

Cloning and GST-fused expression in *E. coli* of mouse β -1,4-galactosyltransferase*

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Abstract: β -1,4-galactosyltransferase (β 4Gal-T) (EC 2.4.1.38) plays a multifunctional role in many aspects of normal cell physiology. By now, several dozens of β 4Gal-T genes have been cloned, separated from mouse, chick, bovine, human, etc. This paper presents the cloning and GST-fused expression of mouse β 4Gal-T gene in *Escherichia coli* (*E. coli*). The target gene was cloned by PCR, followed by identification by DNA sequencing and expression in *E.coli* with isopropyl- β -D-thiogalactoside (IPTG) gradient concentrations, products of which were separated on SDS-PAGE showing that the target protein had the same molecular weight as that of mouse β 4Gal-T. The transcriptional product of β 4Gal-T gene was proved by Western hybridization analysis to be due to GST-fusion.

Key words: β -1,4-galactosyltransferase (β 4Gal-T), Cloning, GST-fusion

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INTRODUCTION

β -1,4-galactosyltransferase (β 4Gal-T) (EC 2.4.1.38), as one of the most researched glycosyltransferases in recent years, produces a β -1,4-linked galactosylated glycan by the transference of galactose (Gal) from an uridine diphosphate-galactose (UDP-Gal) to a monosaccharide N-acetylglucosamine (GlcNAc) or a GlcNAc residue located in the non-reducing terminal of glycans of glycoproteins or glycolipids. In addition to GlcNAc, the enzyme can also use other reducing sugars, such as N-acyl substituted glucosamines and N-acetyl-D-mannosamine, as acceptors.

As a Type II membrane protein, β 4Gal-T exists not only in the Golgi complex, but also on the cytomembrane; and functions differently compared to

other intracellular glycotransferases. β 4Gal-T has multiple functions in many aspects of normal cell physiology, and seems to be associated with sperm binding (Macek *et al.*, 1991; Youakim *et al.*, 1994), cell-cell recognition (Hathaway and Shur, 1996; Wassler and Shur, 2000; Wassler *et al.*, 2001; Shi *et al.*, 2001), cell migration on basal lamina (Maillet and Shur, 1993; Eckstein and Shur, 1992; Appeddu and Shur, 1994) and neurite extension (Begovac and Shur, 1990; Hathaway and Shur, 1992; Huang *et al.*, 1995), embryonic maturation (Bayna *et al.*, 1988), rheumatoid arthritis (Yang *et al.*, 1996; Axford, 1999), and cell development (Maillet and Shur, 1994; Asano *et al.*, 1997; Akimoto *et al.*, 1995).

By now, several dozens of β 4Gal-T genes have been cloned from many vertebrates, such as mouse, chick, bovine and human. Since human β 4Gal-T, as a house-keeping gene, was cloned in 1997 for the first time (Almeida *et al.*, 1997), seven members of

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this human gene family have been found (Vadaie et al., 2002; Fan et al., 2002).

This paper presents molecular cloning and high efficiency expression induced by isopropyl- β -D-thiogalactoside (IPTG) gradient concentrations, which yielded the first GST-fused mouse β 4Gal-T gene in *Escherichia coli*.

MATERIAL AND METHODES

Bacteria and plasmids

E. coli strain of BL21 (DE3) and DH5 α were conserved in our lab, and a strain of TOP10 was presented by the Biochemistry and Cell Institute of the Chinese Academy of Sciences (CAS).

pMGT-239/2615 vector, kindly provided by Professor Shur N.D. of Emory University School of Medicine, was exextracted as described (Sambrook et al., 1989). pMD18-T vector was the cloning and sequencing vector and pGEX-4T-2 vector was the plasmid expression vector.

PCR amplification

The target gene was obtained from pMGT-239/2615 vector through Polymer Chained Reaction (PCR), primered with up-primer (5'-ATCGGATCC ATGAGGTTTCGTGAGCAG-3') and down-primer (5'-ACGGAATTCCTATCTCGGTGTCCCGATG-3'). PCR amplification was conducted in total volume of 300 μ l with 30 μ l reaction buffer, 24 μ l of 2.5 mmol/L deoxynucleoside triphosphates (dNTPs), 12 μ l of 10 mmol/L primers mixture and 1.5 μ l of 5 μ g/ μ l Taq polymerase. Cool-starting method was applied to the PCR product, followed by thermo-cycling at 94 $^{\circ}$ C for 30 seconds, 50 $^{\circ}$ C for 50 seconds, and 72 $^{\circ}$ C for 80 seconds for 32 cycles. DNA Sequencing of the PCR product was performed by the Sequencing Ward, College of Life Sciences, Zhejiang University.

Cloning of the target gene

Plasmids were extracted with WizardR Purification Kit by the method of alkaline lysis. Sub-cloning of the target gene was done with pMD18-T, the vector of which contains Glutathione S-transferase (GST) gene that expresses a fusion protein with

potential for further protein detection and separation.

The nucleotide acid sequence of the target gene was compiled and analyzed on the basis of Genbank databases (<http://www.ncbi.nlm.nih.gov>).

Detection of target protein

IPTG gradient concentrations were applied to induce high efficiency expression of the target protein. Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) was performed as described (Sambrook et al., 1989). Hybridization was proceeded at the presence of 0.6 mg/ml 3,3'-diaminobenzidine (DAB), goat anti-GST antibody as the primary antibody, and mouse anti-goat IgG marked by horse radish peroxidase (HRP) as the secondary antibody. The reaction temperature was held at 25 $^{\circ}$ C.

RESULTS

Cloning of the β 4Gal-T gene

On the basis of the conserved constituents of the amino acids sequence of homologues of β 4Gal-T gene and taking into consideration of the guanosine (G) and cytidine (C) percentage of the β 4Gal-T gene, the primers, including the up-primer containing a cleavage site for BamH I, and the down-primer containing a cleavage site for EcoR I, were designed.

PCR amplification carried out with pMGT-239/2615 vector as a template was detected by 1% agarose gel electrophoresis (Fig.1).

Detection of target gene by gel electrophoresis

pMD18-T vectors probably containing an extra sequence of the target gene were extracted from *E. coli* and cleaved by BamH I, followed by detection by 1% agarose gel electrophoresis (Fig.2). The digested products of the positive plasmids were detected by comparison with the 200 bp ladder of the DNA marker (Fig.3), and identified by PCR-gel electrophoresis (Fig.4).

Sequencing of PCR amplification product

The DNA sequence of the PCR amplification

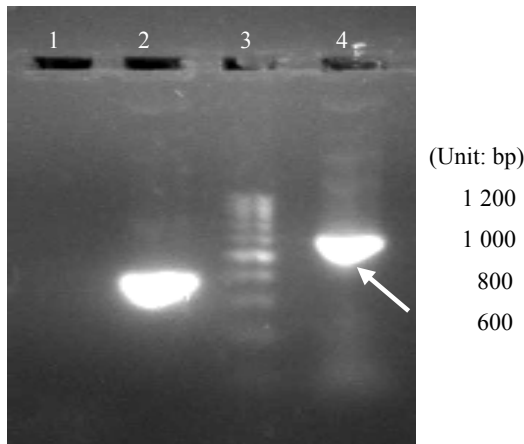


Fig.1 1% agarose gel electrophoresis of PCR amplification product of $\beta 4\text{Gal-T}$

Lane 1 was negative control; Lane 3 shows the 200 bp ladder of the DNA marker, the brightest band of which contained DNA of 1000 bp; Lane 2 shows the product of PCR ($L\beta 4\text{Gal-T}$); and Lane 4 shows the $S\beta 4\text{Gal-T}$ gene (not from PCR)

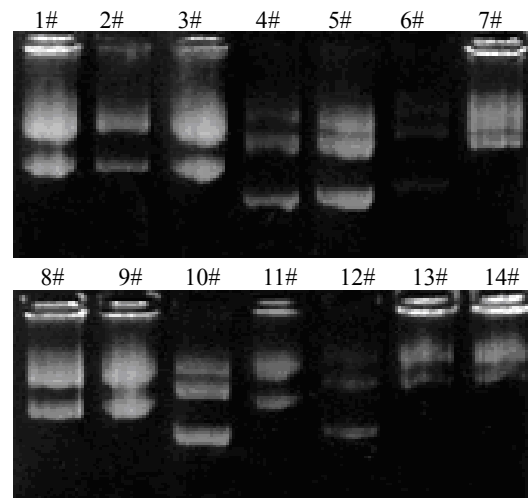


Fig.2 1% agarose gel electrophoresis

Containing an extra gene, the recombinant plasmids (1#, 2#, 3#, 7#, 8#, 9#, 11#) ran slower than the origin plasmids

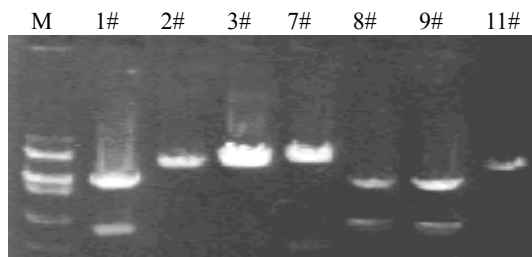


Fig.3 Digest analysis of the recombinant plasmids

Plasmids 1#, 8# and 9# were the candidates, containing DNA band of 1200 bp, while the DNA marker shows the ladder of 6050, 2690, 2160, 1100, 840, 710 and 300 bp

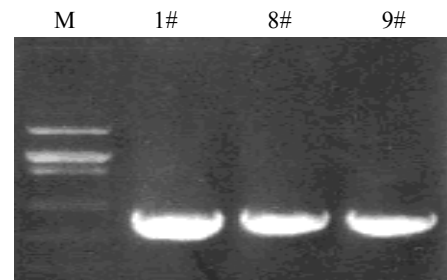


Fig.4 1% agarose gel electrophoresis of PCR amplification products

Plasmids 1#, 8# and 9# showed are positive clones. The DNA marker ladder is the same as that in Fig.3

product is shown in Fig.5. The $\beta 4\text{Gal-T}$ gene of the sequence of 1 216 bp DNA, sub-cloned in the cloning and sequencing vector of pMD18-T, was the same as reported (Nomura *et al.*, 1988), except for guanosine (G) of the +12 site, which was mutated from thymine (T) but did not modify the anticipant sequence of the transcriptional product, because the original codon (CGT) and the mutant codon (CGG) both encoded the amino acid of arginine (Q).

Mouse $\beta 4\text{Gal-T}$ gene encodes two isoforms of mRNA in somatic cells, differentiating in the transcription promoter of the long isoform ($L\beta 4\text{Gal-T}$, 4.1 kb) located upstream the first exon (about 200 bp) while the short one's ($S\beta 4\text{Gal-T}$, 3.9 kb) located downstream. As a result, the long isoform of mouse

$\beta 4\text{Gal-T}$ gene containing 13 more amino acids in the N terminal had transcription pattern different from that of the short isoform. The same happens with the isoforms of bovine and human gene (Masri *et al.*, 1988; Mengle *et al.*, 1991).

Expression pattern of $\beta 4\text{Gal-T}$

GST-fused expression of $\beta 4\text{Gal-T}$ in *E. coli* in the presence of IPTG gradient concentrations, was examined on SDS-PAGE by Western hybridization analysis. The holoproteins on SDS-PAGE showed that the optimum culture condition was 3 hours at 37 °C with 0.4 mmol/L of IPTG (Fig.6). Density Scanning of $\beta 4\text{Gal-T}$ protein on SDS-PAGE (Fig.7) showed that the weight of this protein is 10% that of

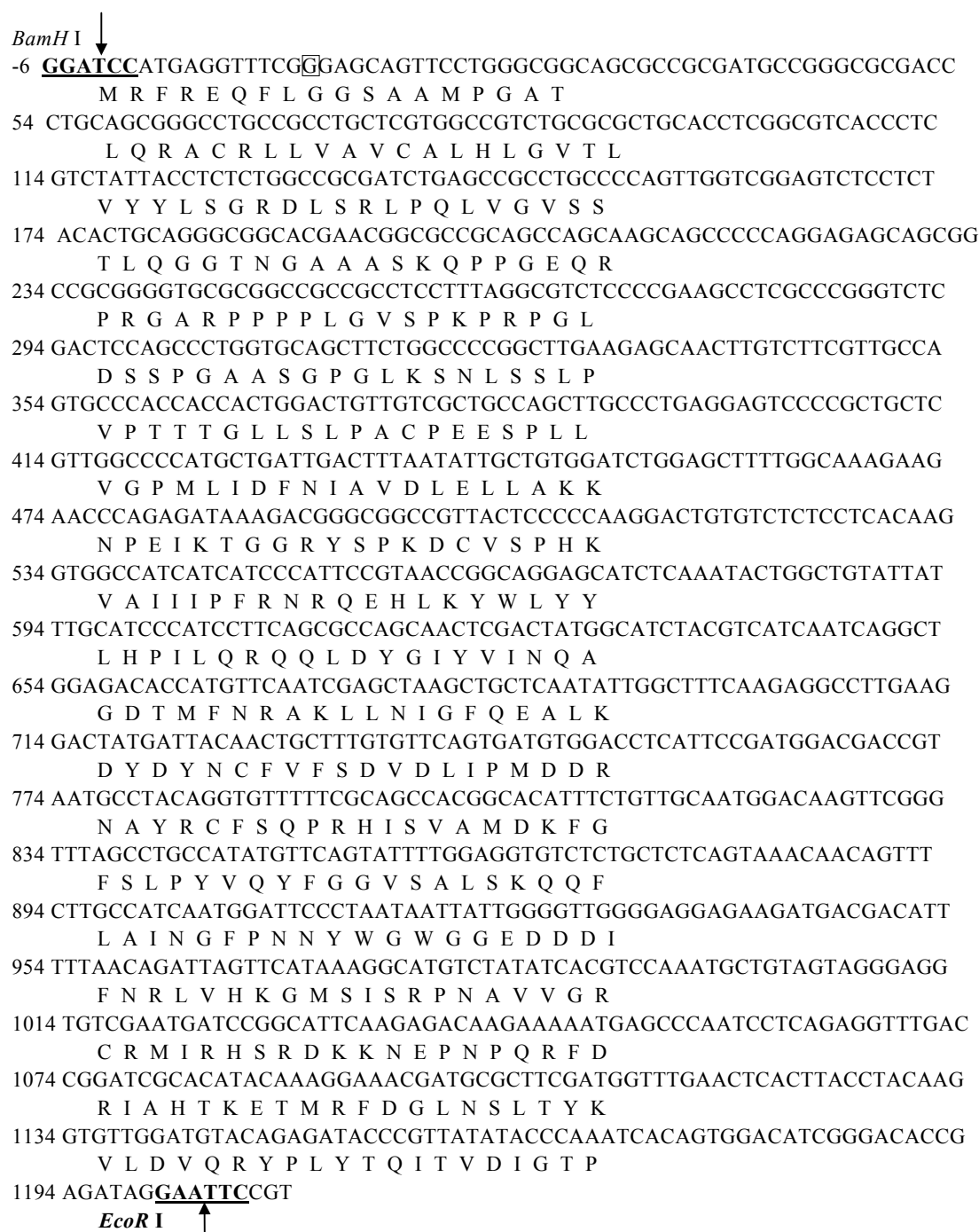


Fig.5 DNA sequence of the PCR product (mouse β 4Gal-T gene) and the amino acid sequence transcript hypothesized from the former DNA sequence

the holoproteins. Western hybridization analysis showed that positive GST-fusion proteins were of the molecular weight of 70 kD (Fig.8). As the mo-

lecular weight of β 4Gal-T is about 45 kD and that of GST is 26 kD, the 70 kD band was confirmed as the β 4Gal-T-GST fusion proteins.

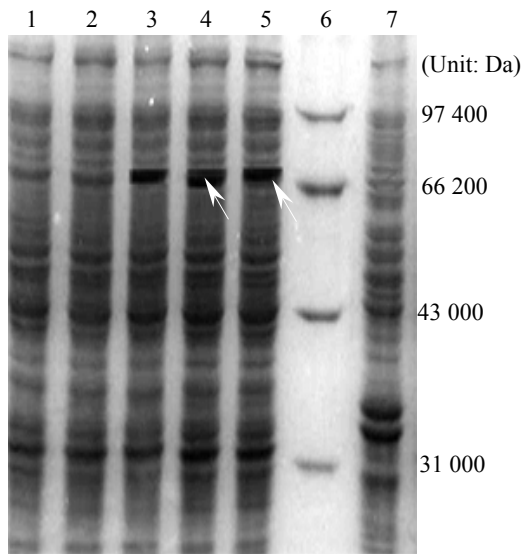


Fig.6 SDS-PAGE analysis for the expression product of $\beta 4$ Gal-T with ladder of IPTG concentration

Lane 6 contains low molecular weight marker, consisting of protein ladder of 97.4, 66.2, 43 and 31 kD. Lane 7 is the negative control of IPTG-induction. Lane 3, 4 and 5, respectively, show a strip on 70 kD, having the same molecular weight as $\beta 4$ Gal-T as reported

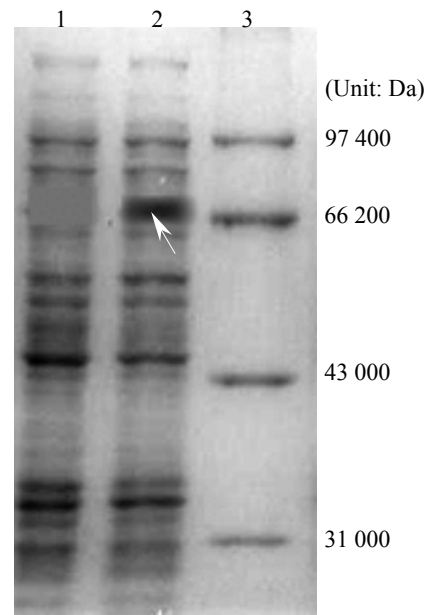


Fig.7 SDS-PAGE analysis for the expression product of $\beta 4$ Gal-T from *E. coli* (BL21)

Lane 3 contains low molecular weight marker, consist of protein ladder of 97.4, 66.2, 43.0 and 31.0 kD. Lane 7 is the negative control of IPTG-induction. Lane 2 shows an apparent strip on about 70 kD in comparison with the marker

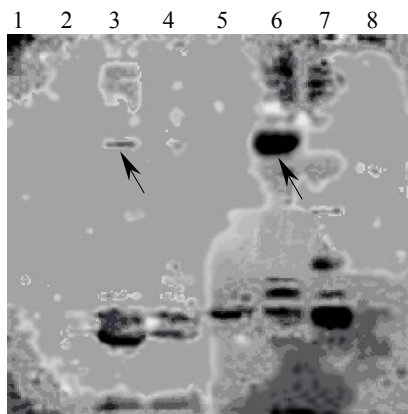


Fig.8 Western blotting of product of pGEX-4T-2-GT

Lane 1 and 2 contain the products of IPTG-free expression. Lane 3 and 6 contain the product of expression with induced by IPTG of 0.4 mmol/L, showing the $\beta 4$ Gal-T-GST fused gene. Lane 4, 5 and 7 are the negative controls. Lane 8 is the low molecular weight marker

DISCUSSION

Hyperstructure of $\beta 4$ Gal-T

The secondary structure of $\beta 4$ Gal-T protein was

measured by DNA Star program (Fig.9).

Masibay *et al.* (1993) found that the secondary structure of $\beta 4$ Gal-T contained 20% β -sheets, 23% α -helices and 2% α - 3_{10} -helices. The catalytic site (Leu131-Ser402) is a $52 \times 50 \times 42$ Å tri-dimensional structure that consists of 11 β -sheets ($\beta 1$ - $\beta 11$), 6 α -helices ($\alpha 1$ - $\alpha 6$) and 2 α - 3_{10} -helices, the central part of which is a twist of 8 β -sheets [Code: 7(81)32465] enclosed by 2 α -helices and 4 α -helices on each side respectively. The central β -sheets ($\beta 2$) structure locating in the N terminus follows an un-affirmatory amino acids terminus and a nude α -helix ($\alpha 1 \beta 1$) hairpin structure. The C terminal domain that begins with a β -sheet ($\beta 8$), consists of α -helix ($\alpha 4$, $\alpha 5$) and a short β -sheet ($\beta 9$). $\beta 9$ trifolds parallelly in reverse upon the central β -sheets are enclosed by an α -helix (3_{10} - $\alpha 6$) and a long β -hairpin ($\beta 10 \beta 11$) respectively. The tapered hyperstructure of $\beta 4$ Gal-T has a big and deep capacious pocket with a diameter of 13 Å (Vrieling *et al.*, 1994); and shows no structural homology with T4 DNA-modified β -glucotransferase (Holm and Sander, 1983).

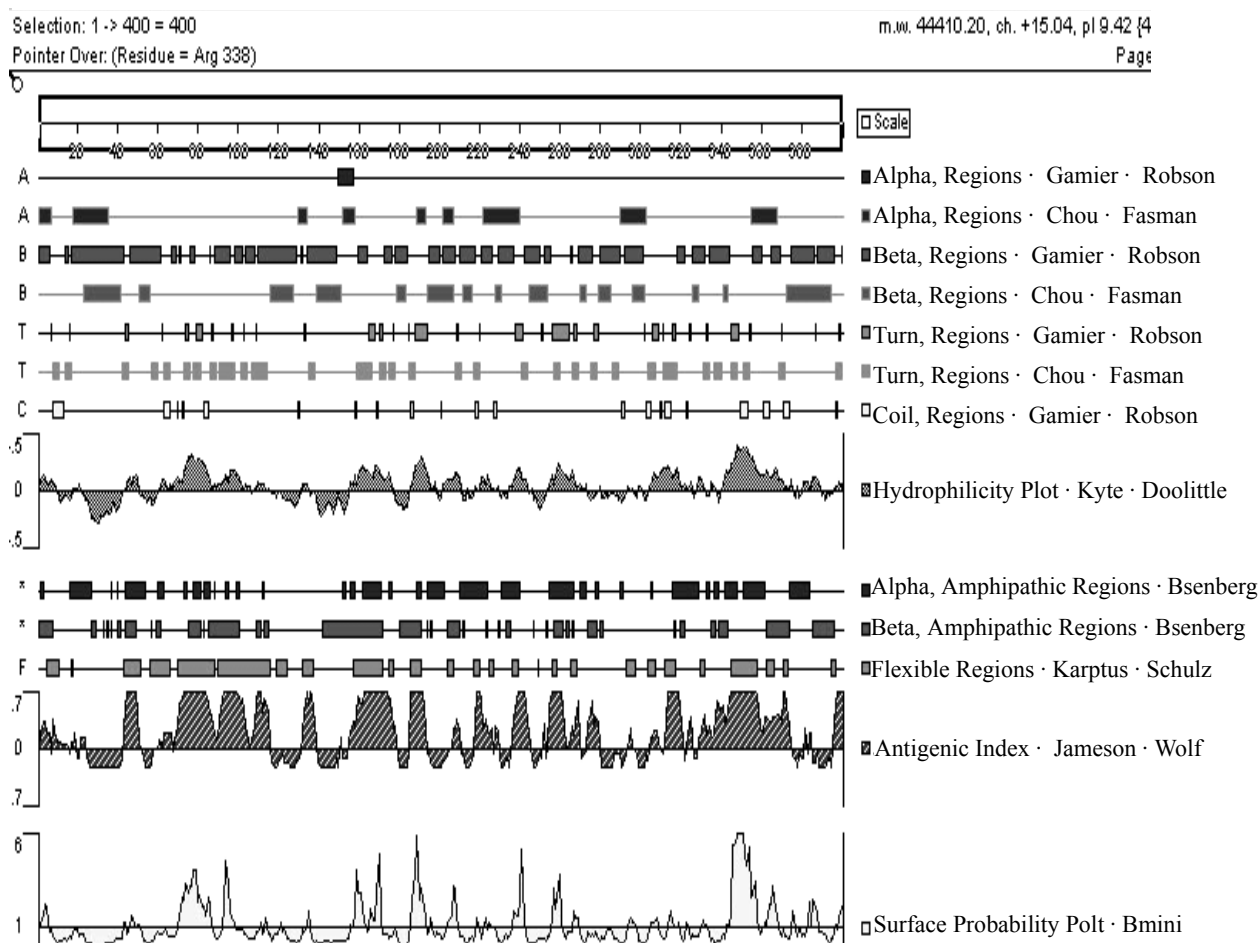


Fig. 9 The secondary structure of the $\beta 4$ Gal-T protein

Phylogenetic tree of Gal-T homologues from vertebrates

$\beta 4$ Gal-T encoded by a DNA fragment of 1 200 bases contains 399 amino acids. The cloned gene of $\beta 4$ Gal-T originally separated from cellular plasma contained 13 more amino acids than the $\beta 4$ Gal-T gene in the Golgi complex; the short sequence functions like a leader sequence to the former gene while transiting through the cellular membrane (Lopez *et al.*, 1991; Evans *et al.*, 1995).

Originating from mouse, the expression product of the cloned gene of $\beta 4$ Gal-T shares structural homology with $\beta 4$ Gal-Ts from other vertebrates, such as human, bovine and chick. But Uehara (1998) found that mouse testis $\beta 4$ Gal-T gene had some apparent phylogenetic difference with from genes of $\beta 4$ Gal-T and other glycosyltransferases.

Characterization of pMD 18-T vector

Modified from pUC18 vector, pMD 18-T vector is an ampicillin selective marker with a new cleavage site for EcoR V added between the cleavage sites of Xba I and Sal I in the polycloning site of pUC18 vector. After cleaved by EcoR V, thymines (T) were added to both of the 3'-ends of the linear fragment respectively.

pMD18-T vector was specially designed for cloning of PCR products, which helped to promote the efficiency of linkage of PCR products and sub-cloning.

DNA recovery from agarose gel

Among methods for recovery of DNA from agarose gel, DNA recovery kits are highly efficient but expensive. We developed an easy, economical

but efficient and stable method similar to low melting point gel processing but had quicker response time, which is suitable for subcloning of PCR products into pMD18-T vector.

Expression and characterization of β 4Gal-T

As reported, the expressions of human β 4Gal-T I in *E. coli* and yeast (Malissard *et al.*, 2000), and the expressions of human β 4Gal-T II in a transformed insect cell line (Tomiya *et al.*, 2003), had been successful.

This paper presents a GST-fused expression vector of mouse β 4Gal-T in *E. coli*. As an expression vector, pGEX-4T-2 engenders a GST fusion protein stabler than natural protein and with larger molecular weight than other natural proteins of *E. coli*; and have potential for detection and purification of target protein. Gradients of temperature, IPTG concentration and time were designed for the highly efficiency expression of proteins in the recombinant pGEX-4T-2 vector (pGEX-4T-2 β 4-Gal-T).

Studies on the structure and biophysics of β 4Gal-T were prevented by the propensity of recombinant human β 4Gal-T I proteins to aggregate when their concentrations were larger than 1 mg/ml. Malissard and Berger (2001) developed a mutated form of the catalytic domain of human β 4Gal-T I, which was proved much more soluble in aqueous solutions.

Park *et al.* (2002) got an expression of β 4Gal-T from *Neisseria meningitidis* and *Neisseria gonorrhoeae*, showing that most of the β 4Gal-Ts were insoluble and proteolysed into inactive enzymes that lacked C terminus residues (29.5 kD and 28 kD) when purified. The β 4Gal-Ts exhibited optimum activity at pH 6.5–7.0 in the presence of the Mn^{2+} ions. But in the presence of Fe^{2+} , Zn^{2+} or Cu^{2+} ions, Mn^{2+} ions could not show activity, so the β 4Gal-T activity did not increase any more. Interestingly, the β 4Gal-T activity could be stimulated to 150 per cent in the presence of non-ionic detergent Triton X-100 at the concentration of 0.1%–5 %.

Function of GST fusion protein

Glutathione S-transferase (GST; EC 2.5.1.18), encoded by the parasitic helminth *Schistosoma ja-*

ponicum, was a 26 kD protein with C terminal of Sj26. GST-fusion proteins are soluble in aqueous solutions and can be easily purified by affinity chromatography on immobilised glutathione. The GST part of the fusion protein can be cleaved off by digestion with site-specific proteases, such as thrombin or blood coagulation factor X, followed by absterion by glutathione-agarose, which can absorb the cleaved-off GST part and the uncleaved fusion protein (Smith and Johnson, 1988).

GST-fusion proteins can be applied as probes for qualitative detection in Western hybridization (Posern *et al.*, 1998). After separation on SDS-PAGE, GST-fusion proteins can be transformed to the detection membrane directly, followed by reaction of sealant, such as 5% bovine plasma albumin (BPA) or phosphate buffer solution (PBS) (Gillespie and Hudspeth, 1991). Goat anti-GST antibody as the primary antibody, and mouse anti-goat IgG marked with horse radish peroxidase (HRP) as the secondary antibody, were applied to the succeeding hybridization with GST-fusion proteins.

In addition, GST-fusion protein can be applied for sensitive quantitative detection or other complicated molecular analysis (Posern *et al.*, 1998; McDonald *et al.*, 1999).

CONCLUSION

As β 4Gal-T has important functions in many physiological aspects of different kinds of cells, especially in cell-cell recognition (Hathaway and Shur, 1996; Wassler and Shur, 2000; Wassler *et al.*, 2001) and rheumatoid arthritis (Yang *et al.*, 1996; Axford, 1999), further experimental research on this protein must be conducted. But it is complex and separating β 4Gal-T with classical biochemical approaches is hampered by low efficiency. We present the cloning and GST-fused expression of β 4Gal-T gene in *E. coli*, which have potential for further studies, and subsequently facilitating the medical application of β 4Gal-T.

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