Yellow fever virus envelope protein expressed in insect cells is capable of syncytium formation in lepidopteran cells and could be used for immunodetection of YFV in human sera

Maria CES Barros¹, Tatiane GCM Galasso¹, Antônio JM Chaib², Nicolas Degallier³, Tatsuya Nagata¹ and Bergmann M Ribeiro¹*

Abstract

Background: Yellow fever is an haemorrhagic disease caused by a virus that belongs to the genus Flavivirus (Flaviviridae family) and is transmitted by mosquitoes. Among the viral proteins, the envelope protein (E) is the most studied one, due to its high antigenic potential. Baculovirus are one of the most popular and efficient eukaryotic expression systems. In this study a recombinant baculovirus (vSynYFE) containing the envelope gene (env) of the 17D vaccine strain of yellow fever virus was constructed and the recombinant protein antigenicity was tested.

Results: Insect cells infected with vSynYFE showed syncytium formation, which is a cytopathic effect characteristic of flavivirus infection and expressed a polypeptide of around 54 kDa, which corresponds to the expected size of the recombinant E protein. Furthermore, the recombinant E protein expression was also confirmed by fluorescence microscopy of vSynYFE-infected insect cells. Total vSynYFE-infected insect extracts used as antigens detected the presence of antibodies for yellow fever virus in human sera derived from yellow fever-infected patients in an immunoassay and did not cross react with sera from dengue virus-infected patients.

Conclusions: The E protein expressed by the recombinant baculovirus in insect cells is antigenically similar to the wild protein and it may be useful for different medical applications, from improved diagnosis of the disease to source of antigens for the development of a subunit vaccine.

Background

Yellow fever (YF) is an haemorrhagic disease caused by a virus and transmitted by mosquitoes through two distinct cycles: the urban YF, transmitted by Aedes aegypti and the sylvatic YF, maintained in an enzootic cycle by Haemagogus and Sabethes mosquitoes with monkeys as main hosts [1]. No cases of urban YF have been reported in Brazil since 1942 [2]. The sylvatic YF is mostly restricted to wild and rural areas but recent outbreaks amongst human visitors and travelers together with the reinfestation of urban areas with the vector mosquito Aedes aegypti have concerned health authorities about the reurbanization of YF.

The main mechanisms for YF control consists of vaccination and insect vector control in urban areas. Yet, in 2008, 228 YF epizootic cases were reported and 64 cases of dead monkeys occurred just in January. These are more than the 104 YF epizootic cases and the 17 cases of dead monkeys during the whole year of 2007. Until July of the same year, 45 cases of YF amongst humans were confirmed with 25 deaths, which represents 55.6% case fatality rate [3]. Similar to what happens to other flavivirus diseases, the investment on improved diagnosis techniques for YF with faster and precise methods is crucial for the early detection and correct identification of yellow fever infection for prevention and control of
disease spread by public health authorities besides correct epidemiological reports [4].

Although clinical diagnosis is sufficient during an epidemic, laboratory diagnosis is the definite method to confirm yellow fever infection mainly in sporadic cases because yellow fever main symptoms may be confused with a broad range of related diseases that vary from severe malaria to dengue or leptospirosis. Viral isolation in mosquito cells cultures is a sensitive technique for the first days of infection, during the viremic period of the disease, but currently yellow fever diagnosis is based on serology, mainly enzyme-linked immunosorbent assays (ELISA) [5]. Yet, serological tests still use the whole virus as the antigen pool which could be improved with specificity by the use of antigenic parts of the virus, minimizing the risk of cross reactions with other flavivirus and risk of infection by health professionals.

Yellow fever virus (YFV) is an enveloped virus with a positive sense, single stranded RNA genome of 10,862 bases coding for a single ORF of 10,233 bp. This ORF encodes three structural proteins (Capsid, pM, and E) and seven non-structural proteins (NS1, NS2a, NS2b, NS3, NS4a, NS4b, NS5) [6]. Among the viral proteins, the E protein is the most studied one, due to its high antigenic potential.

E protein is involved in many events, such as receptor binding site for viral attachment [7], fusion, penetration, hemagglutination, host range and cell tropism [8]. It also has an important role in immunological anti-virus response, eliciting neutralizing antibodies and inducing protective response [9].

Native E protein presents itself as homodimers and its activity is intimately linked to its structure that suffers conformation rearrangements changing the native homodimer into a fusogenic homotrimer after entering cells by receptor-mediated endocytosis [7]. The conformational change occurs in the lower pH environment of the endosome where viral lipid envelope fusion with endosomal membrane, releasing the nucleocapsids into the cells cytoplasm [10]. Each E protein monomer has a molecular mass of 50-55 kDa and presents 3 distinct domains: domain I, II and III. Domain III is the immunoglobulin-like receptor binding domain [9] and is recognized by virus-neutralizing antibodies, being considered a target for diagnosis assays [8].

On the purpose of isolating viral parts and expressing them separately, different heterologous expression systems may be used and within these, the Baculovirus Expression System is one of the most popular and efficient.

Baculoviruses are large (30-60 × 250-300 nm), rod-shaped, double-stranded DNA (80-180 Kbp) viruses that are highly specific and only capable of replication in arthropod hosts [11]. This is why baculoviruses were first studied as biological control agents to protect crops and forests [12]. The Autographa californica multicapsid nucleopolyhedrovirus (AcMNPV) is the most studied baculovirus at the molecular level and most expression vectors are based on this virus [13]. There are many advantages of using the Baculovirus Expression System such as high expression levels and post-translational modifications that allows the expressed heterologous proteins to be correctly folded and biologically active [14].

In order to evaluate the potential antigenic activity of the Yellow fever E protein, in this study, we constructed a recombinant baculovirus containing a cDNA encoding the env gene of yellow fever virus. This recombinant virus was used to infect insect cells and larvae of susceptible insects and the protein produced was tested for its antigenicity.

Methods

Virus and Cells

Autographa californica multiple nucleopolyhedrovirus (AcMNPV), Anticarsia gemmatalis multiple nucleopolyhedrovirus (AgMNPV), and the recombinant viruses vSynScathL [15] and vSynYFE (constructed in this work) were propagated in the Trichoplusia ni cell line BTI-Tn5B1-4 (Tn5B) [16]. Tn5B cells were maintained in TC100 medium (Invitrogen) supplemented with 10% fetal bovine serum.

Envelope Gene Amplification and Cloning

Standard molecular cloning techniques were used as described in Sambrook et al. [17]. The DNA sequence encoding YFV env gene was isolated by RT-PCR. The RNA was extracted with TRIzol reagent (Invitrogen) from a sample of lyophilized sucking mice intracerebral tissue inoculated with YFV (kindly supplied by Evandro Chagas Institute, Brazil) and the complementar DNA (cDNA) was synthetized with ImPromII reverse transcriptase system (Promega) using only the reverse oligonucleotide YFE2460R (5'-CATGACCTCCTTAATCCGCCCAACTCC-3') which anneals downstream of the env gene. This cDNA was used in a PCR reaction using the specific oligonucleotides YFE878F (5'-GATCTCCTTAATCCGCCCAACTCC-3') and YFE2460R (5’-GA TATCTCCTAAATCCGCCCAACTCC-3’) located upstream and downstream of the env gene. The oligonucleotides were designed to incorporate the N and C-terminal hydrophobic regions of the E protein responsible for the ER translocation (amino acids 271 to 285) and membrane anchoring properties (amino acids 740 to 778), respectively [18]. The PCR program used was 94°C/1 min; (94°C/1 min, 50°C/2 min, 72°C/3 min) × 35; 72°C/10 min. An expected 1,590 bp PCR fragment
corresponding to env gene was amplified and cloned into the pGEM-T®-Easy vector (Promega), following the manufacturer’s instructions. This recombinant plasmid was designated pGEMYFE and utilized to amplify the envelope gene in E. coli DH5-α cells (Invitrogen).

**Construction of the transfer vector pSynYFE**

The 1,590 bp DNA fragment corresponding to the env gene was obtained by EcoRI digestion of the pGEMYFE DNA which was separated by electrophoresis in an agarose gel (0.8%) and the fragment corresponding to the env gene was purified from the gel using the GFX DNA extraction kit, according to the manufacturer’s instructions (GE Healthcare Life Science). The purified fragment was then ligated with the pSynXIV V1+X3 plasmid [19] previously linearized with EcoRI, resulting in the recombinant plasmid pSynYFE. This plasmid has the env gene under the control of two promoters in tandem (pSyn and pXIV). The pSyn/pXIV-env gene cassette region is flanked by baculovirus homologous regions that permit allelic replacement into the baculovirus genome during the co-transfection of insect cells with DNA from pSynYFE and the baculovirus vSynVIgal, a recombinant AcMNPV with the lacZ gene under the control of two promoters in tandem (AcMNPV [19]). The viral DNA was purified from disrupted virions as described in O’Reilly et al. [13]. The viral DNA was purified from disrupted virions as described in O’Reilly et al. [13], and used to perform a PCR reaction with the specific env oligonucleotides YFE878F and YFE2460R as described above.

**Heterologous protein expression analysis**

Tn5B cells were seeded at 1 × 10^6 cells per well in six well plates (TPP), mock infected and infected with wild type AcMNPV, vSynVI gal, *Anticarsia gemmatalis multiple nucleopolyhedrovirus* (AgMNPV), and the recombinants vSynYFE and vSynScathL. Cells were harvested at 96 h.p.i. and centrifuged at 750 × g for 7 min. The pellet was washed in PBS buffer (137 mM NaCl, 2.7 mM KCl, 10 mM Na₂HPO₄, 2 mM KH₂PO₄, pH 7.4) and aliquots from the samples were then analysed in a 12% SDS-PAGE gel [20] using a Mini Protean Tetra Cell apparatus (BioRad) following the manufacturer’s instructions. After electrophoresis, the gel was stained in a mixture of acetic acid:methanol:water (10:40:50) containing 0.1% Coomassie brilliant blue and destained afterwards in boiling water.

**Fluorescence analysis**

Tn5B cell monolayers (1 × 10^6 cells), grown over glass laminules placed inside sixty millimeter plates, were infected with vSynYFE (10 pfu/cell) or mock-infected (negative control). At 72 h.p.i, the laminules containing the cell monolayers were used for immunostaining. Cells were fixed with ice cold acetone for one hour (-20° C) and then incubated for 30 min with a PBS/BSA 10% solution at room temperature. The laminules containing cells were then incubated for 2 h with an E specific monoclonal antibody [21] which reacts with the envelope protein of the wild (Asibi) and vaccine strains of yellow fever virus with no cross reaction against other flavivirus or for 30 min with a polyclonal antibody for detection of flavivirus (anti-FLAV fluorescein isothiocyanate-conjugated antibody), currently used in a public diagnosis laboratory of reference (LACEN). The laminules incubated with the monoclonal antibody were washed 3 times with PBS (pH 7.4) and stained with DAPI (4,6-diamidino-2-phenylindole). Cells were again washed 3 times with PBS and incubated with the Alexa488-conjugated anti-mouse antibody (Invitrogen). Finally, cells were embedded with 1 drop of A31632 Component B Image IT TM Fx Signal Enhancer (Invitrogen) and 1 drop of N-Propyl gallate (0.2 g in 4 mL of PBS with glycerol-1 mL) and observed with a fluorescence (Axiohot, Zeiss) and/or confocal (The Broadband Confocal Leica TCS SP5, Leica DMI Series EL6000) microscopes with excitation and emission settings appropriate for the dyes used.
Structural and ultrastructural analyses of infected cells

Sixty millimeter plates (TPP) were seeded with Tn5B cells (1 × 10^6) and infected with vSynYFE (10 pfu/cell). Mock infected cells and vSynYFE infected cells were observed 72 h.p.i and photographed in a light Axiosphot microscope (Zeiss) and processed for electron microscopy. In brief, samples were fixed for 30 minutes (2% glutaraldehyde, 2% paraformaldehyde in 0.1 M sodium cacodylate buffer pH 7.4 with 5% sucrose), centrifuged at 750×g for 5 min, the pellet washed in the same buffer, post-fixed (1% osmium tetroxide, 0.8% potassium ferricyanide in the same buffer), contrasted in block with 0.5% uranyl acetate, dehydrated in acetone, and embedded in Spurr’s resin. The ultrathin sections were contrasted with uranyl acetate/lead citrate and observed in a TEM JEOL 100C and JEOL 1011 at 80 kV.

MAC-ELISA Immunoassay

The antigenic properties of the recombinant yellow fever virus envelope protein present in vSynYFE-infected insect cells were tested in a MAC-ELISA modified assay [22]. In brief, Tn5B cells were seeded at 1 × 10^6 cells per well in six well plates, mock infected and infected with wild type AcMNPV and the recombinant vSynYFE. Cells were harvested after 96 h.p.i and centrifuged at 750 × g for 7 min. The pellet was washed in PBS buffer and this procedure was repeated thrice. Cells were resuspended in a solution of PBS and human serum and reserved for a later time.

Two hundred μL of serum from two patients positively diagnosed for yellow fever were diluted 1:40 in PBS containing 0.5% BSA (bovine serum albumin). A volume of 50 μL of this sample was added to the wells and the plate was incubated for 1 hour in a moist chamber at 37°C. The plate was then washed five times with PBS. A volume of 50 μL of antigen was added to the plate. The different antigens used in this assay were described above and consists of the 3 insect cell extracts (mock-infected, AcMNPV-infected and vSynYFE-infected). The plate was incubated overnight in a moist chamber at 4°C and washed for 5 times with PBS. The anti-FLAV peroxidase conjugated antibody (Monoclonal Antibody 6B6C-1 Mab, HRP Conjugated, currently used for diagnosis of Flavivirus infection at LACEN, Brasília, Brazil) was added to the wells diluted 1/3000 in PBS plus 20% human serum. The plate was incubated for 1 h in a moist chamber at 37°C and then washed for 5 times with PBS. Peroxidase substrate solution, (100 μL of H_2O_2 and 2,2’ azinobis-3-ethylbenzthiazoline-6-sulphonic acid) was added and after 30 minutes at 37°C the OD (492 nm) of each well was measured thrice using a plate reader. All samples with an optical density above 0.2, higher than the negative control, were considered positive.

The same experiment was repeated with a pool of sera from three patients positively diagnosed for dengue fever (pool for dengue virus, DENV, sorotypes 1, 2 and 3, D1, D2, D3). Positive control was based in the reaction between the same pool of sera with a pool of DENV1, 2 and 3 (D1D2D3).

Results

Construction of the recombinant baculovirus vSynYFE

In order to construct the recombinant baculovirus vSynYFE, the DNA sequence encoding YFV envelope protein was isolated by RT-PCR from a sample of lyophilized suckling mice intracerebral tissue inoculated with YFV as a 1,590 bp band (Figure 1) and cloned (not shown) into the pGEM-T®-Easy vector (Promega). After transformation and amplification in E. coli (DH5-α, Invitrogen), the plasmid DNA was digested with EcoRI (data not shown) and the 1,590 bp fragment was cloned into the transfer vector pSynXIV V1+X3 generating the recombinant plasmid pSynYFE. The correct insertion of the env gene into the plasmid was confirmed by restriction enzyme digestion (data not shown). Finally, DNA from this plasmid was used to construct the vSynYFE recombinant baculovirus, by homologous recombination, after co-transfection with DNA from the vSynVI gal virus inside insect cells (Figure 1). To confirm the presence of the env gene into the recombinant virus, a PCR reaction was performed with viral DNA, released from the disrupted virions which amplified the expected 1,590 bp fragment (data not shown).

Detection of the recombinant protein expression

Insect cells extracts infected with vSynYFE were used in a SDS-PAGE gel to detect envelope protein and, when compared to wild type infected cells, they presented a protein band of around 54 kDa (Figure 2b). This band was not present in Tn5B cells mock infected neither in cells infected with AcMNPV, AgMNPV and a recombinant AcMNPV containing the ScathL gene from Sarco phaga peregrina (vSynScathL) [15]. The 54 kDa recombinant protein produced has the expected size for the recombinant protein.

Fluorescence analysis

In order to confirm the immunogenicity of the E recombinant protein expressed in insect cells, fluorescence and confocal microscopy were performed. Polyclonal and monoclonal antibodies against the E protein were able to detect the presence of the recombinant E protein in the citoplasm of vSynYFE-infected Tn5B cells (Figure 3 and 4) indicating that the recombinant protein is antigenically similar to its authentic counterpart.
Figure 1 Amplification of the envelope gene (env). Amplification of the Yellow Fever env gene by RT-PCR. (A) Agarose gel (0.8%) electrophoresis showing the RT-PCR amplification product (lane 2) using the oligonucleotides YFE2460R e YFE878F. Lane 1 shows the molecular mass marker, 1 Kb plus DNA ladder (Invitrogen). (B) Diagram of the Yellow Fever genome (Genebank accession # YFU17066) showing the main ORF (polyprotein ORF) and the position of the structural (Cap, pM and E) and non-structural (NS1, NS2a, NS2b, NS3, NS4a and NS5) proteins. The position of the oligonucleotides YFE2460R e YFE878F, used for the amplification reaction are also shown.

Figure 2 Construction of recombinant virus and production of the E protein. (A) Schematic representation of the construction of the recombinant virus vSynYFE, showing homologous recombination between the plasmid pSynYFE and the lacZ region (polh locus) in the virus vSynVI-gal. (B) SDS-PAGE analysis of E protein expression in insect cells. Lane 1, molecular weight marker (BenchMark™ Prestained Protein Ladder - Invitrogen); Lane 2, TnSB cells mock infected; Lane 3, TnSB cells infected with wild type AcMNPV; Lane 4, TnSB cells infected with vSynVI-gal; Lane 5, TnSB cells infected with Anticarsia gemmatalis nucleopolyhedrovirus (AgMNPV); Lanes 6, 7 and 8 TnSB cells infected with a recombinant AcMNPV containing the ScathL gene from Sarcophaga peregrina (vSynScathL) and Lane 9, TnSB cells infected with the recombinant vSynYFE. Arrow indicate a major polypeptide band of around 54 kDa in vSynYFE-infected TnSB cells.
Strutural and ultrastructural analysis of infected insect cells
Insect cells infected with vSynYFE showed multinucleated syncytial cells, which is a typical cytopathic effect of Flavivirus infection, which was not observed in Tn5B non-infected cells by light and electron microscopy (Figure 5).

MAC ELISA Immunoassay
In this assay, we used two different yellow fever-infected patients sera for the reaction with mock-infected, AcMNPV- and vSynYFE-infected insect cells extracts as antigens. OD values shown in figure 6 are averages of the three repetitions of experiment measures and error bars were determined using Graph Pad Prism Software (Graphpad Software, San Diego, CA). Mock- and AcMNPV-infected cells extracts showed low OD values (0.113 and 0.121 respectively) when compared to the vSynYFE-infected insect cells (0.214) extracts (Figure 6A). AcMNPV-infected cells extracts also showed low OD values using the second yellow fever-infected patient sera (0.081) when compared to the vSynYFE-infected insect cells (0.294) extracts (Figure 6B). We also carried out a second assay with a pool of sera from dengue virus-infected patients and purified dengue virus as antigens (Figure 7). No cross reactivity was detected with the vSynYFE-infected insect cells extracts tested, indicating that vSynYFE-infected insect cells extracts could be used in substitution of the whole yellow fever virus as antigen in this assay.

Discussion
In order to check the antigenic activity of a recombinant YFV E protein, we expressed this protein via a recombinant baculovirus (vSynYFE) in insect cells. Mammalian cells are very efficient to express E protein from yellow fever virus [23,24] and other flaviruses, such as DENV [25]. However, its use is much more expensive and cumbersome than insect cells which are also not infected by human pathogenic viruses [26]. Tn5B cells infected with vSynYFE expressed the predicted 54 kDa recombinant protein. The recombinant virus infected cells also showed the formation of multinucleated syncytial cells which is a typical cytopathic effect found in mosquito cells infected with Flavivirus [1]. In fact, this is one of the diagnosis techniques available for yellow fever. This infected cells were also positively immunostained with monoclonal and polyclonal antibodies raised against the E protein of the original Yellow fever virus and to flavivirus, respectively, and currently used in a diagnostic centre (LACEN) in the city of Brasilia, Brazil. Furthermore, in a MAC ELISA assay, two different
yellow fever-infected patients sera reacted positively with vSynYFE-infected insect cells extracts as antigens. This experiment would be more reliable if more patients sera were tested. Unfortunately, yellow fever positive serum is rare and most of it is distributed amongst reference centers of diagnosis and were not available. On the other hand, the recombinant virus infected insect cells extracts did not cross react with a pool of sera from dengue virus infected patients (Figure 7) which indicates that the recombinant envelope protein may respond antigenically very alike to what happens with the intact virus which is very relevant when considering that this might turn diagnosis tests more specific and less dubious. Cross reactivity with different Flavivirus is quite common during serological tests [27]. The use of recombinant proteins of the virus or even smaller parts of it, may diminish this kind of problem. It also could reduce costs too, since nowadays the use of the whole yellow fever virus as antigen for diagnosis is quite

---

**Figure 4** E protein detection of vSynYFE-infected insect cells by confocal microscopy. Tn5B cells were infected with vSynYFE (10 pfu/cell) and at 72 h p.i., cells were fixed with ice cold acetone and incubated with an E specific monoclonal antibody. Bound monoclonal anti-E antibodies were detected using Alexa488-conjugated anti-mouse antibodies by confocal microscopy (green). Cell nuclei were visualized by DAPI staining (blue). (A) Tn5B cells mock infected (bright field). (B) Tn5B cells mock infected (DAPI staining). (C) Tn5B cells mock infected (merge). (D) Tn5B cells infected with vSynYFE (brightfield). (E) Tn5B cells infected with vSynYFE (DAPI). (F) Tn5B cells infected with vSynYFE (merge). In A, B, C, D, E and F, Bars = 50 μm. Blue staining (DAPI) indicates cells nuclei.

**Figure 5** E protein structural and ultrastructural analysis of vSynYFE-infected insect cells. Tn5B cells were infected with vSynYFE (10 pfu/cell) and at 72 h p.i., the cells were photographed in a light microscope and processed for electron microscopy. Multinucleated syncytial Tn5B cells were easily observed by light (A) and electron microscopy (B). Bar in (A) = 14 μm and in (B), 5 μm. Arrows indicates cells nuclei.
expensive since it requires a biosafety structure, with animal use and care.

Quicker diagnosis would mean quicker disease control. Subnotificated cases leads to delayed emergency response and the risk of reurbanization of the disease since the urban vector has adapted to living among humans in domestic environments. The use of recombinant envelope protein could be extended in the development of a safer vaccine which did not include the active virus at all. Especially when severe adverse events are being associated with the actual yellow fever vaccine with the attenuated virus, such as hypersensitivity reactions, associated neurotropic disease and associated viscerotropic disease [28].

Després et al. [29] used baculovirus to express and characterize yellow fever proteins E and NS1 (a non-structural yellow fever virus protein identified as the soluble complement-fixing antigen) and showed that insect cells infected with the recombinant baculoviruses synthesized E and NS1 proteins that were similar in size and antigenicity to those expressed in Vero cells infected with YFV. Their work resembled the methods used is this work but differed in the host cell type used, which were *S. frugiperda* insect cells, the vector used for the recombinant virus construction and the length of the E protein gene used. In their work, no syncytial formation in virus-infected *S. frugiperda* cells was reported. This could be explained by the expression of a truncated version of the E protein without the C-terminal hydrophobic region (amino acids 740 to 778) responsible for the transmembrane localization of the protein inside intracellular membranes which may be important for the oligomerization and folding of the E into a biological active protein [18].

The same research group injected recombinant virus-infected insect cell extracts into mice and conducted a fatal YFV challenge [29]. The result was a solid protection against lethal YFV encephalitis, which reinforce the recombinant protein antigenicity when compared to the whole virus. Shiu et al. [30], also expressed recombinant E protein in insect cells using recombinant baculovirus. The recombinant virus genome contained a cDNA coding for amino acids 109 to 820 of the YFV polyprotein. This region has a N-terminal truncated pM, the complete M and E proteins and the first 42 amino acids of the NS1 protein. This recombinant protein expressed in insect cells was shown to be antigenically indistinguishable from the E protein of yellow fever vaccine virus.

The actual yellow fever vaccine strain is considered to be very safe, in fact, its considered to be one of the safest vaccines ever developed [31]. Its production has been administered to man since the late 1930s [24] and a single dose of 17D yellow fever vaccine confers long-term immunity, with some data confirming cases exceeding the expected 10 years of protection [32]. However, as it still makes use of attenuated virus, some adverse events might result from its use [33] causing viscerotropic and neurotropic adverse events and anaphylaxis [34]. Yellow fever vaccine associated viscerotropic disease (YEL-AVD) resembles wild yellow fever symptoms with a similar fatality rate while yellow fever vaccine associated neurologic disease (YEL-AND) may manifest encephalitis, meningitis, neuropathy, Guillain-Barre syndrome, acute disseminated encephalomyelitis or spinal myelitis [35]. Their occurrence varies from 0.4 to 7.9 cases per 100,000 vaccine doses and some
alternatives are being studied in order to minimize the risks of vaccine administration specially for the ones with a compromised immunologic system such as elderly people, immunosuppressed or thymectomized patients, infants under 9 months of age and pregnant or nursing women. One alternative would be the administration of inactivated virus which is already being tested in clinical trials [34]. Other would be to use viral-like particles as immunogens.

Viral-like-particles (VLP) produced in insect cells have been used in subunit vaccine trials for the influenza infection and preliminary studies demonstrated that this VLPs are capable of inducing immunity against some strains of influenza virus [36]. Gut-Winiarska et al. [37] developed a direct sandwich blocking ELISA diagnostic test using a recombinant glycoprotein B (gB) expressed in baculovirus to detect pseudorabies virus. It showed high sensitivity and specificity when compared to two other commercial tests using the whole virus. Recently, a baculovirus vector based vaccine conferring protection against 

Human papillomavirus (HPV) (causative agent of cervical neoplasia) was approved for use in Brazil and the Brazilian Ministry of Health is considering to adopt it in its current annual vaccination program [38].

Conclusions

In conclusion, our data shows that the recombinant E protein derived from YFV expressed in insect cells is recognized by monoclonal and polyclonal antibodies used to detect YFV infection in human sera and therefore, may be useful for different medical applications, from improved diagnosis of the diseases to source of antigens for the development of a subunit vaccine, which might be useful in the future to promote immunization of individuals contraindicated to receive the current yellow fever attenuated virus vaccine.

Acknowledgements

Many thanks to LACEN-DF for gently supplying of anti-FLAV fluorescein isothiocyanate-conjugated antibody. This work was supported by the following Brazilian agencies: CNPq, CAPES, FAPDF.

Author details

1 Cell Biology Department, University of Brasília, Brasília, DF, CEP 70910-970, Brazil. 2 Laboratório Central do Distrito Federal - LACEN-DF, Brasília, DF, CEP 70830-010, Brazil. 3 Institut de Recherches pour le Développement UNIR182, LOCEAN-IJPSL UPVLC, 4 pl. Jussieu, case 100, Paris, 75252 Cedex 05, France.

Authors’ contributions

MCESB and TGCMG carried out the study, performed analysis of data and drafted the manuscript. TN helped with the construction of recombinant viruses and with the confocal analysis of virus-infected cells. AJMC helped with the MAC ELISA assay protocol and provided hyperphized sucking mice intracerebral tissue inoculated with YFV ND provided part of research funds and discussed partial results. BMR conceived the study, provided research funds, supervised students and revised the manuscript. All authors have read and approved the final manuscript.

Competing interests

The authors declare that they have no competing interests.

Received: 3 February 2011 Accepted: 27 May 2011 Published: 27 May 2011

References

1. Vasconcelos PFC. Febre amarela: reflexões sobre a doença, as perspectivas para o século XXI e o risco da reurbanização. Rev Bras Epidemiol 2002, 5(3):244-258.
2. Bryant JE, Barnett ADT. Comparative phylogenies of yellow fever isolates from Peru and Brazil. FEMS Immunol Med Microbiol 2003, 39:103-118.
3. Situação da Febre Amarela Silvestre no Brasil, 2007 e 2008, [http://portal.saude.gov.br/portal/paroquis/pdf/boletim_sms_febre_amarela_040408.pdf].
4. De Paula SO, Fonseca BAL. Dengue: a review of the laboratory tests a clinician must know to achieve a correct diagnosis. Braz J Infect Dis 2004, 8(6):390-398.
5. Vasconcelos PFC. Febre amarela. Rev Soc Bras Med Trop 2003, 36(2):275-293.
6. Fauguet CM, Mayo MA, Maniloff J, Dessellenger U, Ball LA. Virus Taxonomy: eighth report of the International Committee on Taxonomy of Viruses San Diego: Elsevier Academic Press, 2005.
7. Mayhob AS, Khaliq M, Kuhn RJ, Cushman M. Design, Synthesis, and Biological Evaluation of Thiocolesterol Targeting Flavivirus Envelope Protein. J Med Chem 2011, 54:1044-1054.
8. Chávez JH, Silva JR, Amariolla AA, Figueiredo LTM. Domain III peptides from flavivirus envelope protein are useful antigens for serologic diagnosis and targets for immunization. Biologicals 2010, 38:613-618.
9. Bonaldo MC, Garrart RC, Freire MS, Galler R. Expression of Foreign Protein Epitopes at the Surface of Recombinant Yellow Fever 17D Viruses Based on Three-Dimensional Modeling of its Envelope Protein. Cell Biochemistry and Biophysics 2006, 44:313-324.
10. Gardner CL, Ryman KD. Yellow Fever: A Reemerging Threat. Cln Lab Med 2010, 30:237-260.
11. Rohrmann GF. Baculovirus Molecular Biology. Bethesda (MD): National Library of Medicine (US), National Center for Biotechnology Information, 2008 [http://www.ncbi.nlm.nih.gov/bookshelf/br.fcgi?book=bacvir].
12. Micocci F. Assessment of the application of baculoviruses for control of Lepidoptera. Annu Rev Entomol 1999, 44:257-289.
13. O’Relly DR, Miller LK, Luckow VA: Baculovirus Expression Vectors: A Laboratory Manual New York: WH, Freeman and Company, 1992.
14. Kost TA, Cordray JP, Jarvis DL: Baculoviruses as versatile vectors for protein expression in insect and mammalian cells. Nat Biotechnol 2005, 23(9):567-575.
15. Gramkow AW, Perecmanis S, Sousa RL, Noronha EF, Felix CR, Nagata T, Ribeiro BM: Insecticidal activity of two proteases against Spodoptera frugiperda larvae infected with recombinant baculoviruses. Viral J 2010, 29(7):143.
16. Granados RR, Guoxun L, Derskove AC, McKenna KAA: A new insect cell line from Trichoplusia ni (BTI-Tn-5B1-4) susceptible to Trichoplusia ni single nuclear polyhedrosis virus. J Invertebr Pathol 1994, 64:260-266.
17. Sambrook J, Fritsch EF, Maniatis T: Molecular Cloning: a laboratory manual. New York: Cold Spring Harbor Laboratory Press, 1989.
18. Ruiz-Linares A, Cahour A, Després P, Girard M, Bouloy M: Processing of yellow fever virus polyprotein: role of cellular proteases in maturation of the structural proteins. J Virol 1989, 63(10):4199-4209.
19. Wang X, Ooi BG, Miller LK: Baculovirus vectors for multiple gene expression and for occluded virus production. Gene 1991, 100:131-137.
20. Laemmli UK: Cleavage of structural proteins during assembly of the head of bacteriophage T4. Nature 1970, 227:680-685.
21. Schlesinger JJ, Brandis MW, Monath TP: Monoclonal antibodies distinguish between wild and vaccine strains of yellow fever virus by neutralization, hemagglutination inhibition, and immune precipitation of the virus envelope protein. J Virol 1983, 125(1):8-17.
22. Vázquez S, Valdés O, Pupo M, Delgado I, Alvarez M, Pelegirno JL, Guzmán MG: MAC-ELISA and ELISA inhibition methods for detection of antibodies after yellow fever vaccination. J Virol Methods 2003, 110(2):179-184.
23. Després P, Cahour A, Wychowski C, Girard M, Bouloy M: Expression of the yellow fever virus envelope protein using hybrid SV40/yellow fever viruses. Ann Inst Pasteur Virol 1988, 139(1):59-67.
24. Nogueira RT, Nogueira AR, Pereira MCS, Rodrigues MM, Galler R, Bonaldo MC: Biological and immunological characterization of recombinant Yellow Fever 17D Viruses expressing a Trypanosoma cruzi Amastigote Surface Protein-2 CD8+ T cell epitope at two regions of the genome. Virol J 2011, 8:127-140.

25. Jaiswal S, Khanna N, Swaminathan S: Replication-defective adenoviral vaccine vector for the induction of immune responses to Dengue virus type 2. J Virol 2003, 77(23):12907-12913.

26. Morton CI, Potter PM: Comparison of Escherichia coli, Saccharomyces cerevisiae, Pichia pastoris, Spodoptera frugiperda, and COS7 Cells for Recombinant Gene Expression: Application to a Rabbit Liver Carboxylesterase. Mol Biotech 2000, 16(3):193-202.

27. Monath TP: Yellow fever. Medicine 2005, 83(7):21-23.

28. Barnett ED: Yellow fever: epidemiology and prevention. Clin Infect Dis 2007, 44(6):850-856.

29. Després P, Dietrich J, Girard M, Bouloy M: Recombinant baculoviruses expressing yellow fever virus E and NS1 proteins elicit protective immunity in mice. J Gen Virol 1991, 72:2811-2816.

30. Shiu SYW, Monicka S, Buckley A, Higgs S, Karunakaran N, Blachere C, Gould EA: 17D yellow fever vaccine virus envelope protein expressed by recombinant baculovirus is antigenically indistinguishable from authentic viral protein. J Gen Virol 1991, 72:1451-1454.

31. Struchiner CJ, Luz PM, Dourado I, Sato HK, Aguiar SG, Ribeiro JG, Soares RC, Codeço CT: Risk of fatal adverse events associated with 17DD yellow fever vaccine. Epidemiol Infect 2004, 132(5):939-946.

32. Bodilis HC, Benabdelmoumen G, Gergely A, Goujon C, Pelicot M, Poujol P, Consigny PH: Persistance à long terme des anticorps neutralisants de la fièvre jaune chez les personnes âgées de 60 ans et plus: Long term persistence of yellow fever neutralising antibodies in elderly persons. Bull Soc Pathol Exot 2010, 1:1-6.

33. Eventos adversos pós-vacinação com a vacina contra febre amarela. [http://portal.saude.gov.br/portal/arquivos/pdf/nota_FA_19112007.pdf].

34. Monath TP, Fawler E, Johnson CT, Baiser J, Morin MJ, Sisti M, Trent DW: An Inactivated Cell-Culture vaccine against Yellow Fever. N Engl J Med 2011, 364(14):1326-1333.

35. Hayes EB: Is it time for a new yellow fever vaccine? Vaccine 2010, 28:8073-8076.

36. Quan FS, Huang C, Companis RW, Kang SM: Virus-like particle vaccine induces protective immunity against homologous and heterologous strains of influenza virus. J Virol 2007, 81(7):3514-3524.

37. Gut-Winiarska M, Jacobs L, Kerstens H, Bienkowska-Szewczyk K: A highly specific and sensitive sandwich blocking ELISA based on baculovirus expressed pseudorabies virus glycoprotein B. J Virol Methods 2000, 88:63-71.

38. European Medicines Agency: Cervarix suspension for injection - Human Papillomavirus vaccine [Types 16, 18] (Recombinant, adjuvanted, adsorbed). European public assessment Reports [http://www.ema.europa.eu/docs/en_GB/document_library/EPAR_-_Product_Information/human/000721/WC500024632.pdf].

Cite this article as: Barros et al. Yellow fever virus envelope protein expressed in insect cells is capable of syncytium formation in lepidopteran cells and could be used for immunodetection of YFV in human sera. Virology Journal 2011 8:261.

DOI: 10.1186/1743-422X-8-261