reading frame encoding a polyprotein. This polyprotein is cleaved into 4 different nonstructural proteins (nsP1, nsP2, nsP3, and nsP4) in the 5' two-thirds of the viral genome and a 4 kb frame that encodes the terminal 3' region of the genome encoding the structural polyprotein in the remaining one-third of the alphavirus genome [62, 63, 64]. The structural polyprotein is cleaved into 5 structural proteins including a nucleocapsid protein (C), two small cleavage products (E3 and 6K) which do not exist in all alphaviruses, and two major envelope glycoproteins (E1 and E2). These E1 and E2 glycoproteins are important components in viral replication [21, 63, 65]. The E1 glycoprotein is necessary for viral fusion with the host cell and the E2 glycoprotein facilitates attachment of the virion to cell surface receptors and entry of viral particles to the host cell through clathrin-dependent endocytosis [55, 60, 66]. The nonstructural proteins play key roles in various stages of genome replication and are essential for the transcription and translation of viral mRNA in the host cell cytoplasm. The non-structural protein 1 (nsP1) is a mRNA capping enzyme that functions as a guanine-7-methyltransferase (MTase) and guanylyltransferase (GTase), where they add a methylguanosine cap to newly synthesized genomic and sub-genomic viral RNAs [58, 65, 67]. The resulting cap structure is necessary for translation of viral mRNA. Non-

structural protein 2 (nsP2) is a multifunctional protein that possesses several enzymatic functional roles such as protease, helicase, and nucleoside triphosphatase (NTPase) activities [55, 57, 64]. This nsP2 protein has also been characterized as a virulence factor that shuts off transcription and translation in infected host cells and inhibits interferon (IFN) mediated antiviral responses contributing to the control of the translational system, by viral factors [65, 68].

Of all alphavirus proteins, the function of nonstructural protein 3 (nsP3) is the least characterized. This protein is a necessary component of the replication complex (RC) and is highly phosphorylated. It has three known domains: a N-terminal macrodomain with nucleic acid binding capability and phosphatase activity, a C-terminal hypervariable domain, and an alphavirus unique domain (AUD) [21, 62, 67]. It has been demonstrated that deletion of the C-terminal hypervariable domain in SFV nsP3 leads to low viral pathogenicity, highlighting its importance in the regulation of viral RNA transcription. The final linear non-structural protein, NsP4, is the most highly conserved protein in alphaviruses and it is an RNA-dependent RNA polymerase [21, 57, 61]. Studies on antiviral strategies against alphaviruses generally target the E1 and E2 proteins responsible for viral binding to the cell during endocytosis and nsP proteins as the replication machinery. Old world alphaviruses prevent transcription of host proteins by their nsP2, whereas new world alphaviruses deploy an alternative mechanism to inhibit transcription of the host which is mainly defined by their capsid protein (CP). Accordingly, CP could be a significant target protein for potential antivirals [57, 68, 69]. Viral CP has several functions such as acting as serine protease for cleavage and binding of viral genomic RNA and binding to other capsid proteins during formation of the nucleocapsid and interacting with viral proteins during the formation and budding of virion particles [57, 70].

Natural inhibitors of arboviruses

Since, there are no effective antiviral therapeutic agents to treat infections caused by arboviruses, the only available treatments involve amelioration of symptoms of the disease by analgesics and non-steroidal anti-inflammatory agents [5, 22]. Arising from recent epidemic outbreaks, it is important to recognize any natural products that might potentially treat these virulent pathogens. This is particularly relevant since they are readily available inexpensive treatment options and may also be considered as templates for the development of efficacious derivative antivirals [15].

Plants are a rich reserve of natural chemical compounds exhibiting antiviral properties. Over the centuries, natural products from plants have been used to treat diseases and during the last few decades, these compounds have been studied for their antimicrobial and more specifically, antiviral activities. A number of positive findings have been reported on the antiviral activity of phytochemical compounds like saponins, flavonoids, alkaloids, sterols, and other compounds [23, 71, 72]. Flavonoids are the largest group of plant compounds. These secondary metabolites widely occur in edible plants such as fruits, vegetables, seeds, nuts, spices and plant stems. Over the years, various types of flavonoids have been studied for their broad spectrum of medicinal benefits such as antioxidant, anti-inflammatory, antitumor, antimicrobial and pro-apoptotic activities which are accompanied by minimal toxicity [73, 74]. Several recent studies have unveiled anti-flavivirus and anti-alphavirus properties of flavonoids and other plant compounds. Herein we describe some natural products that display inhibitory propensities against flaviviral and alphaviral infections.

In 2016, Carneiro *et al.*, reported that epigallocatechin gallate (EGCG), a natural compound found in a wide variety of foods, especially green tea, had inhibitory properties against the entry of the Zika virus into host cells *in vitro*. The study also revealed that the anti-Zika activity of EGCG was related to a direct interaction with the lipid envelope, resulting in destruction of viral particles [75]. Other studies indicated that EGCG prevented CHIKV transduction by blocking cell entry of CHIKV Env-pseudotyped lentiviral vectors and suppressed CHIKV attachment [76] whilst catechin-5-O-gallate was predicted by *in silico* analysis to interact with CHIKV proteins thereby showing antiviral promise [70].

Quercetin-3-O-glucoside (Q3G) is a glycosylated derivative of the plant flavonol quercetin. This natural polyphenolic compound has been evaluated for an ability to prevent the initiation of ZIKV infection in both tissue culture and knockout mice by blocking Zika virus entry into the host cell. [77]. Investigations on fisetin (a related plant flavonoid present in many vegetables and fruits), disclosed a dose-dependent inhibitory proclivity against CHIKV replication with minimal toxicity [78, 79] suggestive of conceivably wider antiviral activity. This was substantiated by Zandi *et al.*, when they tested this compound on dengue virus type-2 and ascertained that fisetin considerably decreased the copy number of DENV-2 RNA in infected cells [80].

In order to distinguish novel inhibitors of DENV, an ethyl acetate bark extract of *Cryptocarya chartacea* has been shown to have significant inhibitory properties against DENV NS5 polymerase. The outcome of this elegant

phytochemical study resulted in the isolation of a series of active 6-mono and 6,8-dialkylated flavanones, named chartaceones. In addition, from an extract of *Trigonostemon cherrieri*, a series of chlorinated daphnane diterpene orthoesters (DDO): trigocherrins (non macrocyclic) and trigocherriolides (macrocyclic) were also isolated. These natural compounds possessed a selective and potent inhibitory effect against chikungunya virus replication in cells [21].

Rosmarinic acid (RA) is an ester of caffeic acid, which is commonly found in several members of the *Lamiaceae* family [81]. This polyphenolic phytochemical diminishes the replication of Japanese encephalitis virus in an *in vivo* model [82]. What is more, virtual screening of a natural compound library predicted not only the inhibitory effects of rosmarinic acid on CHIKV capsid protein but also a suppressive activity of Arjungenin, (a triterpenoid isolated from *Terminalia arjuna*/T. chebula), on CHIKV E3 structural protein [70].

Baicalein (5, 6, 7-trihydroxyflavone) is an extractable flavonoid found in *Scutellaria baicalensis* and baicalin (5,6-dihydroxy-7-O-glucuronide flavone) is also a flavonoid which can be isolated from *S. baicalensis* root [83]. In 2016, Hassandarvish and colleagues conducted an *in silico* study to identify the potential of baicalein and baicalin as possible antiviral candidates in the treatment of DENV. It was deduced that both baicalein and baicalin are potential inhibitors of DENV by interaction with one of the vital dengue virus proteins, namely, E, NS3–NS2B and NS5 (baicalein interacting with DENV NS3/NS2B and baicalin having binding affinity for viral NS5) affecting the viral replication cycle. [84]. In accord with this, Zandi *et al.*, showed that baicalein exerts potent virucidal effects against DENV-2 replication in Vero cells [85]. Apart from DENV, an antiviral effect of baicalin on CHIKV has been investigated computationally by Seyedi *et al.* who conceptualized a compelling interaction with nsP3 [86]. Also, the *in vitro* virucidal effects of baicalein on Japanese encephalitis viral (JEV) replication have been evaluated though any underlying molecular or cellular mechanism(s) of action remain unknown [87]. Possible inhibitory effects of baicalein and baicalin have been investigated against ZIKV in cell lines. Thus, baicalein was highly repressive during intracellular ZIKV replication whereas baicalin was most potent against viral entry and from computer modelling, both plant products divulged convincing binding affinities towards ZIKV NS5 protein [88].

The flavanones naringenin and hesperetin, are natural commonly occurring plant secondary metabolites derived from *Citrus aurantium* reported to have both antioxidant and anti-inflammatory properties [89]. They also block Sindbis (SINV, neurovirulent strain) viral replication *in vitro* [90]. In a more recent study carried out by Ahmad *et al.*, both flavanones exhibited antiviral properties against CHIKV *in vitro* by preventing viral replication at the post-entry stage [91]. Aside from SINV and CHIKV, naringenin and hesperetin also display antiviral properties against yellow fever likewise by blocking viral replication and reducing the number and size of viral plaques [92]. In addition to this, naringenin can affect the entry of Semliki Forest Virus (SFV) into host cells [93].

Betulin, also known as betuline, betulinol, and betulinic alcohol, is a naturally occurring triterpene. It is commonly obtained from the bark of the white birch tree (*Betulaceae* family) but can also be isolated from punica, zizyphus, asparagus and ocimum [94]. A range of heterocyclic betulin derivatives were found to prevent SFV replication by Pohjala *et al.* The free or acetylated OH group at C-3 was recognized as a substantial structural contributor to anti-SFV activity, 3, 28-di-O-acetylbetulin being the most potent derivative in this respect [95]. Berberine is an isoquinoline alkaloid found in a variety of plants including, goldenseal, goldthread, Oregon grape, European barberry and tree turmeric [96]. This compound was evaluated as a potential anti-CHIKV agent affecting several of circulating strains of CHIKV. In this regard, berberine is capable of suppressing one or more host factors important for the replication of CHIKV including viral synthesis and protein expression. Furthermore, berberine also exhibits similar effectiveness against other alphaviruses such as SFV, SINV and O'nyongnyong viruses [97].

Antiviral properties of several polyphenols present in plants (delphinidin, cyanidin, catechin, epicatechin, epigallocatechin, and epigallocatechin gallate) have been evaluated against WNV, DENV, and ZIKV. In this context, delphinidin and EGCG act on WNV production mainly by affecting the attachment and entry stages of the viral life cycle. Moreover, both these polyphenols exhibited a direct effect on virus particles and they also impaired the infectivity of ZIKV and DENV [98].

In order to identify novel inhibitors of DENV-2, an *in silico* investigation was performed to evaluate a total of over 2000 plant-derived secondary metabolites in preventing DENV protease (NS2B-NS3pro), helicase (NS3 helicase), methyltransferase (MTase), and the viral envelope. The study analysis revealed that 24 compounds from the entire

collection appeared to exhibit avid docking properties with DENV NS2B-NS3 protease and 13 phytochemicals showed a pronounced capability for attachment to DENV MTase. Additionally, 21 of the compounds displayed a pronounced likelihood to attach to DENV NS3 helicase, 8 plant-derived components had a strong proclivity to dock with DENV RNA dependent RNA polymerase and 32 of the compounds possessed a significant capacity to dock with the DENV envelope protein [99].

Curcumin is a plant-derived polyphenolic compound (non-flavonoid polyphenol) which occurs in the rhizome of turmeric (*Curcuma longa*). It has been reported that this compound mediates its antiviral properties through a variety of mechanisms [100]. Hence, it reduces the production of infectious particles of Japanese encephalitis virus via a dysregulated ubiquitin-proteasome system and an accumulation of ubiquitinated proteins [101]. Curcumin also has an impact on *in vitro* ZIKV and CHIKV replication by suppressing viral cell surface binding [102]. Sophoraflavenone G (SFG) is another flavanone compound naturally obtained from *Sophora Flavecens*, *Sophora alopecuroids*, *Sophora pachycarpa*, and *Sophora exigua* [103]. The capacity of SFG to restrict the replication of two viruses in the flavivirus genus; DENV and ZIKV was tested. In a dose-dependent manner, the number of expressed cells significantly reduced. This finding also suggested that the polymerase of both flaviviruses was severely inhibited by SFG [104].

In 2016, Zamora and collaborators probed the antiviral potential of a combination of monoterpene alcohols (CMA) derived from *Melaleuca alternifolia* as a candidate to treat WNV infection. The *in vitro* results signified that CMA produced anti-WNV activity and significantly reduced viral titers and the number of infected cells. Detailed analysis of the antiviral mechanism of CMA indicated an affect at the early step of the viral life cycle between the entry and cleavage of the polyprotein, resulting in a reduction in NS1-expressing cells during WNV infection. The results suggested that CMA could be an effective candidate to treat pathogenic and non-pathogenic (in human) strains of WNV [105].

Palmatine is one of four main protoberberine alkaloids extracted from several plants including *Corydalis yanhusuo*, *Enantia chlorantha*, *Coptis Chinensis*, *Phellodendron amurense*, among others [106]. It has potential as an antiviral agent by virtue of its ability to inhibit WNV NS2B-NS3 protease [107]. Furthermore, palmatine can also suppress DENV-2 and YFV in a dose-dependent manner by blocking the function of NS3 protease [108, 109]. Several reports have signified that *Boesenbergia rotunda* (L.) compounds such as chalcones and flavanoids, have pharmaceutical properties. Chalcones are aromatic ketones that form the central core of a diverse array of biological compounds which are the biosynthetic precursors of flavonoids and isoflavonoids in plants [110]. These natural products possess antimicrobial, antifungal, anti-inflammatory, and antitumor properties. In this connection, the cyclohexenyl chalcone derivatives, 4-hydroxypanduratin A and panduratin present in *B. rotunda* have been reported to competitively inhibit DEN-2 virus NS3 protease [111]. Studies have also revealed that two flavonoids, namely kaempferol and daidzin, yield antiviral effects against JEV. Kaempferol is a natural flavonol which has been isolated from a variety of plants and plant-derived foods such as tea, broccoli, delphinium, witch-hazel, grapefruit, Brussel sprouts, apples and medicinal plants while daidzin is an isoflavone found mainly in soybeans. The antiviral action of both these natural products involve prevention of JEV replication and E viral protein expression and in this respect, kaempferol is more effective than daidzin, though their overall action is augmented by combination [112, 113].

An *Isatis indigotica* extract and its constituent alkaloidal compounds (indigo and indirubin) is potently virucidal against JEV and this antiviral action originates from a blockade of viral attachment, suppressed JEV replication and a reduction in viral yield [114, 115]. Also, the phenylpropanoid dibenzylbutyrolactone arctigenin which is found in some species of the *Asteraceae* family, including *Bardanae fructus*, *Saussurea medusa*, *Torreya nucifera*, *Saussurea heteromalla*, *Ipomea cairica*, *Forsythia intermedia* and *Arctium lappa*. This biologically active lignan has antioxidant, anti-inflammatory, and antiviral activities and in both *in vivo* and *in vitro* studies, it has been established that arctigenin reduces the severity of JEV-induced infection by diminishing replication and the viral titer [116, 117]. Finally, the flavonol quercetagenin, which has been isolated from *Citrus unshiu*, is a hexahydroxyflavone that has an additional 6-OH group on the molecular structure of the flavone backbone. This phytochemical compound is considered to have significant potential as an antiviral agent against extracellular CHIKV particles by virtue of its docking capacity at the active site of nsP3 [86, 118].

Table 2. Natural compounds that show inhibitory activity against arboviruses

Compound	Class	Source	Virus	Mechanism of Action	References
Epigallocatechin gallate (EGCG)	flavonoid	green tea and black tea Some other foods such as strawberries, plums, hazel nuts, raspberries, blackberries, plums, peaches, kiwi, and avocado	Zika virus (ZIKV)	lipid envelope	[75, 76]
			Chikungunya virus (CHIK),	capsid protein and CHIKV attachment	
			West Nile virus (WNV)	attachment and entry stages of the virus	
			Dengue virus (DENV)	E protein	
Quercetin-3-O-glucoside (Q3G)	polyphenolic compound	<i>Lepisorus contortus</i>	ZIKV	virus entry	77
Fisetin	Flavonoid	many vegetables and fruits such as strawberry, persimmon, apple, and many others	CHIKV	replication of CHIKV and DENV	[78, 79, 80]
			DENV type-2		
Chartaceone	Flavonoid	<i>Cryptocarya chartacea Kostermans</i>	DENV	NS5 polymerase	[21]
Trigocherrins (non macrocyclic) and trigocherriolides (macrocyclic)	diterpene	<i>Trigonostemon cherrieri Veillon</i>	CHIKV	viral replication	
Rosmarinic acid	phenolic compound	several members of the Lamiaceae family	CHIKV	capsid protein	[70, 81, 82]
			Japanese encephalitis virus (JEV)	viral replication	
Arjungenin	triterpenoid	<i>Terminalia arjuna/ T. chebula,</i>	CHIKV	E3 protein.	
Baicalein	flavone	<i>Scutellaria baicalensis,</i>	DENV	NS3/NS2B protein	[83-88]
			JEV	viral replication	
			ZIKV	high inhibitory activity during intracellular ZIKV replication AND	

				the strongest binding affinities towards Zika virus NS5 protein	
Baicalin	flavone	<i>S.baicalensis</i>	CHIKV	nsP3	
			DENV	NS5 protein	
			ZIKV	. most potent against virus entry. And the strongest binding affinities towards Zika virus NS5 protein.	
Betulin	triterpene	white birch tree, <i>Punica</i> , <i>Zizyphus</i> , <i>Asparagus</i> , <i>Ocimum</i>	Semliki Forest virus (SFV)	virus replication	[94, 95]
Naringenin and hesperetin	flavanone	<i>Citrus aurantium</i>	Sindbis virus (SINV)	Viral replication	[89-93]
			Yellow fever virus (YFV)	viral replication.	
			CHIKV	viral replication in post-entry stage	
			SFV	entry of Virus into the host cell	
Berberine	isoquinoline alkaloid	goldenseal, goldthread, Oregon grape, European barberry, and tree turmeric.	CHIKV	reduce the virus synthesis and protein expression	[96, 97]
			SFV		
			SINV		
			O'nyongnyong viruses (ONNV)		
delphinidin	flavonoid	<i>Leschenaultia</i> cv. Violet Lena	DENV	E protein	[98, 99]
			WNV	attachment and entry stages of the virus	
			ZIKV	E protein	
Curcumin	polyphenolic compound	turmeric (<i>Curcuma longa</i>) rhizome	JEV	viral replication	[100-102]
			ZIKV	suppressing virus binding to the cell	

			CHIK	surface	
Sophoraflavenone G	flavanon	<i>Sophora Flavecens, Sophora alopecuroids, Sophora pachycarpa, and Sophora exigua.</i>	DENV	polymerase of both flavivirus was severely inhibited by SFG	[103, 104]
			ZIKV		
monoterpene alcohols (CMA)	monoterpene	<i>Melaleuca alternifolia</i>	WNV	affect the early step of viral life cycle between the entry and cleavage of the polyprotein, resulting in the reduction in NS1-expressing cells	[105]
Palmatine	alkaloid	<i>Corydalis yanhusuo, Enantia chlorantha, Coptis Chinensis, Phellodendron amurense, and other plants</i>	YFV	NS3 protease	[106-109]
			DENV	NS3 protease	
			WNV	NS2B-NS3 protease	
Cyclohexenyl chalcone derivatives 4-hydroxy panduratin	chalcone	<i>Boesenbergia rotunda (L.)</i>	DENV	NS3 protease	[110, 111]
kaempferol	flavonol	variety of plants and plant-derived foods such as tea, broccoli, delphinium, witch-hazel, grapefruit, Brussel sprouts, apples, and medicinal plants	JEV	viral replication and E viral protein expression	[112, 113]
daidzin	isoflavone	soybeans			
indigo and indirubin	indole alkaloid	<i>Isatis indigotica</i>	JEV	blocking virus attachment, reducing virus yield in cell culture and inhibition JEV replication	[114, 115]
Arctigenin	lignan	species of the Asteraceae family, including the <i>Bardanae fructus, Saussurea medusa, Torreya nucifera, Saussurea heteromalla, Ipomea cairica</i> and <i>Forsythia intermedia, Arctium lappa</i>	JEV	reduction of viral titer and replication	[116, 117]
Quercetagetin	flavonol	genus <i>Eriocaulon</i>	CHIKV	nsP3.	[86, 118]
silymarin	flavonoid	<i>Silybum marianum</i>	CHIKV	suppressing post-entry stages of viral replication, most likely CHIKV RNA	[119]

				replication significantly with a dose dependent manner and also expression of viral structural E2 protein were down-regulated	
Luteolin	flavone	camomile tea, perilla leaf, green pepper, celery and <i>Cynodon dactylon</i>	JEV	inhibits JEV infection at the virus post-entry stag and E protein was significantly reduced	[120]
Apigenin	flavone	<i>Melissa officinalis</i> , <i>Cynodon dactylon</i> and <i>Aspalathus linearis</i>	CHIKV	inhibited CHIKV replication	[121]
Trigowiin A	diterpenoid	<i>Trigonostemon howii</i>	CHIKV	inhibited CHIKV replication	[122]
labranine and 7-O-methyl-glabranine	flavonoid	<i>Tephrosia sp</i>	CHIKV	inhibited a viral replication of dengue virus	[123]
Harringtonine	alkaloid	<i>Sephalotaxus harringtonia</i>	SINV	Viral protein syntesis	[124]
			CHIKV		
3- α -tigloyl-melianol and melianone	triterpene	<i>Melia azedarach L</i>	YFV	Virus entry process	[125, 126]
			WNV		
			DENV		
Saponin	Saponin	<i>Achyrocline satureioides</i>	Western equine encephalitis virus (WEEV)	Viral replication	[127]
affeic acid	polyphenol				
chlorogenic acid					
quercetin	flavonoid				
pinostrobin	flavonoid	<i>Boesenbergia rotunda (L)</i>	DENV	NS2B-NS3 protease	[128]
pinocembrin					
alpinetin					

phenylpropanoid cardamonin					
cyclohexenylchalcone -					
hydroxypanduratin A					

Conclusion

Arboviral infections are a serious global public health problem and they are a progressively common cause of severe febrile disease that can progress to long-term physical or cognitive impairment and even lead to early death. Since, there are no antiviral treatments against these infections, there is a pressing need to identify new inhibitors of these viruses with structural diversity. Attention has been drawn to natural compounds that may conceivably be used to contribute towards the development of effective antiviral treatments. To date, only a limited number of natural products capable of inhibiting arboviral replication have been identified while the bulk of earlier studies have focused on nucleoside analogs often with non-optimal selectivity. Consequently, biologically active phytochemicals warrant further attention in this field. The antiviral mechanism(s) of plant compounds are largely based on their inhibitory effects on viral RNA synthesis and viral entry. In the context of alphavirus inhibitors, there have been several reports on non-structural proteins having substantial roles in the stages of the viral life cycle in the quest for novel drugs. Additionally, identification of inhibitors of structural and non-structural flavivirus proteins, replication and entry steps are notable starting points as targets for the design of novel anti-flavivirus agents because the damage caused by virulence factors during replication can be minimized. Finally, this review indicates that identification of plant-derived compounds could play a crucial role in the discovery and development of new antiviral treatments.

Conflict of interests

The authors declared no competing interests.

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