

**Optimization of small-scale production of Zika virus envelope glycoprotein  
by transient expression in HEK293 cells for ELISA**

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Running title: Zika virus envelope protein production for ELISA

## Abstract

Zika virus (ZIKV) is an emerging mosquito-borne flavivirus which has recently caused global epidemics with its association with congenital Zika syndrome such as severe microcephaly. The recombinant ZIKV envelope (Env) glycoprotein is useful for immunological applications such as serodiagnosis of ZIKV infection and for monitoring immune responses in preclinical and clinical ZIKV vaccine developments. In this chapter, we describe the optimization of production of Zika virus envelope glycoprotein in Human Embryonic Kidney (HEK 293T) cells by small-scale expression followed by large-scale protein production. Small-scale expression of HEK 293T cells allows screening of a large number of vectors simultaneously to select the vectors with best secretory profiles for scale-up in Expi293 mammalian system to maximise the protein yield followed by purification for research and clinical applications.

Keywords: Zika virus, envelope protein, HEK 293, CD4 fusion tag, protein production, protein purification.

## 1. Introduction

Optimum expression of proteins with high purity levels can be useful reagents for a wide array of applications such as immunological assays, protein-based vaccines or therapeutics and structural studies. Protein expression in mammalian cells is preferred to *E. coli* cells for correct post-translational modifications. Various commercially available Zika virus (ZIKV) envelope (Env) glycoproteins are expressed in insect cells, reaching a purity of approximately 85% but yet they are not easily affordable in developing countries endemic with ZIKV.

In this chapter, protocols for the production, purification and characterisation of secreted ZIKV Env glycoproteins by transient transfection of HEK 293 cells and optimisation for its use in ELISA for ZIKV serodiagnosis are described. The methods are exemplified by reference to the work by Kim et al [1]. ZIKV Env is the primary target for neutralizing antibodies during ZIKV infection and involved in the receptor binding and viral entry into host cells [2, 3]. In order to produce recombinant ZIKV Env glycoproteins for ZIKV serodiagnosis and monitoring of humoral responses in both mice and humans vaccinated with Env-based vaccines, the secretory profile of various constructs (prM-Env, Env only, and truncations of Env) with or without a rat CD4 fusion tag at the C-terminus were tested in small-scale expression screening to select the construct with the best secretion profile. The best secretion vector was chosen to be expressed at larger scale using the Expi293 mammalian transient expression system and purified by immobilized metal affinity chromatography (IMAC) and size exclusion chromatography (SEC) to obtain highly purified fractions containing ZIKV envelope proteins (Env-CD4 and Env) as seen in the Figure 1. These ZIKV Env-CD4 recombinant proteins were characterised by western blotting (WB) using a pan flavivirus antibody, the sera from ChAdOx1 ZIKV prME\_ΔTM vaccinated mice [4] and a ZIKV Env monoclonal antibody (Figure 2). ZIKV Env recombinant protein was then used as coating reagent for ELISA assay using sera from ZIKV-infected patients to detect patient-derived ZIKV antibodies and optimised for sero-diagnosis of ZIKV infection (Figure 3).

## **2. Materials**

### **2.1 Small-Scale Cell Transfection**

1. HEK 293T cells.
2. Cell culture medium: Dulbecco's Modified Eagles Medium (DMEM) supplemented with 10% Foetal Calf Serum (FCS), non-essential amino acids (1:100) and L- Glutamine (1:100).

- 78 3. Trypsin- Ethylene dinitrilotetra-acetic acid (Trypsin- EDTA).
- 79 4. Phosphate buffered saline (PBS): 0.01 M phosphate buffer, 0.0027 M potassium chloride,
- 80 0.137 M sodium chloride, pH 7.4.
- 81 5. T175 tissue culture flask.
- 82 6. Plasmid DNA (*see Note 1*) with an  $A_{260}/A_{280}$  ratio of greater than 1.8 (*see Note 2*).
- 83 7. Transfection reagent.
- 84 8.V-well micro-titre plate.
- 85 9. 24-well tissue culture plate.
- 86 10. Tissue culture incubator at 37°C and 5% CO<sub>2</sub>.

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## 88 **2.2 Large-Scale Cell Transfection**

- 89 1. Expi293™ cells.
- 90 2. Expi293™ Expression Medium.
- 91 3. OPTI-MEM medium.
- 92 4. Expifectamine™ transfection reagent. (*see Note 3*).
- 93 5. Enhancer 1 and 2.

94

## 95 **2.3 SDS-Polyacrylamide Gel Electrophoresis (SDS-PAGE) and Western Blotting (WB)**

- 96 1. 2× SDS-PAGE sample loading buffer: 100 mM Tris-HCl pH 6.7, 40 mg/ml sodium dodecyl
- 97 sulphate (SDS), 2 mg/ml bromophenol blue, 20% v/v glycerol, in water.
- 98 2. Heating block
- 99 3. SDS-PAGE MOPS running buffer.
- 100 4. SDS-PAGE gel.

- 101 5. Nitrocellulose or PVDF membrane.
- 102 6. Protein stain.
- 103 7. Phosphate buffered saline with Tween (PBST): 0.01 M phosphate buffer, 0.0027M
- 104 potassium chloride, 0.137M sodium chloride, 0.05% v/v Tween 20, pH 7.4.
- 105 8. 5% milk in PBST
- 106 9. Primary antibody solution: 1:2000 Anti-His<sub>6</sub> mouse monoclonal antibody or 1:500 dilutions
- 107 of a mouse anti-flavivirus group antigen monoclonal antibody (Millipore) or 1:500 dilutions of
- 108 mice sera following a vaccination with ChAdOx1 ZIKV prME\_ΔTM [4] or 1:1000 anti-Zika
- 109 Env monoclonal antibody (mouse mAb to Zika Env protein, Aalto BioReagents).
- 110 10. Secondary antibody solution: 1:5,000 Anti Mouse-Goat IgG Peroxidase conjugate.
- 111 11. Western blotting detection system.

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## 113 **2.4 Large Scale Glycoprotein Purification**

- 114 1. 5 ml HisTrap FF column.
- 115 2. HiLoad 16/600 Superdex 200 column.
- 116 3. ÄKTA purification system such as ÄKTAexpress™
- 117 4. Nickel Wash buffer: 50mM Tris-HCl, 500mM NaCl, 30mM imidazole, pH 7.5 (*see Note 4*).
- 118 5. Nickel Elution Buffer: 50mM Tris-HCl, 500mM NaCl, 500mM imidazole, pH 7.5.
- 119 6. Size Exclusion Buffer: 20mM Tris-HCl, 200mM NaCl, pH 7.5.
- 120 7. 96 deep well plate

121

## 122 **2.5 Enzyme-linked immunosorbent assay (ELISA)**

- 123 1. 96-well Elisa plates.

2. Phosphate buffered saline (PBS): 0.01 M phosphate buffer, 0.0027 M potassium chloride, 0.137 M sodium chloride, pH 7.4.
3. Phosphate buffered saline with Tween (PBST): 0.01 M phosphate buffer, 0.0027M potassium chloride, 0.137M sodium chloride, 0.05% v/v Tween 20, pH 7.4.
4. Protein-Free Blocking Buffers.
5. Anti-Human IgG-alkaline phosphatase-conjugated antibody.
6. Buffer solution: Prepare pNPP substrate at 1mg/mL by dissolving each pNPP tablet in 20ml Diethanolamine Substrate Buffer. Leave to dissolve at RT in the dark for one hour.
7. Elisa plate reader.
8. Naïve control sera from healthy volunteers.
9. Zika patient samples.

### **3. Methods**

Small-scale expression allows for rapid assessment of expression level from a large number of plasmids in parallel. This parallel expression screening in HEK 293T was used in the work of Kim et al [1] to screen constructs encoding different lengths of ZIKV prM-Env and Env. A protocol for the comparative small-scale expression screening of plasmids using HEK 293T cells is given in Section 3.1 followed by methods for sample preparation and Western blot analysis of secreted product in Section 3.2. The expression vectors showing the best secretion profile can then be scaled-up for production and purification in Section 3.3 and 3.4.

#### **3.1 Transfection of Cells in 24 well plates for small-scale expression screening**

1. Seed 1 ml of HEK 293T cells into a 24-well tissue culture plate at a density  $\sim 1.5 \times 10^5$  cells / ml to give  $\sim 70\%$  confluency after 24 h incubation at 37°C.

- 148 2. After 24 h incubation, mix an appropriate amount of transfection reagent (depending on  
149 manufacturer's instructions) in DMEM supplemented with 1mM glutamine and 1 × non-  
150 essential amino acids in a V-well micro-titre plate.
- 151 3. Add ~ 1µg of plasmid DNA and mix thoroughly by mixing up and down for ~10 times and  
152 incubate for 30 min at room temperature.
- 153 4. Carefully aspirate the medium from the cell layer in the 24-well plate and discard.
- 154 5. Add sufficient cell culture medium to the DNA / transfection reagent cocktail to achieve a  
155 final volume of 1 ml. Add to the plated cells.
- 156 6. Incubate the cells for 72 h at 37°C in a 5% CO<sub>2</sub>/95% air incubator.

157

### 158 **3.2 Harvest and Western Blot Analysis of Secreted Proteins**

- 159 1. To analyse the expression of secreted proteins, harvest the culture supernatant from the cells  
160 in the 24 well plate by centrifuging at 17,000 × g for 10 min. Mix an equal volume of  
161 supernatant and 2× SDS-PAGE sample loading buffer and denature the protein by heating to  
162 95°C for 3 minutes.
- 163 2. Run the samples on an SDS-PAGE gel and transfer to a nitro-cellulose or PVDF membrane  
164 using standard Western blotting procedures (*see Note 5*).
- 165 3. Block the membrane for at least 30 min with 5% milk in PBST at room temperature.
- 166 4. Remove the blocking solution and add 1:2000 Anti-His<sub>6</sub> mouse monoclonal antibody as the  
167 primary antibody and incubate for 1h at room temperature on a shaker (*see Note 6*).
- 168 5. Wash the membrane three times in PBST for 5 min each time.
- 169 6. Remove the wash buffer and incubate the membrane for 1 h in the secondary antibody  
170 solution at room temperature on a shaker.

171 7. Wash the membrane three times in PBST for 5 min followed by two washes for 5 min each  
172 in PBS.

173 8. To visualise the protein, remove excess buffer from the membrane and cover with Western  
174 blotting detection system reagent for 5 min to develop the blot.

175 9. Signal is detected using a chemiluminescent Western blot imaging system.

176

### 177 **3.3 Large-Scale Transient Expression**

178 1. Seed Expi 293 cells in a roller bottle with 300 ml culture volumes at a density of  $\geq 1.0$ - $1.5$   
179  $\times 10^6$  cells/ml at  $\sim 95\%$  viability for 24 h.

180 2. Determine cell count and viability and prepare the transfection as below. Transfect if cell  
181 count  $\geq 1.5$ - $2.0 \times 10^6$  cells/ml at  $\sim 95\%$  viability.

182 3. Prepare transfection cocktail as follows: Solution A - Add 800  $\mu$ l of Expifectamine  
183 transfection reagent in a Falcon tube and dilute with 15 ml OPTI-MEM medium. Incubate for  
184 5 min at room temperature. Solution B - Add 300  $\mu$ g of DNA in a tube and dilute with 15 ml  
185 OPTI-MEM medium.

186 4. Mix solution A and B and incubate for 10 min at room temperature (*see Note 7*).

187 5. From the 300 ml suspension culture to be transfected, remove 30 ml of cells before adding  
188 the transfection cocktail to make up the final volume of 300 ml culture.

189 6. Incubate transfected cells for 16 - 18 hours at 37°C.

190 7. After 16-18 h, add 1.5 ml of Enhancer 1 and 15 ml Enhancer 2 to the culture and return to  
191 the incubator and incubate for 2 days (*see Note 3*).

192 8. Determine cell count and viability. Harvest media by spinning in 500 ml centrifuge bottle  
193 for 45 min at 6000 x g. Filter sterilise using a 500 ml 0.45  $\mu$ m bottle top filter and proceed to  
194 purification.



### **3.4 Automated purification protocol using the ÄKTAexpress™ for large-scale glycoprotein purification**

1. Using an AKTAexpress™ unit, equilibrate the HiLoad16/600 Superdex 200 column with 2 column volumes of Size Exclusion Buffer.
2. After equilibration of the size exclusion column, insert the inlet tubes A1 and A2 into Nickel Wash Buffer and wash the pumps manually to fill the lines with buffer.
3. Insert A3 into Nickel Elution Buffer and insert the outlet tube F3 into a duran glass flask that is at least 1.5 times the sample volume and place a 96 deep well plate in the fraction collector.
5. Insert a pre-charged 5 ml HisTrap FF column into column position 1.
6. Carefully remove the line A2 from the Nickel Wash Buffer and insert it into the flask containing the sample.
7. Using a Method Run in the System Control section of the Unicorn™ software, run the glycoprotein purification programme described in Nettleship *et al.* [5] (*see Note 8*).

### **3.5 SDS-PAGE Analysis of Large-Scale Purified Proteins**

1. Analyse the A280 trace of the size exclusion profile and make a note of the fractions that contain protein, as represented by a peak in the A280 trace.
2. Take 10 µl from each fraction-of-interest and mix with 10 µl of 2× SDS-PAGE sample loading buffer. Denature the protein by heating the sample to 95°C for 3 min.
3. Run the samples on an SDS-PAGE gel until the dye reaches the bottom. Wash the gel three times in water for 1 min each time and add 20 ml of protein stain and incubate at room temperature with rocking for at least 15 min.
4. To de-stain the gel, remove the stain and incubate the gel in water at room temperature with rocking for 15 min or more until a clear background is obtained.

5. Combine fractions containing the protein of interest by SDS-PAGE.

### **3.6 Western Blot Analysis of Large-Scale Purified Proteins**

1. To characterise the purified proteins, similar western blot is carried out according to the protocol in Section 3.2 using a pan flavivirus antibody, the sera from ChAdOx1 ZIKV prME\_ΔTM vaccinated mice and a ZIKV Env monoclonal antibody as the primary antibody.

### **3.7 Serodiagnosis of ZIKV by ELISA**

Purified ZIKV Env is used as a coating reagent for immunoassay (ELISA) using sera from ZIKV-infected patients from an endemic area to detect patient-derived ZIKV antibodies and compared to the commercially available ZIKV Env (Figure 3).

1. Coat Elisa plates with 5 µg/ml antigen diluted in PBS; add 50µL per well (*see Note 9*).
2. Leave plates at room temperature on bench overnight.
3. Wash plates 6 times with PBST by adding PBST to completely fill plates to the top and discarding it.
4. Block plates with 300 µl per well Blocking Buffer for 2h at room temperature.
5. Dilute sera sample starting at 1:300 in PBST.
6. Discard blocking solution.
7. Add 50 µl of PBST to wells in rows B-H.
8. Add 75 µl of sera to wells in row A and serially dilute 3-fold down plate in PBST by transferring 25 µl into row below, mixing 20 times and transferring again. Throw away the 25 µl taken out of row H. Final volume should be 50 µl per well.
9. Leave 1h at RT.

- 243 10. Wash plates 6 times with PBST by adding PBST to completely fill plates to the top and  
244 discarding it.
- 245 11. Add 50  $\mu$ l per well the detecting Ab– Anti-Human IgG-alkaline phosphatase-conjugated  
246 antibody diluted 1/5000 in PBST (5 mL per plate + 1  $\mu$ L Ab). Leave 1h at RT.
- 247 12. Wash plates 6 times with PBST by adding PBST to completely fill plates to the top and  
248 discarding it.
- 249 13. Add pNPP substrate, 100  $\mu$ l per well. (*see Note 10*).
- 250 14. Cover with foil and leave for 15 minutes to develop at room temperature. (*see Note 11*).
- 251 15. Read OD at 405nm on Elisa plate reader.
- 252 16. Serum antibody endpoint titers were defined by absorbance value three standard deviations  
253 greater than the average OD405 of control sera pool.

254

## 255 **Notes**

- 256 1. The selected prM-Env gene was cloned into pOPINTTGneo or pOPINTTGneo-3C-CD4  
257 expression vectors which produce soluble ZIKV Env proteins with a C-terminal His tag or a  
258 C-terminal rat CD4 fusion and His tag.
- 259 2. DNA for transfection needs to have an  $A_{260}/A_{280}$  ratio of  $> 1.8$ . This can be obtained using  
260 standard commercial kits such as the PureLink HiPure Expi Plasmid Megaprep kit.
- 261 3. As an alternative to using the ExpiFectamin™ kit, 300 ml culture volume of cells are  
262 transfected by adding 30 ml of OPTI-MEM containing 300  $\mu$ g of DNA and 1.6 ml of PEI Max  
263 40kDa (1 mg/mL). The following day, 5 ml of 350 mM Valproic acid, 2 ml 1M sodium  
264 propionate and 5.5 ml of 45% w/v glucose are added to enhance expression.
- 265 4. The buffers are vacuum filtered using 0.45  $\mu$ m filters to remove unwanted dust and bubbles.

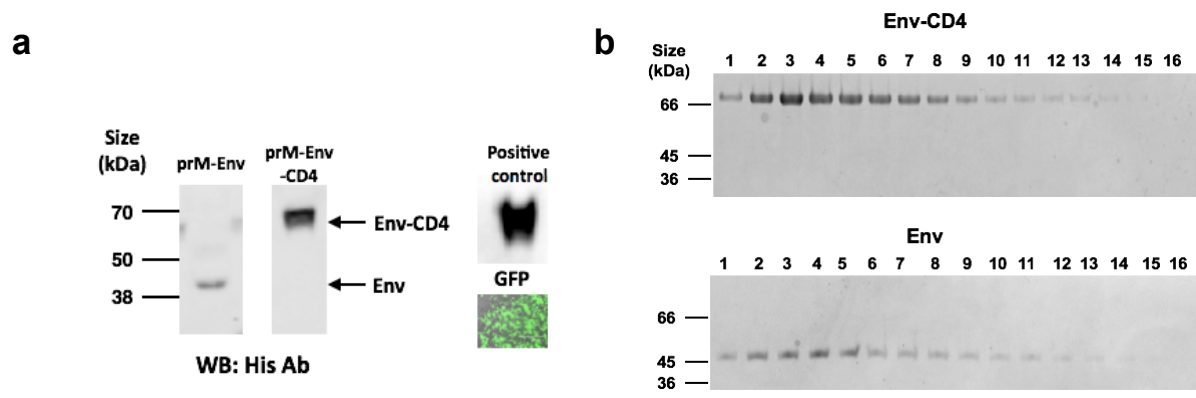
5. Trans-Blot Turbo Mini Nitrocellulose Transfer Packs were used to transfer Env proteins from gels using the Trans-Blot Turbo Transfer System in Kim et al [1]. Alternatively, standard western blotting method using PVDF membranes can be used as described in Mahmood et al [6].
6. The amount of antibody depends on the size of the container the membrane is in, for example, 1:2000 dilution of Anti-His<sub>6</sub> mouse monoclonal antibody in 30 ml 5% milk PBST can be made by adding 15 µl of antibody in 30 ml 5% PBST.
7. Incubation longer than 10 min can result in loss of transfection efficiency.
8. The full method for the glycoprotein purification is written out in Nettleship et al [5] and can be copied into the Method Editor section of the Unicorn<sup>TM</sup> software.
9. Concentration of coating ZIKV Env may vary depending on antigen used.
10. Add substrate at least 1 minute apart to each plate so that all plates can be developed for the same length of time and OD values are read sequentially on the ELISA reader.
11. Development time is antigen/user dependent

## References

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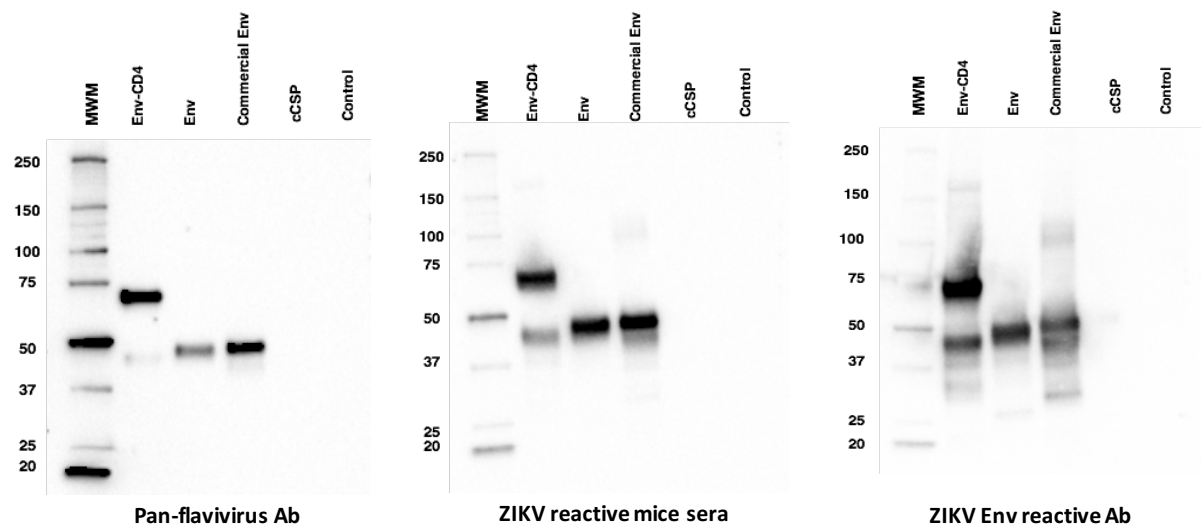
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## Figures



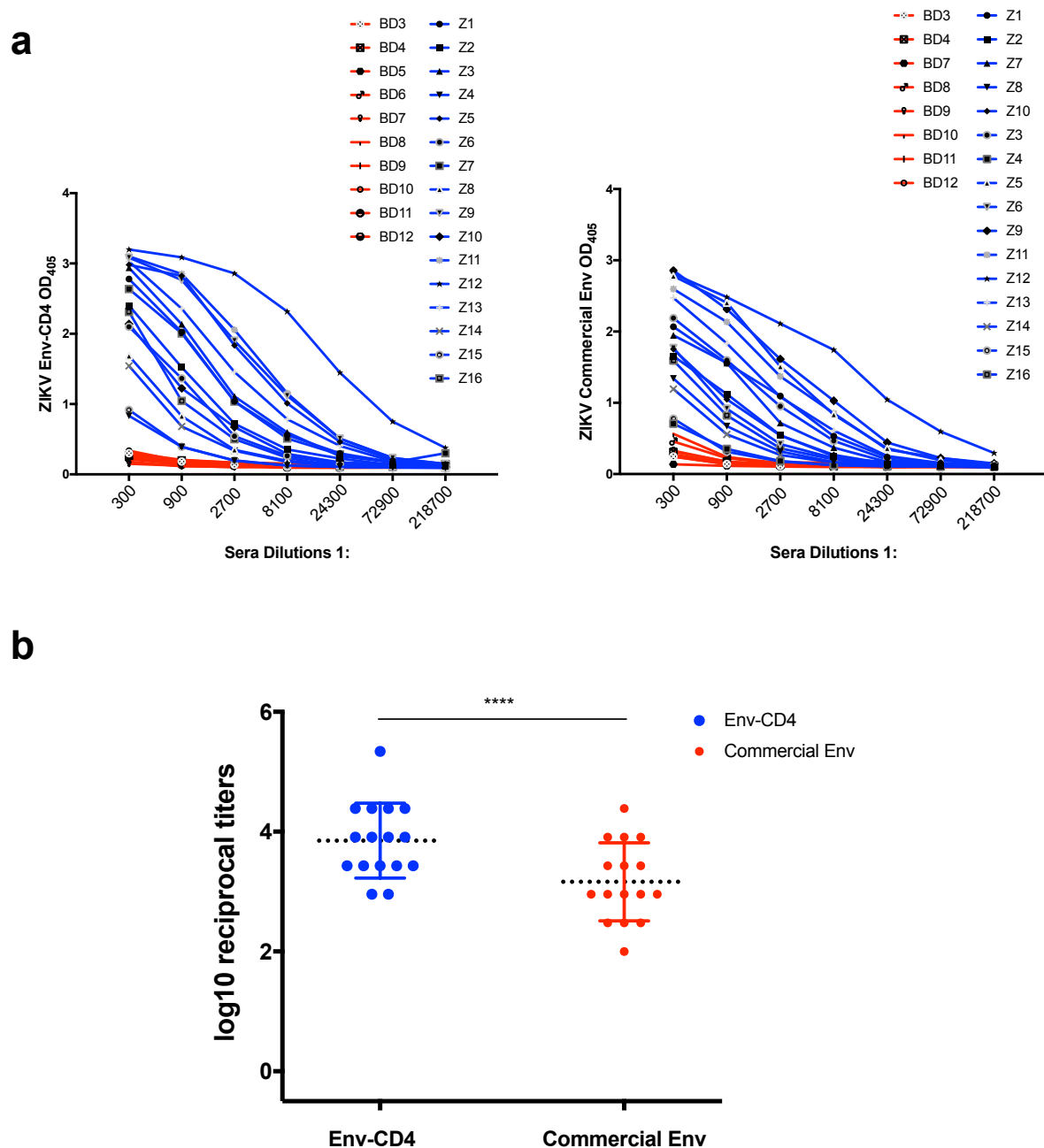
**Figure 1 Production and purification of Zika Envelope proteins**

a). An example of western-blot of Env-CD4 and Env secreted from HEK293T in small-scale expression screening detected with anti-His antibodies. The ectodomain of cell surface receptor sFcεRIα was the positive control and GFP was the negative control. b). Coomassie Brilliant Blue staining (CBB) of the fractions of purified Env-CD4 and Env proteins are shown (reproduced from [1] with permission from Nature Springer).



**Figure 2 Characterisation of ZIKV Env proteins**

Western blot of the purified ZIKV Env-CD4 and Env proteins using a pan-flavivirus antibody, sera from ChAdOx1 ZIKV prME\_ΔTM immunized mice and Zika Env monoclonal antibody. For Env-CD4, the top band corresponds to Env-CD4 protein and the bottom band to Env protein which is formed following cleavage from Env-CD4. An unrelated malaria protein (cCSP) was used as a negative control for the ZIKV-Env specific antibodies. Negative control is cell-free media. (reproduced from [1] with permission from Nature Springer)



**Figure 3 ELISA assays to assess the reactivity of human sera from patients in a ZIKV endemic region of Mexico.** Sera from Zika patients from Mexico (blue) and sera obtained from healthy donors (red). a). The graph of ZIKV Env-CD4 and commercial Env OD405 against sera dilutions. b). The graph shows the endpoint reciprocal titers for ZIKV patients using Env-CD4 and commercial Env proteins. P value ( $<0.0001$ ) was determined by pairwise t-tests. (reproduced from [1] with permission from Nature Springer)