

# Dengue Virus Cellular Receptors and Tropism

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**Abstract** Viral entry into host cells primordially defines tropism and represents an attractive target to counteract infection either by antiviral agents or by immune mediated mechanisms. Research on Dengue virus entry presents interesting challenges. Whatever the mechanism dengue virus exploits to gain entry into cells, this had to be evolutionarily conserved, so that it is now present in arthropod and human cells. Until now, dengue cellular receptors were not completely unraveled. However, we have clues about the key steps dengue virus is relying on. Initially a group of factors that interact with the virus through carbohydrate interaction assure its adherence and further contact with a protein receptor complex, which is held together thanks to its special interaction with cell membrane lipidic platforms. This interaction may be so intimate that it may trigger not only viral entry through receptor-mediated endocytosis, but also activation of cell signaling pathways that the virus is going to subvert to its advantage.

**Keywords** Dengue · Receptor · Tropism · Viral entry · Viral tropical medicine · Virus · Dengue cellular receptor

## Introduction

### Importance of Dengue Virus Infection

The increase in dengue's worldwide prevalence has made this disease the most important mosquito-borne viral infection for the past six decades. Dengue virus (DENV) inflicts a significant health burden in endemic countries, among the most populated areas worldwide. According to the World Health Organization (WHO), Southeast Asia and southwest Pacific regions reported nearly 75 % of the total cases of dengue infections (WHO, 2009). With an annual average of 574,000 cases reported, the economic cost to eight countries in America and Asia is at least \$587 million [1]. Indeed, WHO classifies dengue as a major international public health problem [2].

DENV has one of the simplest infectious cycles among arthropod-borne viruses. A person gets the virus from an infected mosquito of the *Aedes* genus. After a replication period of seven to ten days, DENV reaches levels in peripheral blood (viremia) that allow transmission to an uninfected mosquito. After infecting its digestive tract and circulatory system, DENV reaches the vector salivary glands and the cycle repeats. Dengue prevalence has increased not only because geographic areas where *Aedes* mosquito dwells coincide with unplanned human settlements, but also because of mutations in circulating DENV and climatic factors [3]. It follows then, that the analysis of DENV entry into host cells of both, human and mosquito nature is of paramount importance for the design of control strategies.

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## Dengue Virus Tropism

**Human cells** In vitro DENV is able to infect a large variety of cell lines from murine, hamster, canine, primate, and naturally mosquito and human hosts lineages. However, in vivo, few human cell types have been shown to support DENV replication. Samples from patients traditionally come from complicated, severe cases and not from initial phases of infection. In consequence, histopathological studies from patients' biopsies documenting the localization of DENV proteins might be confused with cells that can actively replicate the virus with those cells that acquire dengue antigens by endocytic or phagocytic uptake of viral particles or proteins [4, 5]. The initial steps following an infected mosquito inoculation have been studied ex vivo, where a skin explant is infected transcutaneously [6]. DENV is able to infect resident cutaneous Langerhans dendritic cells (DCs) at the site of mosquito inoculation. The infection of these cells results in two important events. First, their activation and migration through the lymphatic system brings DENV to sites where a robust viral replication occurs (secondary lymphoid tissue) and, secondly, in the expression of cytokines and chemokine-mediated recruitment of immune competent cells [6]. The viral infection primary targets in humans and murine models are monocytes/macrophages, as well as B and T lymphocytes [7–9, 10••]. Although hematopoietic lineage cells are major sites of DENV replication, some other cells of non-hematopoietic origin such as hepatocytes and Kupffer cells or neurons and microglia can be permissive to DENV during natural infection [5, 11]. Furthermore, autopsies have documented infection of endothelial cells (ECs), a finding that has been reproduced in a mouse model of DENV infection [4, 12]. Infection of ECs can contribute to the pathogenesis of the disease by increasing viremia, cytokine secretion, and modulating host's immune responses [13•]. In this respect, it is not clear if the tropism observed in fatal cases of DENV infection or in the mice model is the same that the one detected in patients with dengue fever [4, 14, 15].

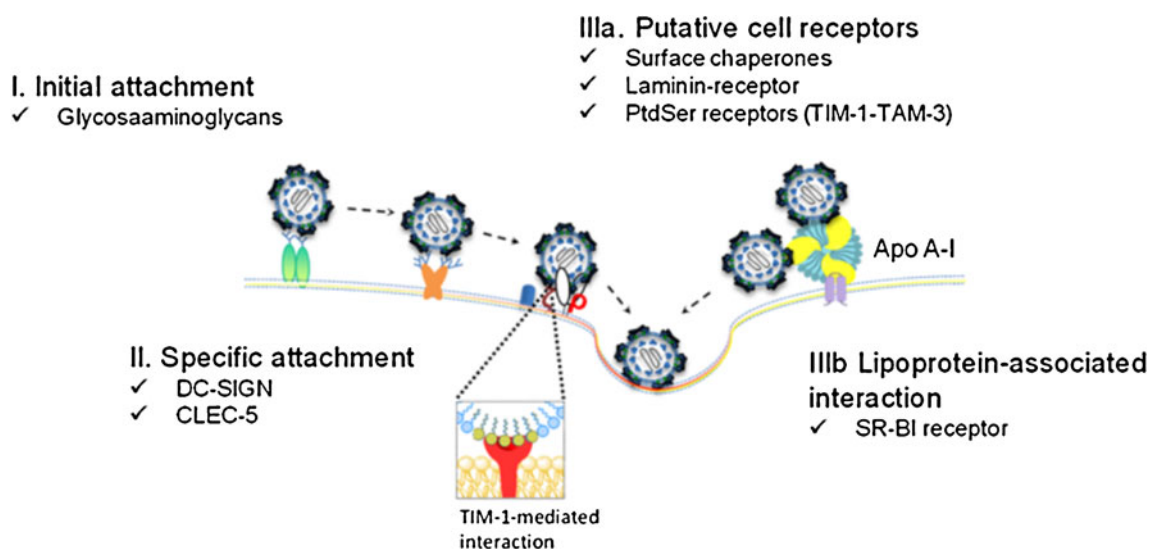
**Mosquito cells** Two arthropod vectors from the *Aedes* genus are responsible of DENV transmission to humans, *Ae. aegypti* and *Ae. Albopictus* [16]. The haematophagous female necessitates a blood meal for oviposition, therefore, *Aedes* mosquitos usually breed in close proximity to humans. This fact has enormous implications for DENV epidemiology and prevention strategies. The first mosquito organ where DENV replicates is the midgut. Then the virus spreads using the circulatory system to other organs/tissues such as the tracheal system, fat body, salivary glands, nervous system, esophagus, hemocytes, ommatidia of compound eyes, and malphigian tubules [17]. This period of replication that is to reach the salivary glands is known as intrinsic incubation period and lasts approximately five to

seven days, afterwards the mosquito becomes a life-long potential virus transmitter.

## Dengue Virus Receptors in Human Cells

**Dengue virus entry into human cells** The first step in any viral life cycle is viral entry. This dynamic process is usually carried on in multiple steps and the nature of each step depends on the particular characteristic of the virus (depicted in Fig. 1). DENV is an enveloped virus, with a major glycoprotein (E), arranged in ninety dimers with a quasi-icosahedral symmetry. The structure of DENV is very conserved in the *Flaviridae* family, which has prompted several groups to postulate that the entry process is conserved, but not the particular players. The viral membrane derives from the previous infected cell plasmatic membrane, and it is possible that the lipid components budded off with the virus to play a role to facilitate the process of viral entry [18•], which in the case of dengue and other flaviviruses, is rather fast and estimated to occur in minutes [19]. The study of viral receptors traditionally starts with a susceptible cell line and through biochemical or genetic methods a surface molecule(s) is identified and postulated as putative receptors. Next, if the experimental induction of expression of the putative molecule in non-permissive cells renders them susceptible to infection, the putative molecule is considered a *bona fide* cell receptor. This process has found some hurdles for DENV and other arthropod borne flaviviruses. The range of susceptibility of many cell lines of different origins is very wide, and finding a non-susceptible cell line has proven to be a non-trivial task. On the other hand, the molecules identified to be putative receptors are either of not a protein nature (hence, no induction of expression) or with an important intracellular housekeeping function besides their potential role on the cell membrane (for instance surface chaperones). In sum, no single protein or proteins have been singled out as the DENV cell receptor, and what follows is a recounting of the molecules that have been implicated in DENV entry into mammalian cells.

**Glycosaminoglycans (GAGs) and other attachment factors** The first molecule that was identified to participate in DENV entry into mammalian cells was heparan sulfate [20–22]. Highly sulfated GAGs are ubiquitous molecules present on the surface of several types of cells, also mediating attachment for many viruses [23]. It is documented that there are electrostatic attractions among a dengue virus E glycoprotein and the negatively charged carbohydrate moieties present in GAGs [24]. Due to the documented evidence that the GAGs-DENV interactions are stronger when the virus has been passaged in cell-culture repetitively, and due to its correlation with in vivo attenuation, it has been suggested its role is artifactual. The most relevant attachment factor for DENV entry identified so far is the calcium-dependent lectin,



**Fig. 1** The process in multiple steps and the nature of each step depends on the particular characteristic of the virus

dendritic cell-specific intercellular adhesion molecule 3-grabbing non-integrin (DC-SIGN) [25–27]. This receptor has high affinity for high-mannose ligands [28]. During the mosquito blood-meal, DENV is deposited in the dermis where its interaction with resident dendritic cells (Langerhans cells) has been documented [6]. Upon entry, dendritic cells migrate to draining lymph nodes where the infection is spread to immune-competent myelo-monocytic cells. Whether DC-SIGN is the only factor responsible for dengue virus entry or the virus replicates actively in dendritic cells is still unknown. Another lectin that may play a role in dengue virus attachment and release of pro-inflammatory mediators important in the pathogenesis of severe forms of dengue [31, 32]. In sum, it is clear that the first step in dengue virus entry involves attachment of the virion to the cell surface.

**Cell surface chaperones** Among DENV entry factors identified thus far, there is a group that has been described by independent groups, the surface chaperones. Experimentally, these molecules have been found through binding assays using the envelope glycoprotein as a ligand. HSP-90, HSP-70 and GRP-78 are thought to be part of a receptor complex that mediates dengue entry into human cells from monocytic, neural and hepatic origin [33, 34]. It is not known whether their extracellular function is similar to their well-known intracellular, ATP-dependent protein folding activity, or how they associate with the cell membrane. It has been long been reported that lipopolysaccharide (LPS) inhibits DENV entry into monocytes [35]. This effect is dose dependent. On this regard, surface HSPs were also documented as part of an interacting cluster of plasmatic proteins mediating physiological effects of LPS sensing on cells [36]. It is tempting to

propose that heat shock as an effect on DENV entry through the increase of heat-shock-responsive molecules on the surface of infected cells. This effect was documented in the infection of a monocytic cell line [37]. Another protein with chaperone function reported as a DENV receptor is the protein disulfide isomerase (PDI). This has been implicated in DENV entry into endothelial cells [38•]. Besides its important role in the endoplasmic reticulum, it has been detected on the cell surface of lymphocytes [39], platelets [40, 41], and endothelial cells [38•]. PDI has been implicated in entry of several viruses such as polyomavirus [42], New Castle disease virus [43], and the human immunodeficiency virus [39, 44, 45].

**Lipid receptors** The only gain-of-function screening in the search for DENV receptors found that phosphatidylserine (PtdSer) receptors TIM-1 (T cell immunoglobulin domain and mucin domain) directly and indirectly through TAM ligands, are relevant on DENV entry into the human 293 T cell line [46•]. These receptors mediate binding of apoptotic cells to be scavenged and immunotolerance. Therefore, an intimate interaction of cell receptors with lipids present on the virion membrane envelope suggests the structure of entering particles needs to be revisited. In any case, the lipid content present on the target cell has long been demonstrated as important to optimize DENV entry.

#### Lipids in DENV Entry

**Role of lipid rafts in dengue virus entry** Recent evidence suggests that lipid rafts, defined as detergent resistant membrane microdomains enriched in cholesterol and sphingolipids [47], might serve as platforms to concentrate virus receptors. Since several of the DENV cell receptors are residents of lipid rafts or are relocated to them after DENV binding [48–50],

these platforms traffic the virus to a proper intracellular environment to favor conformational changes in the envelope proteins during the fusion process [51, 52].

The role of lipid rafts in DENV entry has been studied in several mammalian cell lines. In macrophages it has been demonstrated that lipid rafts play an important role in DENV entry in the absence and also in the presence of facilitating antibodies [33, 53••]. The cholesterol dependence of DENV entry and post entry steps was also observed in the mouse neuroblastoma cell line N18, in the hepatocarcinoma cell line Huh 7 [54, 55] and in the endothelial cell line HMEC-1 [56].

However, the involvement of cholesterol in DENV entry does not seem to be a general event, because it has been described that, for green monkey epithelial kidney cell lines (Vero and BS-C-1 cells), hepatocarcinoma cell line HepG2 and ECV304 cells, cholesterol is not required [18•, 19, 57–59]. Furthermore, contradictory data exist about the cholesterol dependence in U937, K562 (hematopoietic) and A549 cells (lung carcinoma) [18•, 33, 57, 60]. On the other hand, DENV entry in the mosquito cell line C6/36 seems to be cholesterol and lipid rafts independent because pretreatment of these cells with lipid rafts disrupters (M $\beta$ CD, filipin and Nystatin) did not inhibit DENV 1 and 2 infection [61, 62]. Similarly, it has been observed that the infection and fusion capacities of DENV remain unaltered in a cholesterol depleted-insect cell system [63].

*Role of lipids in viral fusion* Several observations have suggested that, in addition to low pH, lipid cofactors are required for DENV fusion. The outer leaflets of the plasma membranes of mosquito cells and the membranes of late endosomes of mammalian cells have high concentrations of the anionic lipids phosphatidylserine (PS) and bis(monoacylglycero)phosphate (BMP), respectively. It has been demonstrated that the DENV fusion occurs successfully only upon the addition of anionic lipids (PS and BMP) to liposomal, plasma and intracellular membranes, suggesting that when DENV get into contact with the late endosomes membranes containing anionic lipids, the envelope protein (E) acquires its fusogenic conformation by action of endosomes acidification [64••].

On the other hand, the addition of cholesterol to artificial receptor-free lipid membranes (liposomes) consisting of phosphatidylcholine and phosphatidylethanolamine, increases the membrane fusion capacity of DENV, Tick Borne Encephalitis Virus (TBEV) and West Nile Virus (WNV) at low pH [65••]. Furthermore, the role of cholesterol in viral trafficking was evidenced using U18666A, a cholesterol transport inhibitor, which causes a delay in the release of viral genome into the cytoplasm [66•].

Finally, lipids present on the DENV envelope, specifically cholesterol and PS, have also been involved in the fusion process. Carro et al., demonstrated that the cholesterol present in the viral particle is a critical factor during fusion process because DENV binding and entry is inhibited when viral particles were treated with M $\beta$ CD [18•]. Similarly, as mentioned by Meertens et al., PS present on DENV particles surface is recognized directly by TIM proteins and indirectly by TAM proteins that mediate the PS-dependent phagocytic removal of apoptotic cells and can be exploited by the virus as entry factors [46•].

*Role of lipoproteins in viral infection* Recently, it has been reported that DENV C protein is able to interact with Very Low Density Lipoprotein (VLDL), specifically with Apo E, which is the main surface apolipoprotein in VLDL. This fact is supported by the strong structural similarities between Apo E and perilipin 3, the DENV C protein ligand in lipid droplets [67•, 68]. It has been proposed that the interaction between C and VLDL ApoE may allow the formation of Lipovirparticles (LVP), as has been reported for HCV [69]. They might be transported to the extracellular medium and infect neighboring cells through lipoproteins receptors. However, the presence of LVP during DENV infection in vivo has not been observed, and the direct role of VLDL or LDL receptors in DENV entry has not been determined. On the contrary, there is a report where the interaction of the apolipoprotein A-I (Apo A-1), the main protein component in High Density Lipoprotein (HDL) with DENV particle was confirmed by co-immunoprecipitation assays [70•]. The authors suggest that this interaction is able to promote DENV infection through Apo A-I receptor, the Scavenger Receptor class B type I (SR-BI), and provides insights into the functional importance of lipoproteins in dengue pathogenesis.

#### Dengue Virus Receptors in Mosquitoes

The fact that DENV enters mosquito cells as efficiently as mammalian cells implies that the molecules, strategies, or mechanisms used by the virus are either very conserved or fungible. Nevertheless, DENV is very selective regarding the arthropod it uses, only mosquitoes from the *Aedes* genus are susceptible. This is by no means a rule for other arthropod-borne viruses from the Flaviviridae family, like the West Nile virus (WNV), or the Japanese encephalitis virus (JEV), which are more “promiscuous” in this regard. Next, a brief description of some of the molecules identified as putative DENV receptors in mosquito cells is offered (see Table 1).

*Heparan sulphate* Treatment of C6/36 cells (a line derived from *Ae. Albopictus* larva, [71] with heparinase III prior to infection with the four DENV serotypes showed a variable inhibition, from 25 % for DENV 2 to 60 % for DENV 1 [72].

**Table 1** Summary of the putative receptors for DENV identified in mosquitoes or mosquito cells

Receptor	DENV Serotype	Model	Reference
Hepara sulfate	1,2,3, and 4	C6/36 cells	[72]
high-affinity laminin related receptor	2, 3, and 4	C6/36 cells	[72]
Prohibitin	2	CCL-125 and C6/36 cells <i>Ae aegypti</i> mosquitoes	[76•]
HSP70	4	C6/36 cells	[83•]
Hsc70	2	A7 and C6/36 cells <i>Ae aegypti</i> mosquitoes	[75]
GRP78 (BiP)	4	C6/36 cells	[83•]
PDI	4	C6/36 cells	[83•]
Vav-1	2	A7 and C6/36 cells <i>Ae aegypti</i> mosquitoes	[75]
ATP synthase $\beta$ subunit	2	A7 and C6/36 cells <i>Ae aegypti</i> mosquitoes	[75]
Orisis	2	A7 and C6/36 cells <i>Ae. aegypti</i> mosquitoes	[75]
67 and 80 kDa proteins	1, 2, 3, and 4	C6/36 cells <i>Ae. Aegypti</i> mosquitoes	[94] [93] [95]
77, 58, 54, and 37 kDa proteins	1, 2, 3 and 4	<i>Ae. Aegypti</i> mosquitoes	[96]
67, 56, 54, 50, and 48 kDa proteins	1, 2, 3 and 4	<i>Ae. Polynesiensis</i> mosquitoes	[96]

Interestingly, a similar treatment reduced 40 % the infectivity of JEV [73] whereas treatment with glycosidases, heparinases, lectins and phospholipases did not affect the infection by WNV [74] suggesting they use different molecules for cell surface attachment in mosquito cells.

**Laminin receptor** Homology with the 37/67-kDa human high-affinity laminin-like receptor was documented in a 50-kDa protein present in C6/36 cells that binds DENV 2, 3 and 4. To support its role as receptor, immune sera raised against this molecule or soluble laminin was able to inhibit the infectivity of the four serotypes of DENV with differing efficiency [72]. In contrast, soluble laminin only reduced JEV infection up to 20-30 % in C6/36 cells suggesting that this receptor plays a minor role for JEV internalization [73].

**Prohibitin** Recently, two independent research groups identified the protein prohibitin (PHB) as a receptor for DENV 2 in cells lines from *A. albopitcus* and *A. aegyptii* (C6/36, CCL-125, and A7, respectively) as well as in midgut mosquito cells [75, 76•]. Silencing of prohibitin with specific siRNA reduced virus infection in both mosquito cell lines, and specific immune sera competed with both virus infection and cell binding. PHB is a ubiquitously expressed and highly conserved protein complex in eukaryotic cells composed of two proteins: PHB1 (~30 kDa) and PHB2 (~37 kDa) [77]. They have more than 50 % amino-acid identity and are prone to form oligomers that have been proposed to enhance the stability of the complex [77, 78]. PHB is present in many compartments of the cell, mainly

the mitochondria [79] but it is also present in the cytoplasm and nucleus [78]. Early evidence indicated that this protein acts as an inhibitor of cell proliferation, and that it plays a role in cell signaling at the plasma membrane and in transcriptional regulation [80]. Interestingly, it has also been described as a cell receptor for the Chikungunya virus (CHIKV), an arthropod borne alphavirus, in microglial cells [81•].

**Chaperones and heat shock-related proteins** Similarly to mammalian cells, C6/36 cells increase their susceptibility to DENV upon heat shock (HS). A 74-kDa protein that binds DENV 2 and is strongly expressed under HS conditions [82] was recently isolated and identified as three proteins: 78-kDa glucose-regulated protein (GRP78 or BiP), the 70-kDa heat shock protein (HSP70) and the 70-kD heat shock cognate protein (HSC70) [83•]. Specific antibodies against GRP78, and HSP70/HSC70 were able to inhibit DENV4 infectivity. These proteins are ER resident chaperones of the HSP70 family [84, 85]. They are located ubiquitously in cytoplasm, mitochondria, nucleus and plasma membrane [86–88]. Independently, another group also found HSC70 interacting with DENV2 in C6/36 and A7 cells as well as in *Ae aegypti* females [75]. Furthermore, the chaperone HSC70 has been implicated in JEV entry into C6/36 [89] and mammalian cells [90]. Additionally, a protein (40-kDa) with homology to protein disulfide isomerase (PDI) was also identified among the C6/36 molecules that interacted with DENV4 by affinity chromatography [83•]. Interestingly, PDI has been implicated in DENV entry to endothelial cells [38•] and is also a putative

receptor for other viruses (see above). Several groups through different approaches have documented the participation of cell-surface chaperone proteins in dengue virus entry into mosquito cells.

**Unidentified proteins** Several other proteins in mosquito cell lines or midgut cells from competent vectors have been proposed to function as receptors for DENV. Among them, a 45-kDa glycoprotein present on the surface of C6/36 was recognized by DENV4 by virus overlay protein blot assay (VOPBA) [91]. Its presence was also documented in extracts of different development stages of *Ae. aegypti* mosquitoes as well as in different susceptible mosquito tissues (salivary gland, ovaries and midgut), but not in a non-dengue vector as *Anopheles albimanus*, suggesting the participation of this molecule in DENV infection as well as in vertical transmission [92]. Other groups reported molecules with weights of 67 and 80 kDa important for DENV2 entry [93, 94]. Interestingly, the presence of the 67 kDa protein in midgut cells of three varieties of *Ae. aegypti* mosquitoes correlated with their susceptibility to DENV infection, thus, it has been proposed as a vector competence marker [95]. Finally, proteins ranging in molecular weight from 77 to 37 kDa obtained from *Ae. aegypti* and *Ae. polynesiensis* salivary glands extracts were identified to interact with the four DENV serotypes. Interestingly, proteins with similar molecular weights were recognized by DENV 4 clinical isolate, suggesting that no modification in receptor usage occurred in viruses propagated for a long time in vitro [96].

## Conclusions

Arthropod-borne viruses constitute an excellent model to study evolutionarily conserved mechanisms exploited for a virus to gain cell entry. Even though specific steps in DENV entry have been progressively well characterized in recent years, the specific role of host cell proteins in viral attachment and entry is not completely understood. However, it is possible to infer that a group of factors that interact with the virus through carbohydrate interaction assure its adherence and further contact with a protein receptor complex, which is held together thanks to its special interaction on lipidic platforms on the cell membrane. All these steps allow viral particle entry into early endosomes very rapidly. Chaperones constitute one of the host cell proteins used for DENV in both mammalian and insect cells. The use of several molecules during viral attachment and entry reveal the outstanding requirements and the complex mechanisms that DENV employs to enter the cell. Further studies directed to

understand the initial steps in DENV entry have the potential to provide insights into possible therapeutic targets to control DENV infection.

## Compliance with Ethics Guidelines

**Conflict of Interest** Jorge Reyes-del Valle, Juan Salas-Benito, Rubén Soto-Acosta, and Rosa M. del Angel declare that they have no conflict of interest

**Human and Animal Rights and Informed Consent** This article does not contain any studies with human or animal subjects performed by any of the authors.

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- Of importance
- Of major importance

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