

Original Research Article

<https://doi.org/10.20546/ijcmas.2018.712.154>

## Cloning and Over Expression of CVS Rabies Virus Glycoprotein Gene in *Pichia pastoris* by Multimerization

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### ABSTRACT

#### Keywords

Glycoprotein gene;  
*Pichia pastoris*;  
CVS Rabies Virus,  
SDS-PAGE: RT-  
PCR

#### Article Info

Accepted:  
12 November 2018  
Available Online:  
10 December 2018

Rabies is an endemic, fatal zoonotic disease which is transmitted by the bite of an infected animal. The glycoprotein of rabies virus is most antigenic and immunogenic determinant present in rabies virion and can serve as effective target for development of vaccine. The CVS rabies glycoprotein gene (1575 bp) was cloned into cloning vector pTZ57R/T and confirmed by PCR and Sequencing, further the glycoprotein gene was sub cloned into yeast integrative vector pPICZαA for expression and over expressed by using pPIC9K vector. Linearised recombinant vector containing glycoprotein gene was introduced into *Pichia pastoris* (GS115 and X-33) by electroporation and the recombinant yeast clones were identified by PCR analysis. The recombinant protein expressed in yeast was confirmed by SDS-PAGE, ELISA, western blot and over expression was confirmed RT-PCR. These results allow us to conclude that *P. pastoris* is a convenient system for expression as well as for over expression of CVS rabies glycoprotein through multimerization.

### Introduction

Rabies is a widespread disease and constantly producing victims in human beings as well as in domestic and wild animals in many countries. Presently, however, human rabies cases are reported mostly from developing countries (totally 30,000 persons or more are estimated to die of rabies every year (Meslin *et al.*, 1994). The rabies is unique for its peculiar disease symptoms (e.g. fear of water, painful spasms of the throat triggered by swallowing, etc.) and still is a fearful disease of high mortality rate (almost 100 per cent). Rabies is not only a public health problem in

epidemic areas, but also causes serious economical loss due to outbreaks in cattle of livestock farming in Latin American countries (Arellano-Sota, 1988; Escobar-Cifentes, 1998).

Rabies virus belongs family, Rhabdoviridae, order, Mononegavirales and genus Lyssa virus. The primary structure of the 3' half of rabies viral genome, present in order, the genes encoding for (+) leader RNA, the nucleoprotein 'N', the phosphoprotein 'M1', the matrix protein 'M2', the glycoprotein 'G' and the beginning of the larger protein 'L' which is an essential component of the rabies

virus RNA dependent RNA polymerase (Kawai, 1977). The native G protein is 524 amino acids long, consisting of 19-residue long signal peptide at N-terminus (Anilionis *et al.*, 1981), a 22 amino acid transmembrane domain and a 44 amino acid cytoplasmic domain (Wunner *et al.*, 1983). Glycosylation of the G protein is required for immune protection by the rabies vaccines (Foley *et al.*, 2000).

Rabies vaccines currently used in the developed countries are purified killed virus, which are produced from the infected tissue culture materials. Human diploid cell rabies vaccines have been given to more than 1.5 million people worldwide as of 2006. Newer and less expensive purified chick embryo cell vaccine and purified Vero cell rabies vaccine are now available. The purified Vero cell rabies vaccine uses the attenuated Wistar strain of the rabies virus, and uses the Vero cell line as its host. To control rabies in the dogs, oral immunization is being carried out with live attenuated vaccines, modified live virus rabies vaccine, recombinant rabies vaccine in baits. These vaccines though used extensively, suffer from certain flaws such as requirement of cold chain, administration within 24 h. of infection and their high cost. The recombinant rabies virus glycoprotein (rRVGP) has been produced in several expression systems: [transgenic plants (Ashraf *et al.*, 2005), yeast cells (Sakamoto *et al.*, 1999), mammalian cells (Kankanamge *et al.*, 2003), and insect Sf9 cells infected with Baculovirus (Drings *et al.*, 1999)]. Yeast is one of the eukaryotic microbes that was used for producing viral antigens because of their ability to produce the glycosylated form of the antigens. Within the past 15 years *Pichia pastoris* has successfully entered the scene and is now the second most used host for recombinant gene expression. Based on searches of the PubMed citation database, the use of *P. pastoris* as an expression host has

increased from 4 per cent to 17 per cent of the total recombinant genes reported from 1995 to 2009 (Sorensen, 2010),

Optimization of protein expression is often, but not always, includes the isolation of multicopy expression strains. A strain that contains multiple integrated copies of an expression cassette can sometimes yield more heterologous protein than single-copy strains (Thill *et al.*, 1990; Clare *et al.*, 1991). Strains that contain multiple integrated copies of an expression cassette often produce larger amounts of foreign protein than do single-copy strains (Clare *et al.*, 1991). Therefore, after confirming that a single-copy *P. pastoris* strain produces significant amounts of the correct-sized, biologically active protein, it is advisable to construct and examine protein expression by “multicopy strains” (Li *et al.*, 2007). There are three different approaches that can be used reliably to generate multicopy expression strains of *P. pastoris*. The first approach involves constructing a vector with multiple head-to-tail copies of an expression cassette (Brierley *et al.*, 1998). The second approach entails the use of an expression vector that contains both the *P. pastoris* HIS4 and the bacterial kanamycin resistance gene kanR, which also confers resistance to the related eukaryotic antibiotic G418 (Scorer *et al.*, 1994). The third approach involves the use of a vector carrying the bacterial *Sh ble* gene, which confers resistance to the antibiotic zeocin (Higgins *et al.*, 1998). A detailed description of these methods can be found in Higgins and Cregg (Higgins and Cregg, 1998). Multicopy expression strains of all three types have proven to be stable under the selective pressure of production in fermentor cultures (Higgins *et al.*, 1998).

Mansur *et al.*, (2005) followed two consecutive approaches to increase the number of integrated cassettes: the head-to-tail expression cassette multimerization procedure

and re-transformation with a dominant selection marker. This increased expression from 19 to 250 mg /l when about 11 copies have been integrated. Keeping this in view, the present study was taken to produce and over express the CVS rabies virus Glycoprotein in *Pichia pastoris*.

## **Materials and Methods**

### **Isolation of CVS rabies glycoprotein gene**

Total RNA was isolated from cell line BHK-21 infected with CVS rabies virus (obtained from NIMHANS) using RNeasy mini kit (Qiagen company, GmbH, Germany) and gene specific cDNA by RT-PCR using Omniscript Reverse transcriptase (Qiagen company, GmbH, Germany). Total RNA isolation and Reverse transcriptase PCR reaction were done as per the manufacturer's instructions. RNA so isolated was quantified using a spectrophotometer at 260nm.

The RNA samples were diluted to 1µg/µl concentration using DEPC treated water. 1 µg total RNA was used for getting a gene specific cDNA. The procedure does not allow sample cross contamination and cDNA is utilized as a template for the PCR step, For amplifying full length gene of glycoprotein using specific primers.

### **Cloning of PCR product**

The purified PCR amplicon of CVS glycoprotein was ligated to pTZ257R/T cloning vector (2868 bp), as described in InsT/A clone™ PCR product cloning kit (K1214) of MBI, Fermentas, USA. For ligation, an optimal molar ratio of 1:3 vector: insert was calculated. The ligation mixture along with linerised vector and amplicon DNA were mixed in 0.5 mL micro-centrifuge tubes and incubated at 16°C for 16 h. for ligation.

### **Preparation of competent cells and Transformation of *E. coli* DH5a**

The competent cells of *E. coli* DH5 were prepared by following the protocol mentioned by Sambrook and Russell (2001) with minor modifications. About 100ul of freshly prepared competent cells were taken in a chilled centrifuge tube and 10:1 of ligation mixture was added and mixed gently. The mixture was chilled in ice for 45 min and heat shock was given by shifting the chilled mixture to preheated 42°C water bath for exactly 2min. It was immediately transferred to ice to chill for 5 min. The culture was pre incubated and spread on the plates having Luria agar with Amp50, X-gal, IPTG and incubated overnight at 37°C. The recombinant clones were identified by blue/white assay.

### **Confirmation of clones by PCR and by sequencing**

The Confirmation of the presence of cloned insert was done by PCR amplification of recombinant vectors with respective primers. The total DNA and cloning vector were used as positive and negative controls in the process. The full length 1.575 kb of CVS glycoprotein gene amplicon cloned in pTZ257R/T was sequenced using M13 primers walking technique at Chromas Private Ltd., Bangalore. The sequences were subjected to analysis using BLAST algorithm available at <http://www.ncbi.nlm.nih.gov>.

### **Sub cloning of the cloned CVS Glycoprotein gene into a yeast integrative vector**

For analysing the expression of the cloned CVS glycoprotein gene, yeast integrative vector pPICZαA was used. The amplified CVS glycoprotein gene using gene specific primer having enzyme sites from cDNA were digested with *EcoR* I and *XbaI* restriction

enzyme to create cohesive ends. Accordingly expression vector pPICZ $\alpha$ A was also digested with the same restriction enzymes to have compatible cohesive ends. The ligation reaction was carried out with an optimal molar ratio of 1:3 vector : insert. The components of the ligation mixture was mixed into a 0.5 mL microcentrifuge tube and incubated at 16°C for 16 h. The competent cells of *E. coli* were identified and the plasmid was isolated and confirmed as described earlier using gene specific PCR. Further confirmation was done by complete restriction of clones using *EcoR* I and *Xba*I restriction enzymes.

### **Transformation of yeast cells (*Pichia pastoris*, X-33) with recombinant plasmid DNA**

YPD broth (5mL) in a 50mL conical flask was inoculated with single colony from the YPD agar plates and incubated at 30°C for 16h. With shaking (250-280 rpm). This over night culture (0.5mL) was used for inoculating 500mL fresh YPD broth in a 2L flask and allowed to grow at 30°C to reach an OD600 of 1.3-1.5 (overnight incubation). The cells were pelleted at 4000 x g for 5min at 4°C and suspended in 500mL ice-cold sterile distilled water. The cells were pelleted at 4°C as above and suspended in 250mL sterile ice-cold distilled water. The cells were again pelleted at 4°C as above and suspended in 20mL sterile ice cold 1M sorbitol. The cells were again pelleted as above and resuspended finally in sterile ice-cold 1M sorbitol to a final volume of 1.5mL and aliquoted as 80 $\mu$ l in sterile 1.5mL micro centrifuge tubes and used for transferring the *sac*I digested recombinant pPICZ $\alpha$ A having the glycoprotein gene by electroporation. 100-200 $\mu$ l of the electroporated cells were plated on YPDS agar plates having 400-1000 $\mu$ g/mL concentrated Zeocin and incubated at 30°C until the colonies appeared on the plates. Each

transformants colony was grown in 10mL of YPDS broth at 30°C till the A600 of the culture reached 0.6-1.0 (16-18h.s). The genomic DNA was isolated from yeast transformants by HiPurA kit and confirmed by PCR using specific PCR.

### **Induction of protein expression in recombinant *Pichia* clones and characterization**

Induction of protein expression was done by standard procedure (Cregg *et al.*, 1993) with modification. Buffered Glycerol- complex Medium (BMGY), in 250 mL flasks was inoculated with single PCR positive clone of each serotype and with insert using vector transformant of *Pichia* as a control. The flasks were incubated at 28-30°C in a shaking incubator (250-300rpm) to reach an A600 of 2-6 (~16-18 h). The cells were harvested by centrifuging at 4000xg for 5min at room temperature and suspended in Buffered Methanol-complex Medium (BMMY) to get an A600 of 1.0 (about 200 mL medium) in 1L flask to induce expression. The flasks were incubated at 30°C to continue the growth and methanol was added to a final concentration of 0.5 per cent at every 24h. to sustain the induction. Aliquots of 1 mL each were collected in 1.5 mL Eppendorf tubes at various time points i.e., 0, 12, 24, 36, 48 60, 72, 84, 96,108 and 120 hours and centrifuged at 15000xg in a table top microfuge for 5 min at room temperature. The supernatants containing expressed proteins were collected and the polyhistidine-containing recombinant protein was purified from the secreted media by Affinity column using His tag from the fusion protein.

Proteins expressed by *Pichia* clones carrying rabies virus glycoprotein genes were detected by, RT-PCR, qPCR, Sodium Dodecyl Sulfate - Poly Acrylamide Gel Electrophoresis (SDS-PAGE) using discontinuous buffer system as

followed in Western blot, ELISA as per standard procedures (Sambrook *et al.*, 1989;). Clone without insert was used as a negative control.

### **Multimerization of cvs glycoprotein gene in *Pichia pastoris* (GS115) by sub-cloning into a pPIC9k vector**

The amplified Glycoprotein gene from Recombinant pTZ57R/T having Glycoprotein gene were cut with EcoRI and NotI restriction enzymes. Accordingly expression vector pPIC9K was also digested with the same restriction enzymes to have compatible cohesive ends. Ligation was done by following standard protocol and transformation of ligated product into DH5 $\alpha$  competent cells by using earlier mentioned protocol. Recombinant clones were identified and the plasmid was isolated and confirmed by using gene specific PCR. Further confirmation was done by complete restriction of clones using *EcoR I* and *NotI* restriction enzymes.

### **Transformation of yeast cells (*Pichia pastoris*, GS115) with recombinant plasmid DNA**

The yeast cells, *Pichia pastoris*, GS115 were transformed with linearized plasmid with or without insert as per the standard protocols (Hennen *et al.*, 1978; Cregg, *et al.*, 1985). The recombinant plasmid DNA (pPIC9K- rabies virus glycoprotein gene) isolated from 25 mL over night culture and the linearized plasmid DNAs were used for transformation. In order to get Geneticin resistant cells, the recombinant plasmid DNAs (~20 $\mu$ g) in 100 $\mu$ l was digested with Sac I enzyme. After digestion at 37°C for 6 h., the linearized DNA was purified by Fermentas EneJET PCR Purification kit. pPIC9K DNA, 80  $\mu$ l (~20 $\mu$ g) linearized with Sac I enzyme was used as control.

### **Transformation by electroporation and in vivo screening of multiple inserts**

Freshly prepared competent cells (80  $\mu$ l) were mixed with ~12 $\mu$ g of linearized plasmid, transferred to an ice-cold electroporation cuvette (0.2 cm) the cells were incubated for 5 min. on ice and subjected to electric field as described by the manufacturer of the electroporator, Gene Pulser. The charging voltage was 1500V with a capacitance of 25 $\mu$ F at 2000 resistance which gives field strength of ~7500V fern. Immediately after giving the electric pulse (within 8 seconds), 1 mL of ice-cold 1M sorbitol was added to the cuvette and the contents were transferred to a sterile microfuge tube. From this about 100-200 $\mu$ l of the cells were plated on YPDS agar plates having 0.25mg-4mg/mL concentrated Geneticin and incubated at 30°C until the colonies appeared on the plates.

1. Pipetted 1 to 2 mL sterile water over the 0.25mg/mL Geneticin resistant transformants on each plate.
2. Resuspended the 0.25mg/mL Geneticin resistant transformants into the water by using a sterile spreader and running it across the top of the agar.
3. Transferred the cell suspension into a sterile, 50 mL conical centrifuge tube and vortex briefly (5 to 10 sec).
4. Determined cell density using a spectrophotometer and Plated 10<sup>5</sup> cells on YPD plates containing Geneticin® at a final concentration of 0.25, 0.5, 0.75, 1.0, 1.5, 1.75, 2.0, 3.0, and 4.0mg/mL.
5. Incubated the plates at 30°C and checked daily. Geneticin-resistant colonies were taken 2 to 5 days to appear while cells plated on YPD will take 2-3 days.

There were only a few Geneticin resistant colonies, and they may be of different sizes, but the colony morphology was same. Picked all Geneticin resistant colonies and purified by streaking for single colonies. Glycoprotein gene insertion was analyzed by PCR by isolating the genomic

DNA, and done the Real time PCR for colonies grown on higher concentration of Geneticin to know the copy number by comparing with 0.25mg/mL Geneticin resistant colonies. The cybergreen from Biorad was used for Real time PCR and procedure followed as for the instruction of the company manual.

## Results and Discussion

### PCR amplification of CVS rabies glycoprotein gene from cDNA and sequencing

Total RNA was isolated from BHK-21 cell line infected with CVS rabies virus and constructed the gene specific cDNA Using glycoprotein forward primer. Rabies Glycoprotein gene was amplified from gene specific cDNA using Glycoprotein specific primers designed using reported Rabies nucleotide sequence from the database. The amplicons so obtained were separated on 1.0 per cent agarose gel is presented in Plate 1. From the gel, it is clear that an amplicon of 1575 bp was obtained from amplification of Rabies Glycoprotein gene. This amplicon was cloned into cloning vector pTZ57R/T. The recombinant molecules were transferred into *E. coli* DH5 $\alpha$  using 5  $\mu$ l of ligation mixture. The transformed cells were picked up and streaked on Luria agar containing ampicillin (100mg/mL), X-gal and isopropyl -D-thiogalactosidase (IPTG). The clones containing recombinant molecules were selected based on blue-white colonies. Plasmids were isolated from white colonies

contained Rabies Glycoprotein gene and the clones were confirmed through PCR amplification by using specific primers (Plate 2). The confirmed recombinant vectors with Rabies Glycoprotein were named pNRCRG1506.

The construct pNRCRG1506 was sequenced completely using M13 primers and by employing primer walking technique. Figure 1 represent the map of pNRCRG1506, the complete sequence of nucleotide and amino acid sequences of Rabies Glycoprotein gene is presented in Figure 2 and 4. Sequenced Glycoprotein gene was analyzed for the presence frequently used restriction sites and represent in Figure 3.

### Cloning of rabies glycoprotein gene into yeast Integrative vector and transformation into *Pichia pastoris* (X-33)

The Glycoprotein gene was amplified from pNRCRG1506 using Glycoprotein specific forward and reverse primers having *EcoRI* and *XbaI* sites respectively to clone into pPICZ $\alpha$ A directionally by restricting the vector with same enzyme with that of the gene. The pPICZ $\alpha$ A containing Glycoprotein was transformed into DH5 $\alpha$  and confirmed by PCR with pPICZ $\alpha$ A vector DNA as negative control. The recombinant plasmid DNA gave amplicon of 1575bp, and named this construct as pNRCRG1510 (Fig. 5, 6 & 7) while in pPICZ $\alpha$ A vector; there was no amplification (Plate 3). The positive clones were also confirmed by restriction digestion with *EcoRI* and *XbaI* (Plate 4).

The recombinant clone pNRCRG1510 was linearized with *SacI* enzyme to express the full length of Glycoprotein in *Pichia pastoris* and confirmed by running on one per cent gel (plate 5). The linearized pNRCRG1510 was transferred to electrocompetent *Pichia pastoris* by electroporation as described in the material

and methods. Similarly pPIZ $\alpha$ A vector was transferred into *Pichia* and transferred *Pichia pastoris* was plated on YPD having 400-1000 $\mu$ g of Zeocin. The antibiotic resistant *Pichia* colonies were screened for the selection of recombinant *Pichia* for Glycoprotein gene by PCR using Glycoprotein gene specific primers. A single and intense DNA band of size 1575 bp could be seen in case of positive colonies carrying full length glycoprotein gene, which grew on zeocin, indicating the presence of glycoprotein gene insert (Plate.6) in yeast genome.

### **Analysis of *pichia* Expressed glycoprotein by RT-PCR and SDS-PAGE**

Positive and control *Pichia* clones were induced using 0.5per cent methanol for 96 h. and pelleted the cells. The total RNA isolated from both glycoprotein positive *Pichia* clone and controls were quantified using a Biophotometer at 260 nm. Equal amount of RNA was used for construction of cDNA using Reverse Transcriptase subjected to PCR by using specific primers for Glycoprotein gene. The amplicon of 1575 bp was observed in recombinant *Pichia* and it was absent in control (plate 7).

The culture supernatant from shake flask culture of induced Glycoprotein was purified by anti nickel column. The purified protein was analyzed by SDS-PAGE and quantified by Bradford reagent as 100mg/mL. After destaining, the gel was examined under X-ray illumination and documented (Plate. 8). There was a single protein band, which could be seen in the gel. Upon comparison of protein profile of the recombinant clones (lane 2 and 3) with protein ladder, the protein position is around 78 kDa. Since the full length glycoprotein gene insert is 1575bp and signal sequence along with histag the expected size of the protein expressed, should be around 76-78 kDa. Therefore, the band at position 78

kDa may correspond to the cloned gene product. Cells free extract from positive culture supernatant of induced glycoprotein was concentrated by passing through 30kd cut-off centrican column and quantified by Bradford reagent. Rabies Glycoprotein was also precipitated from cells free extract of positive clones by Ammonium sulphate and quantified by Bradford reagent along with standards.

### **ELISA and Western blotting assay for the expressed protein**

The Glycoprotein of CVS rabies virus produced in *Pichia pastoris* was further analyzed by western blot analysis using sera raised against whole virus (Plate.9). The commercial human vaccine was used as a positive control.

The blot showed positive colour reactions with the proteins of both the clones (lane, 2)) and in Commercial vaccine (lane, 1). However the differences in size of the protein between commercial vaccine and purified recombinant Glycoprotein can be attributed to the His tag and  $\alpha$  secretory signal form the yeast integrative vector. While in ammonium precipitated protein, the tag and secretory signal might have been removed during precipitation. The CVS rabies virus Glycoprotein was further confirmed by ELSA.

Western blot positive *Pichia* expressed glycoprotein was further confirmed by coating the 1 $\mu$ g of antigen and antiserum with different dilutions were used as primary antibody as for the protocol described in the material and methods.

At Dilution of 1:500 in clone 8 got maximum OD of 1.533 as compare to clone7 of OD 0.956 and at 1:1000 dilution in clone 8 got OD of 1.008 where as in clone 7 got the OD of 0.512 (Table 1 and Fig. 8).

### **Multimerization of Glycoprotein gene cassette in *Pichia pastoris* (GS115) by cloning into pPIC9K vector**

The Glycoprotein gene was amplified from pNRCRG1506 using Glycoprotein specific primers having sites of *EcoRI* and *NotI* sites in forward and reverse primer respectively to clone into pPIC9K directionally by restricting the vector with same enzyme with that of gene. The vector and insert from pNRCRG1506 was ligated at 1:3 molar ratio, were used. The transformants in *E. coli* DH5 $\alpha$  were picked and streaked on Luria agar plates containing 100 $\mu$ g/mL Ampicilin. The pPIC9K containing Glycoprotein was confirmed by PCR with pPIC9K vector DNA as negative control. The recombinant plasmid DNA gave amplicon of 1575bp, and named this construct as pNRCRG2504 (Fig. 9) while in pPIC9k vector; there was no amplification (Plate10). These clones were also confirmed by restriction digestion with *EcoRI* and *NotI* (Plate 11).

The recombinant pNRCRG2504 vector was linearized with *SacI* enzyme to integrate multicopy Glycoprotein cassette and express the full length of Glycoprotein in *Pichia pastoris* (GS115).

The linearized pNRCRG2504 was transferred to electrocompetent *Pichia pastoris* (GS115) by electroporation as described in the material and methods. Similarly pPIC9K vector was transferred into *Pichia pastoris* (GS115) and Transferred *Pichia pastoris* was plated on YPD having 0.25mg/mL of Genetcin.

### **Screening of recombinant *Pichia* (GS115) clones for multicopy inserts**

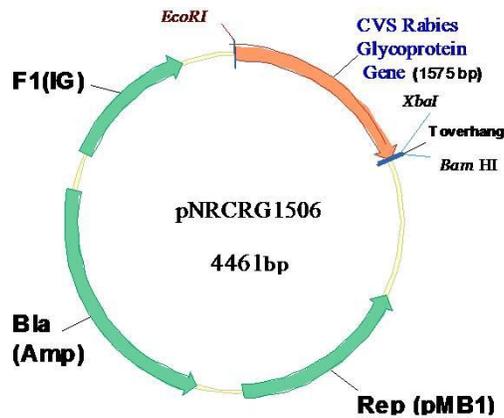
Multicopy Glycoprotein cassette inserted *Pichia* clones were screened by plating the *Pichia* transformants on 0.25mg/mL Geneticin

plates. 0.25mg/mL Geneticin resistant *Pichia* colonies were replated on higher concentrated Geneticin (2mg and 4mg/mL) for the selection of recombinant *Pichia* for Multicopy Glycoprotein gene. Genomic DNAs was isolated from Geneticin resistant colonies transformed by recombinant DNAs and a vector transformant colony were extracted separately as described under material and methods and subjected to PCR amplification with gene specific primers for the presence of the insert. The amplified products were analyzed by agarose gel electrophoresis using standard molecular weight markers (Plate.12). A Single and intense DNA band of size 1575bp could be seen in case of positive colonies carrying full length glycoprotein gene, which grew on Geneticin, indicating the presence of glycoprotein gene insert in yeast genome. While there was no amplification in *Pichia pastoris* transferred with vector without insert.

Real time PCR was carried out using template DNA isolated from 0.25, 2 and 4 mg/mL Geneticin resistant colonies. *Pichia* colony resistant to 4mg/mL Geneticin was crossed the threshold level at 21<sup>st</sup> cycle, 2mg/mL Geneticin resistant colony was crossed at 22<sup>st</sup> cycle and 0.25 mg/mL Geneticin resistant colony was crossed at 24<sup>th</sup> cycle, indicating the 4mg/mL Geneticin resistant colony has got more copy of Glycoprotein insert followed by 2mg/mL Geneticin resistant colony and then with 0.25mg/mL Geneticin resistant colony (Fig. 10). Rabies is currently an incurable disease with a mortality rate of 100per cent, if not immediately treated after the first debilitating nervous symptoms, making it one of the most dangerous zoonotic viral diseases to both humans and animals (NASPHV, 2007; Sudarshan *et al.*, 2007; Zhang *et al.*, 2005; Knobel *et al.*, 2005; Schneider *et al.*, 2007) Yet, efficient diagnostic and safe treatment remains unaffordable in regions where it is most endemic.

**Table1. ELISA absorbance values recorded at 450 nm for CVS Rabies Glycoprotein extracted from Pichia positive clone of seven and eight.**

<b>Serum at different dilution</b>	<b>Glycoprotein from cell free extract of Pichia Clone eight</b>	<b>Glycoprotein from cell free extract of Pichia Clone seven</b>
	<b>OD</b>	<b>OD</b>
1ug/1:500	1.533	0.956
1ug/1:1k	1.008	0.512
1ug/1:2k	0.573	0.289
1ug/1:4k	0.315	0.16
1ug/1:8k	0.186	0.112
Ag Blank	0.066	0.058
Ab Blank	0.099	0.076

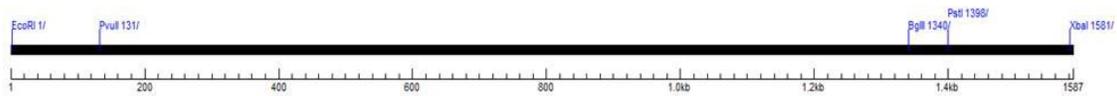


**Fig 1. Construct map of pNRCRG1506 containing full length CVS Rabies Glycoprotein gene in pTZ57R/T**

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GAATTCATGGTTCCTCAGGTTCTTTGTTGTACCCCTTCTGGGTTTTTCGTTGTGTTTC
GGGAAGTTCCTTACACGATACAGACGAACCTGGTCCCTGGAGCCCTATTGACAT
ACACCATCTCAGCTGTCAAATAACCTGGTTGTGGAGGATGAAGGATGTACCAACCTGT
CCGAGTTCTCTACATGAACTCAAAGTGGGATACATCTCAGCCATCAAAGTGAACGG
GTTCACTTGACAGGTGTTGTACAGAGGACAGACCTACACCAACTTTGTTGTTAT
GTCACAACCACATTCAAGAGAAAGCATTTCGCCCCACCCAGACGCATGTAGAGCCG
CGTATAACTGGAAAGATGGCCGGTACCCCGAGATGAAGAGTCCCTACACAATCCATAC
CCCGACTACCACTGGCTTCGAACTGTAAGAACCACCAAGAGTCCCTCATTATCATATCC
CAAAGTGTACAGATTTGGACCCATGACAAATCCCTTCACTCAAGGGTCTCCCTGG
CGGAAAGTGTCTCAGGAATAACGGTGTCTCTACCTACTGCTCAACTAACCATGATTACA
CCATTTGGATGCCCGAGAATCCGAGACCAAGGACACCTTGTGACATTTTACCAATAGC
AGAGGGAAGAGAGCATCAAAGGGAACAAGACTTGCCTGTTGTGGATGAAAGAGG
CCTGTATAAGTCTTAAAGGAGCATGCAGGCTCAAGTTATGTGGAGTCTTGGACTTA
GACTTATGGATGGAACATGGGTGCGGATGCAAACTTAGATGAGACCAATGGTGCC
TCCAGGTGAGTGGTGAATTTGCACGACTTCTCTCAGACGAGATTGAGCATCTCGTTG
TGGAGGAGTTAGTCAAGAAAAGAGAGGAATGTCTGGATGCATTAGAGTCCATCATGAC
CACCAAGTCAGTAAGTTTACAGCTCTCAGTCACTGAGAAACTTGTCCAGGGTTT
GGAAAAGCATATACCATATTCAACAAAACCTTGTGGAGGCTGATGCTCACTACAAGTC
AGTCCGGACCTGGAATGAGATCATCCCTCAAAGGGTGTGAAAGTTAGAGGAAG
GTGCCATCTCATGTGAACGGGGTGTGTTTCAATGGTATAATATTAGGGCTGACGGCC
ATGTCCTAATCCAGAGATGCAATCATCCCTCCTCAGCAACATATGGAGTTGTTGGAAT
CTTCACTATCCCCTGATGACCCCTGGCAGACCTTCTACAGTTTTGAAAGAAAGT
GATGAGGCTGAGGATTTGTTGAAGTTCACCTCCCCGATGTACAAAACAGATCTCAG
GGGTTGACCTGGTCTCCCGAACTGGGGAAAGTATGTATTGATGACTGCAGGGGCCAT
GATTGGCCTGGTGTGATATTTCCCTAATGACATGGTGCAGAAGAGCCAATCGACCCAG
AATCGAAAACAGCGAGTTTTGGAGGGACAGGGAGGAATGTGTCAGTCACTTCCAAA
GCCGAAAAGTCATACCTTCATGGGAATCATATAAGAGTGGAGGTGAGACCAAGTCTC
TCTAGA
    
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**Fig 2. Complete sequence of cloned CVS Rabies Glycoprotein gene along with added base pairs and restriction sites**



**Fig 3. Restriction sites present in Sequenced CVS Rabies Glycoprotein gene**

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MVPQVLLFVPLLGFSLCFGKFPYITIPDELGPWSPIDIHHLSCPNNLVVEDE
GCTNLSEFSYMELKVGYSIAIKVNGFTCTGVVTEAETYTNFVGYVTTTFK
RKHFRTPTDACRAAYNWKMAGDPRYEESLHNPYPDYHWLRTVVRTTKES
LIISPSVTDLDPYDKSLHSRVFPGGKCSGITVSSTYCSTNHDYTIWMPENP
RPRTPCDIFTNSRGKRASKGNKTCGFVDERGLYKSLKGACRLKLCGVLG
LRLMDGTWVAMQTLDETKWCPPGQSVNLHDFLSDEIEHLVVEELVKKR
EECLDALESIMTTKSVSFRRLSHLRKLVPGFGKAYTIFNKTLMEADAHYK
SVRTWNEIIPSKGCLKVRRGRCHPHVNGVFFNGIILGPDGHVLIPEMQSSLL
QQHMELLESAIPLMHPLADPSTVFEEGDEAEDFVEVHLPDVYKQISGVD
LGLPNWGKYVLMTAGAMIGLVLIFSLMTWCRRANRPESKQRSFGGTGR
NVSVTSQSGKVIPSWESYKSGGETRL
    
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**Fig 4. Amino acids sequence of cloned Rabies Glycoprotein gene**

.7% identity in 524 residues overlap; Score: 2763.0; Gap frequency: 0.0%

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Rs-Gly,      1 MVPQVLLFVPLLGFSLCFGKFPYITIPDELGPWSPIDIHHLSCPNLWVEDEGCTNLSEF
352767      1 MVPQVLLFVPLLGFSLCFGKFPYITIPDELGPWSPIDIHHLSCPNLWVEDEGCTNLSEF
*****

Rs-Gly,      61 SYMELKVGYISAIKVNFGFTCTGVVTEAETYTNFVGYVTTTFKRKHFRPTPDACRAAYNWK
352767      61 SYMELKVGYISAIKVNFGFTCTGVVTEAETYTNFVGYVTTTFKRKHFRPTPDACRAAYNWK
*****

Rs-Gly,      121 MAGDPRYEESLHNPYPDYHWRVTRVTTKESLIIISPSVTDLDPYDKSLHSRVFPGGKCSG
352767      121 MAGDPRYEESLHNPYPDYHWRVTRVTTKESLIIISPSVTDLDPYDKSLHSRVFPGGKCSG
*****

Rs-Gly,      181 ITVSSTYCSTNHDYTIWMPENRPRPTPCDIFTNSRGKRASKGNKTCGFVDERGLYKSLKG
352767      181 ITVSSTYCSTNHDYTIWMPEDRPRPTPCNIFTNSRGKRASKGNKTCGFVDERGLYKSLKG
*****

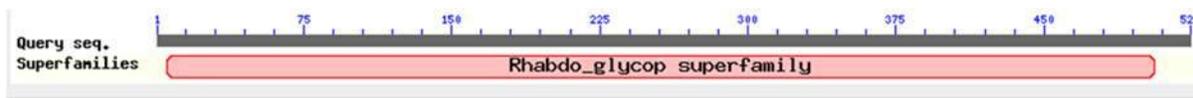
Rs-Gly,      241 ACRLKLCGVLGLRLMDGTWVAMQTLDETKWCPPGQSVNLHDFLSDEIEHLVVEELVKKRE
352767      241 ACRLKLCGVLGLRLMDGTWVAMQTSDETKWCPPDQVLNHLDFHSDEIEHLVVEELVKKRE
*****

Rs-Gly,      301 ECLDALESIMTTKSVSFRRLSHLRKLVPGFGKAYTIFNKTLMEADAHYKSVRTWNEIIPS
352767      301 ECLDALESIMTTKSVSFRRLSHLRKLVPGFGKAYTIFNKTLMEADAHYKSVRTWNEIIPS
*****

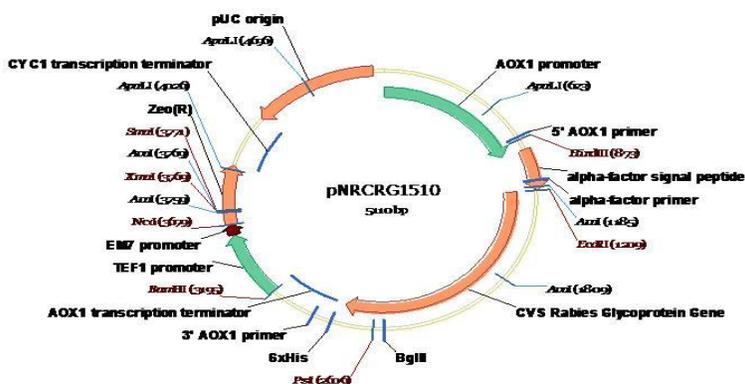
Rs-Gly,      361 KGCLKVGRCHPHVNGVFFNGIILGPDGHVLIPEMQSLLQQHMELLESSAIPLMHPLAD
352767      361 KGCLKVGRCHPHVNGVFFNGIILGPDGHVLIPEMQSLLQQHMELLEKSSVIPLMHPLAD
*****

Rs-Gly,      421 PSTVFEEGDEAEDFVEVHLPDVYKQISGVDLGLPNWGKYVLMTAGAMIGLVLI FSLMTWC
352767      421 PSTVFKEGDEAEDFVEVHLPDVYKRI SGVDLGLPNWGKYVLMTAGAMIGLVLI FSLMTWC
*****

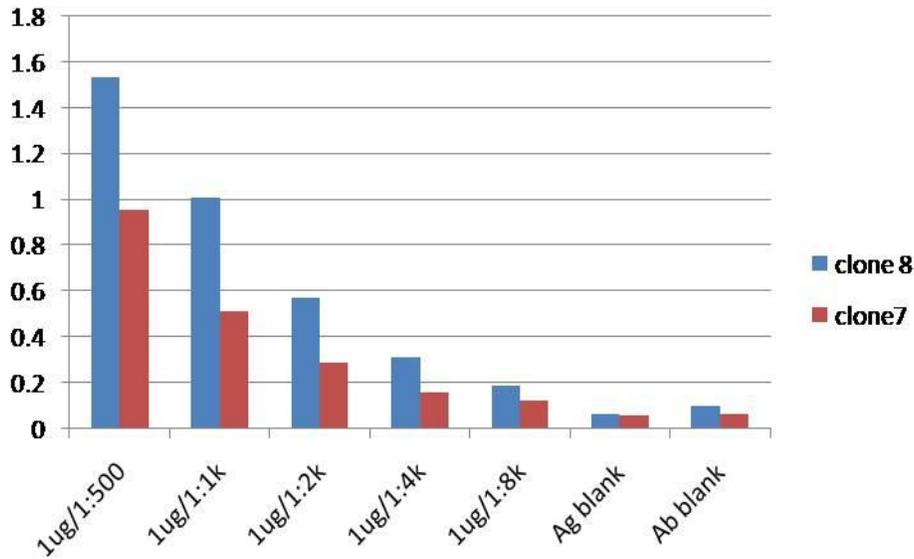
Rs-Gly,      481 RRANRPESKQRSFGGTGRNVSVTSQSGKVI PSWESYKSGGETRL
352767      481 RRANRPESKQRSFGGTGRNVSVTSQSGKVI PSWESYKSGGEIRL
*****
    
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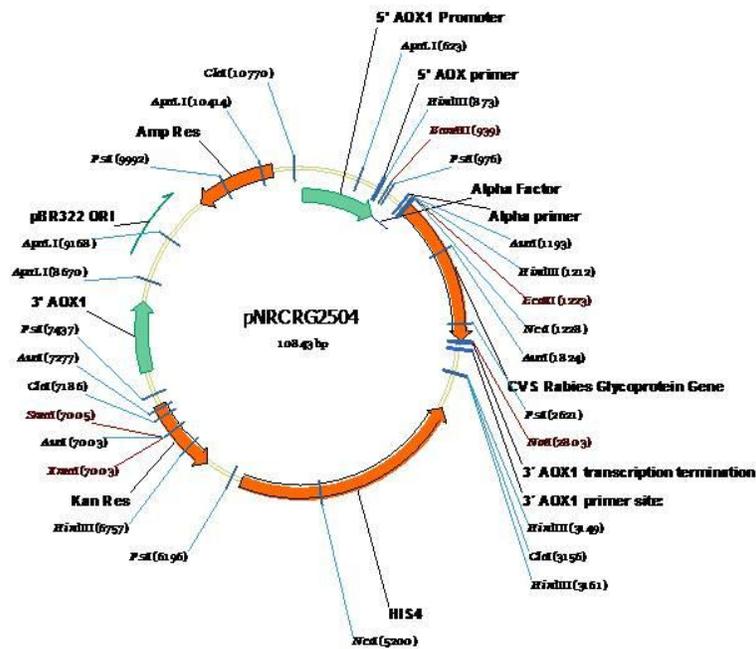
**Fig 6. Conserved protein domain of cloned Rabies Glycoprotein**



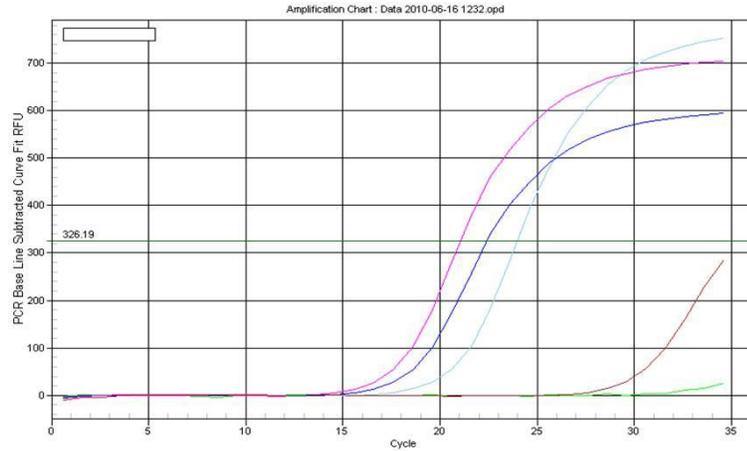
**Fig 7. Construct map of pNRCRG1510 containing full length gene of Rabies Glycoprotein gene**



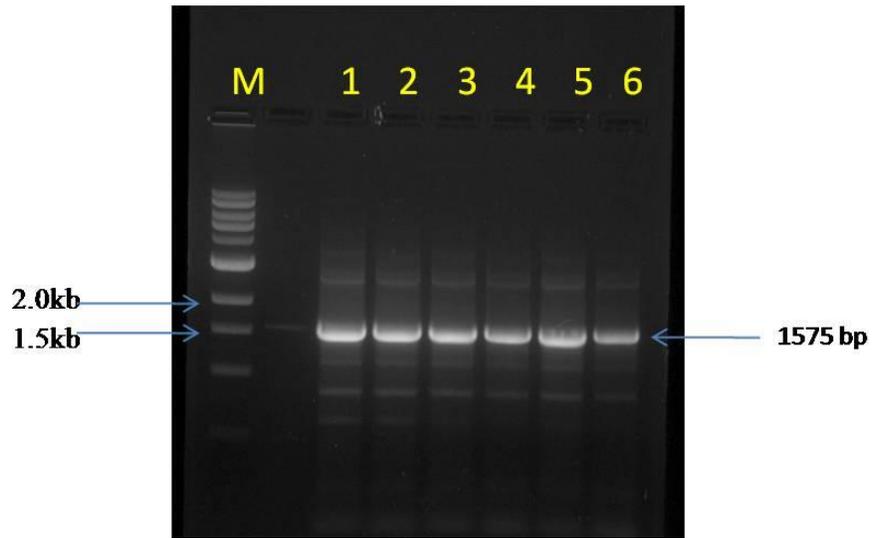
**Fig 8. ELISA of Rabies Glycoprotein using anti sera**  
 X axis – OD  
 Y axis - Serum at different dilution



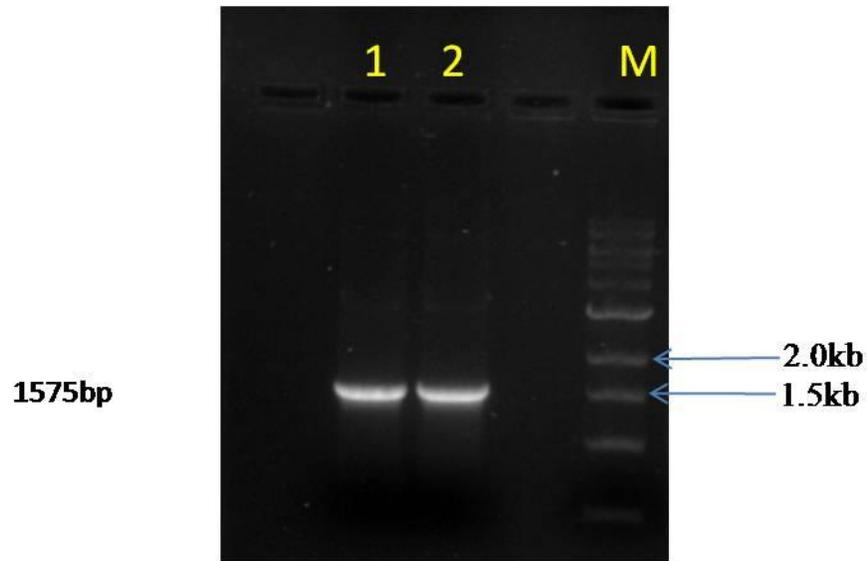
**Fig.9 .Construct map of pNRCRG2504 containing full length gene of Rabies Glycoprotein gene**



**Fig 10. Real time PCR Analysis of *Pichia pastoris* clones containing Rabies Glycoprotein resistant to Geneticin**  
Pink- *Pichia* (GS115) colony resistant to 4mg/ml Geneticin.  
Blue- *Pichia* (GS115) colony resistant to 2mg/ml Geneticin  
Light green- *Pichia* (GS115) colony resistant to 0.25mg/ml Geneticin  
Red – *Pichia* (GS115) host with vector  
Green – water control



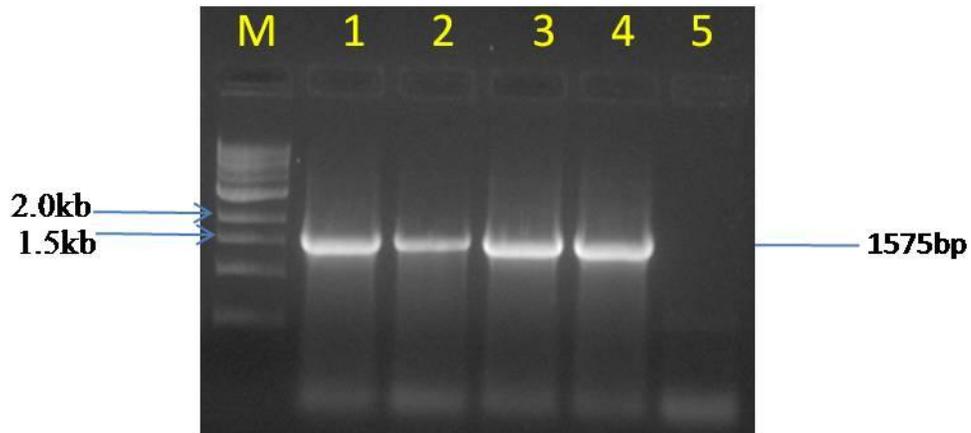
**Plate. 1 PCR amplification of Rabies Glycoprotein gene from cDNA**  
M- 1Kb ladder  
Lane 1,2,3,4,5 and 6- Amplification Glycoprotein with different concentration of cDNA



**Plate .2 PCR confirmation of Construct pNRCRG1506**

Lane M - 1Kb ladder

Lane 1 and 2 – clones

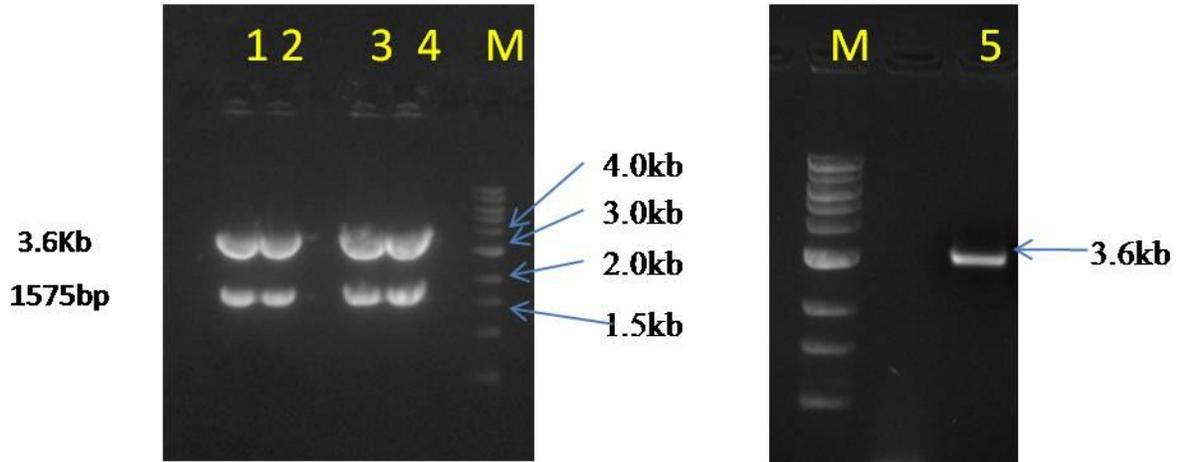


**Plate 3. PCR confirmation of Construct pNRCRG1510**

M- 1 kb ladder

Lane 1,2,3 and 4 – clones

Lane 5- Negative control(vector)

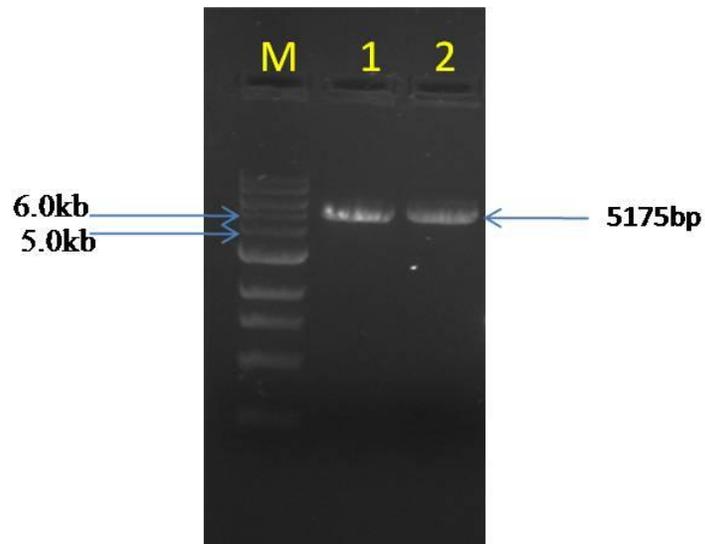


**Plate 4. Restriction digestion of Construct pNRCRG1510**

M- 1 kb ladder

Lane 1,2,3 and 4 – clones

Lane 5 – vector

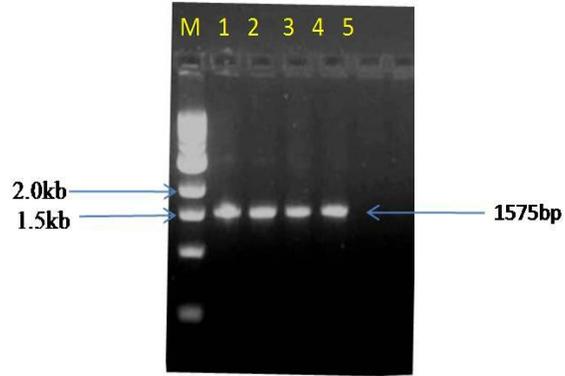


**Plate 5. linearization of Construct**

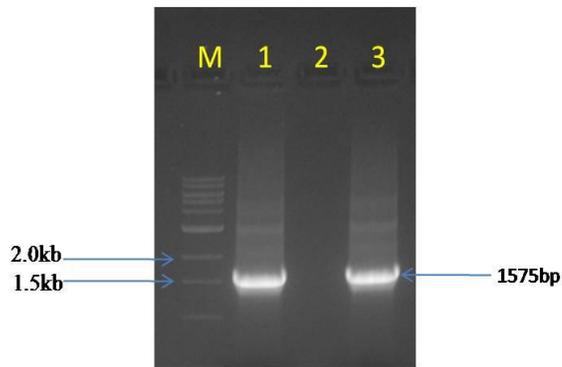
**pNRCRG1510**

M- 1Kb ladder,

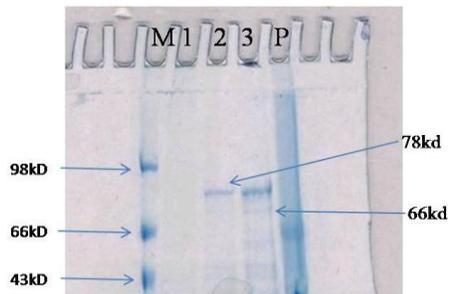
1 and 2 -clones



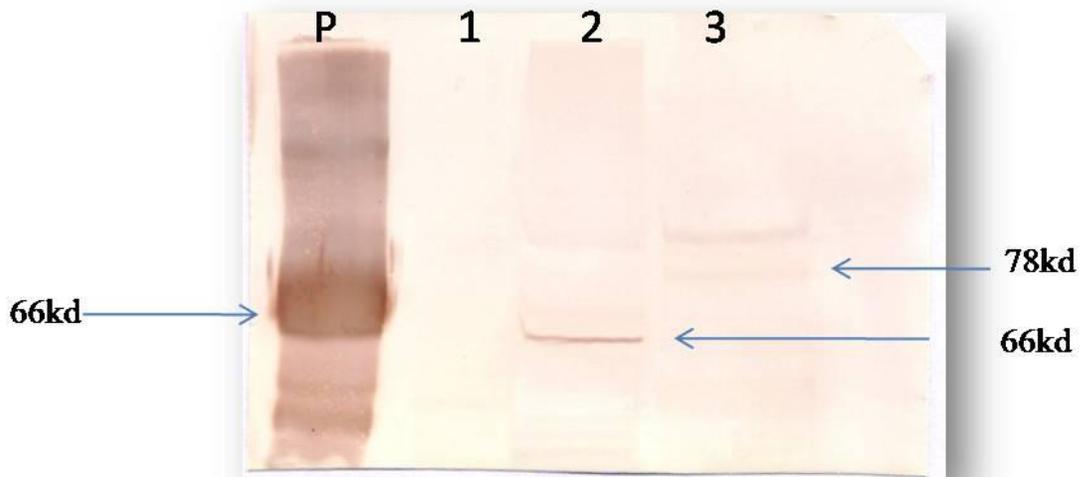
**Plate 6. PCR confirmation of *Pichia* clones for Rabies Glycoprotein**  
1- positive control(pNRCRG1510)  
2,3 and 4- *Pichia* clones  
5- Negative control



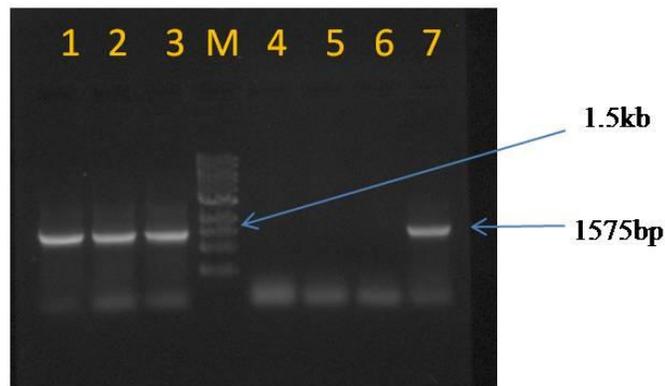
**Plate 7. RT-PCR of Rabies Glycoprotein gene**  
M- 1Kb ladder  
Lane1 and3- clones  
Lane 2- host with vector



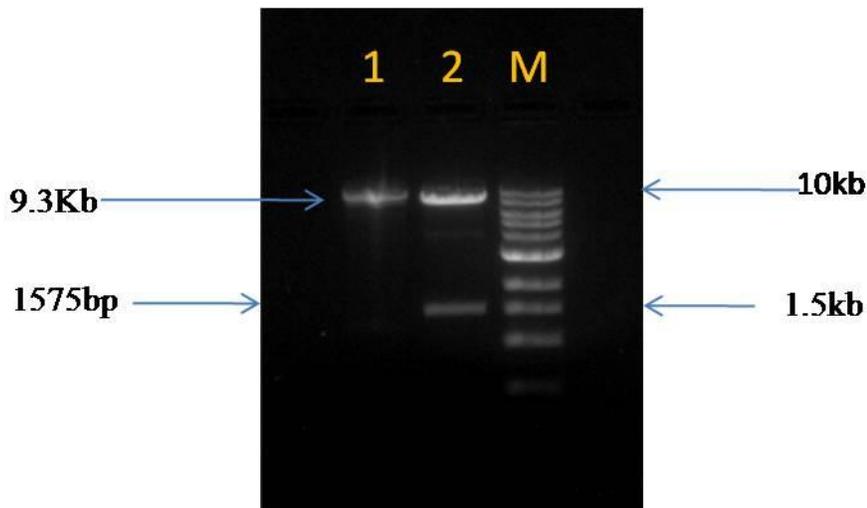
**Plate 8. SDS PAGE analysis of purified Rabies Glycoprotein produced in *Pichia pastoris***  
M- protein marker  
lane1-*Pichia* host with vector  
Lane 2 and 3 – protein from *Pichia* clones  
P- commercial vaccine



**Plate 9. Western blot analysis of purified Rabies Glycoprotein produced in *Pichia pastoris***  
Lane P-commercial vaccine  
Lane 2 –Ammonium sulphate precipitated Rabies Glycoprotein from *Pichia* clones  
Lane3- His tag purified Glycoprotein from *Pichia* clones  
Lane 1 – *Pichia pastoris* host protein.

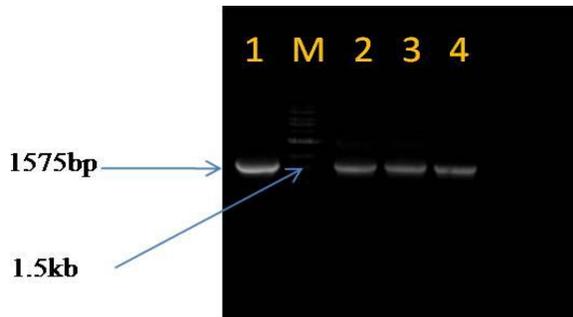


**Plate 10. PCR confirmation of Construct pNRCRG2504**  
Lane 2,3, and 7- clones  
Lane 1- positive control  
Lane 4,5 and 6- Negative control



**Plate 11. Restriction analysis of Construct pNRCRG2504**

Lane 1- digested vector  
Lane 2 –digested pNRCRG2504  
M- 1Kb ladder



**Plate 12. PCR Confirmation of Rabies Glycoprotein gene integration in Pichia pastoris(GS115) clones.**

Lane1 – construct pNRCRG2504  
M-1Kb ladder  
2,3and 4 –clones

The presently available vaccines for rabies include neural tissue derived (NTO), simple vaccine and cell culture derived vaccine. The NTO vaccine is still administered to a large number of people in developing countries, though it causes a lot of serious side effects.

Cell culture vaccines are expensive for routine use in developing countries.

Recombinant gene expression is among the most important techniques used both in molecular and medical research and in

industrial settings. Over the last few decades, the *Pichia pastoris* expression system has been used successfully for production of various recombinant heterologous proteins. The use of the methylotrophic yeast, *Pichia pastoris*, as a cellular host for the expression of recombinant proteins has become increasingly popular in recent times. *P. pastoris* is easier to genetically manipulate and culture than mammalian cells and can be grown to high cell densities. Equally important, *P. pastoris* is also a eukaryote, and thereby provides the potential for producing soluble, correctly folded recombinant proteins that have undergone all the post-translational modifications required for functionality. A further benefit of the *P. pastoris* system is that strong promoters are available to drive the expression of a foreign gene(s) of interest, thus enabling production of large amounts of the target protein(s) with relative technical ease and at a lower cost than most other eukaryotic systems (Daly and Hearn, 2005). This presumption and other success reports on yeast expressed recombinant Polio (Rombaut and Jore, 1997), Bovine herpes virus-1 (Zhu, *et al.*, 1997), Dengue virus (Sugrue *et al.*, 1997) and Hepatitis B Virus Surface Antigen (Liu *et al.*, 2009) vaccines have prompted us to evaluate *Pichia pastoris* as a host to express the Glycoprotein of CVS rabies virus, for using as animal vaccines.

Therefore, the present study was undertaken to clone, express and overexpress the glycoprotein gene coding for transmembrane glycoprotein of CVS rabies virus in *Pichia pastoris* as secretory products and the results of which are discussed here. BHK-21 cells infected with CVS rabies was obtained from the NIMHANS and total RNA was isolated from that cells. Gene specific cDNA was constructed using Forward primer, from this cDNA full length Glycoprotein gene was amplified using gene specific primers. Sakamoto *et al.*, (1999) used the G-cDNA

clone pRNSL, which was isolated from cDNA library that was prepared by reverse-transcribing the mRNA extracted from the hamster HmLu cells infected with rabies virus (sub strain N-HL of the Nishigahara strain was used). The amplified Glycoprotein was cloned into TA cloning vector pTZ57R/T and confirmed the gene by PCR using specific primers. The Rabies Glycoprotein gene was also confirmed by sequencing using M13 primer, and named this recombinant vector as pNRCRG1506. Tomar *et al.*, amplified the RVG gene (1590 bp) using gene specific primers, The amplified Glycoprotein gene was cloned into pTZ57R/T cloning vector by TA cloning. RVG gene was subcloned into pcDNA3.1 (+) expression vector to express rabies virus glycoprotein gene into eukaryotic system and determination of potential T-cell epitopes. The confirmed CVS rabies Glycoprotein gene was amplified from the construct pNRCRG1506 using High fidelity Taq polymerase and gene specific primer having EcoRI and XbaI site. *Pichia* integrative vector pPICZ $\alpha$ A was digested with same enzymes and cloned the digested Glycoprotein amplicon directionally. The ligated vector was transferred to DH5 $\alpha$  cells for maintaining the construct then it was confirmed by PCR and restriction digestion. The recombinant construct of pPICZ $\alpha$ A with Glycoprotein was named as pNRCRG1510.

The pNRCRG1510 was linearized with SacI enzyme and transferred to *Pichia pastoris* (X-33) by electroporation then plated on YPD Zeocin plates. Genomic DNA was isolated from Zeocin resistant *Pichia* clones and confirmed the integration of Glycoprotein gene in *Pichia* genomic DNA by PCR. Su *et al.*, 2009 constructed the recombinant human apolipoprotein E3 in pPICZ $\alpha$ C and subjected to transformation with SacI-linearized version of pPICZ $\alpha$ C-rhApoE3, which favoured its insertion into the *Pichia* (X-33) genome by homologous recombination. Daly and Hearn

(2005) confirmed the Integration of gene of interest in *Pichia pastoris* and the type (insertion or replacement) of events by southern blot analysis hybridized with a probe generated from the AOX1 promoter region.

PCR positive *Pichia pastoris* recombinants for glycoprotein were induced by grown on methanol complex media having 0.5 per cent methanol for 96 hr and checked for the induction of Glycoprotein transcript by isolating total RNA from 0.5 per cent methanol induced *Pichia* cells along with host transferred with vector. The cDNA was constructed from the isolated total RNA. The cDNA was amplified with PCR using Glycoprotein specific primers. Agarose gel electrophoresis indicated that there was an amplicon at 1575bp band in recombinant clones and no band was observed in control. This clearly shown that, the presence of Rabies Glycoprotein transcript in recombinant *Pichia* clones. The same cDNA was used as template for real time PCR and there was a differential concentration of transcript in different Glycoprotein *Pichia* clones. Benmaamar *et al.*, (2009) studied the engineered RNA-based rabies virus glycoprotein (RVGP) expression vector based on the Semliki Forest Virus (SFV) system. A recombinant SFV carrying an RNA coding for RVGP (SFV-RVGP) was constructed and the RVGP expression was evaluated in animal cell cultures. The mRNA coding for RVGP and the RVGP itself were assessed by RT-PCR, qPCR and western blot.

Rabies Glycoprotein purified from Cell free extracts of induced RT-PCR positive clones, by using anti nickel column. Glycoprotein was confirmed by SDS PAGE along with protein ladder, which revealed that the protein size is nearly 76-78Kd. Sakamoto *et al.*, 1999, reported 56 to 66Kd Rabies Glycoprotein produced in yeast cells. The increased in molecular mass weight in our experiment

might be due to alpha secretory signal and his tags which adds nearly 12kd protein. Salah *et al.*, 2009 reported the increased size of the Rabies Glycoprotein is due to posttranslational modification of secreted proteins such as O-glycosylation, lipidation or phosphorylation. In our experiment, the *Pichia* produced G protein is comparable to authentic G protein since it was normally reacted with the confirmation epitope anti-G MAbs. Cells free extract containing rabies glycoprotein was concentrated by passing through 30kd cut-off centrican column and Rabies Glycoprotein was also precipitated from cells free extract of positive clones by Ammonium sulphate and quantified by Bradford reagent along with standards. The concentrated recombinant protein was used for western blotting.

The results of Western blot using a mouse polyclonal immunoglobulin-G, raised against the commercial Rabies vaccine further demonstrated that the protein isolated from cells free extract of induced positive Rabies glycoprotein *Pichia* clones were primarily Rabies Glycoprotein. Western blotting revealed a 66kd band from ammonium sulphate precipitated protein of clone eight and also in commercial vaccine. Whereas protein purified form anti nickel column has shown faint band in higher molecular weight, this may be due to Glycoprotein having alpha secretory signal and histidin tag. In Western-blotting assays, a significant level of RVGP expression was observed in BHK-21 cells infected with SFV-RVGP, at 24 and 48 h. Expression could be monitored by using antibodies against RVGP itself or against the M2-FLAG-tag expressed in fusion with the N-terminal RVGP end. The synthesized RVGP exhibited the expected molecular mass of 80 kDa which corresponds to the recombinant 65 kDa RVGP protein with the additional 15 kDa fusion tags (Benmaamar *et al.*, 2009). Cell free extracts of rabies

Glycoprotein positive *Pichia* clones were further confirmed by ELISA, by coating 1µg Ammonium precipitated rabies glycoprotein of clone seven and eight along with anti sera of rabies commercial vaccine. Clone eight has got the OD of one at 1:1000 anti rabies serum dilution whereas clone seven has got at 1:500 dilution indicating that clone eight is efficient than seven. In a similar study Su *et al.*, (2009) studied the full-length sequence encoding Apolipoprotein E3, gained by RT-PCR, was inserted into the pPICZαC vector and transformed into *P. pastoris* strain X-33, and then confirmed the expressed protein by ELISA using the mouse anti-human ApoE monoclonal antibody.

In vivo multimerization strategy was followed to increase the Rabies Glycoprotein production in *Pichia* by cloning pPIC9K vector, where it provides a Multicopy insertion of Glycoprotein cassette into *Pichia* genome by homologous recombination at AOX site. Insertion of Glycoprotein in *Pichia pastoris* (GS115) was confirmed by PCR. Multicopy inserted colonies were selected by growing on increased concentration of YPD Geneticin plates. Geneticin resistant colonies of 0.25, 2 and 4mg/mL were confirmed by PCR. Real time PCR was performed on these colonies to know the copy number. The *Pichia* colony Grown on 4mg/mL Geneticin has shown higher copy number compared to 2mg/mL and 0.25mg/mL. Abad *et al.*, (2010) also described method used for real-time PCR to quantify the integrated expression cassettes in *Pichia pastoris* to know the copy number.

### Acknowledgement

We gratefully acknowledge the help of the National Institute of Mental Health and Neurosciences, Bangalore, for providing facilities during the initial studies and UAS Banagalore as well as Government of Karnataka for its financial support to carry out the Research.

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**How to cite this article:**

Ningaraju, T.M. and Ramanjini Gowda, P.H. 2018. Cloning and Over Expression of CVS Rabies Virus Glycoprotein Gene in *Pichia pastoris* by Multimerization. *Int.J.Curr.Microbiol.App.Sci*. 7(12): 1233-1255. doi: <https://doi.org/10.20546/ijcmas.2018.712.154>