

Abstract

ST332 is a 332 bp gene marker used in a patented *Salmonella* kit for detection of foodborne pathogen, *Salmonella* Typhi. As the gene is specific in DNA based detection by using polymerase chain reaction (PCR), its specificity in immunoassay is studied. For this purpose, an expression system of the gene is constructed and the expressed recombinant ST332 protein is tested for reactivity by hybridizing it towards typhoid patients sera through dot blotting. Initially, the DNA fragment was amplified with PCR and cloned into pGEX-4T-1 plasmid vector and transformed into *E. coli* BL21. Sequencing of the 332 bp fragment identified it as a hypothetical protein of *S. Typhi* strain CT18 (STY4528), a gene located on *Salmonella* Pathogenicity Island 7 (SPI-7) in *Salmonella*. The recombinant protein was expressed by induction with IPTG and extracted with NP-40 buffer. Analysis of the recombinant protein with dot blot immunoassay revealed weak affinity towards typhoid patients pooled-sera thus made it unreactive as an antigen. In another approach of obtaining specific *S. Typhi* antigen, a library of Novablue (*E. coli* DE3) cells expressing short fragment (150 – 300 bp) of *S. Typhi* genomic DNA was constructed by using Novatope Cloning System (Novagen). Here, about 700 clones were produced and the clones were pre-screened for reactivity towards typhoid patients' pooled-sera by colony blot immunoassay. Clones that showed higher reactivity than negative control were selected and further screened with dot blot immunoassay. Here, about 20 clones showing strong signals on dot blot were selected and further analyzed with ELISA immunoassay. Twice of the assays revealed that at least 3 clones (D1, G35 and I3) were highly reactive towards the typhoid patients antisera with at least 2-fold titre compared to typhoid negative sera.

Abstrak

ST332 adalah gen ‘marker’ bersaiz 332 bp yang digunakan dalam satu kit mengesan *Salmonella* Typhi yang telah dipaten. Memandangkan gen itu adalah spesifik dalam kaedah pengesan berdasarkan DNA iaitu melalui reaksi rantaian polimerasi (PCR), sifat spesifiknya terhadap kaedah pengesan yang lain iaitu assay imun dikaji. Untuk mencapai tujuan itu, satu sistem ekspresi protein itu dibina dan protein rekombinan yang dihasilkan itu disaring dengan sera terkumpul pesakit-pesakit demam kepialu melalui ujian “dot-blotting”. Pada mulanya, fragmen DNA bagi gen itu digandakan dengan teknik PCR, kemudian diklon ke vektor plasmid pGEX-4T-1, dan diubah ke dalam *E. coli* BL21. Analisis turutan nukleotida DNA untuk fragmen gen yang bersaiz 332bp itu mengenalpasti gen itu sebagai protein hipotetikal bagi *S. Typhi* strain C18 (STY4528) yang terletak di “*Salmonella* Pathogenecity Island” (SPI-7) dalam *Salmonella*. Ekspresi protein rekombinan dicetuskan dengan IPTG dan diekstrak dengan buffer Np-40. Analisis assay imun “dot-blot” terhadap protein itu menunjukkan reaksi yang lemah terhadap sera terkumpul pesakit-pesakit demam kepialu. Oleh itu, ia disimpulkan sebagai tidak reaktif sebagai antigen. Dalam pendekatan yang lain untuk mendapatkan antigen yang spefistik terhadap *S. Typhi*, satu kumpulan klon-klon yang membawa fragmen-fragmen DNA *S. Typhi* yang pendek telah dibina menggunakan ‘Novatope Cloning System’ daripada Novagen. Di sini, terdapat kira-kira 700 klon yang telah dihasilkan dan klon-klon tersebut diprasaring untuk kereaktifan terhadap sera terkumpul pesakit-pesakit demam kepialu menggunakan assay imun “colony-blot”. Klon-klon yang menunjukkan kereaktifan yang lebih tinggi berbanding kawalan negatif telah dipilih dan disaring sekali lagi dengan assay imun dot-blot. Terdapat dalam 20 klon yang menunjukkan signal yang kuat dalam ujian itu telah dipilih dan seterusnya dianalisis dengan assay imun ELISA. Assay yang diulang sebanyak dua kali itu

menunjukkan bahawa sekurang-kurangnya terdapat 3 klon (D1, G35 dan I3) mempunyai reaksi yang tinggi terhadap sera terkumpul pesakit-pesakit demam kepialu, dengan sekurang-kurangnya dua kali ganda titre berbanding sera yang negatif terhadap demam kepialu itu.

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