

## REFERENCES

1. White NJ. Melioidosis. Lancet 2003; 361: 1715-1722.
2. Dance DAB. Melioidosis as an emerging global problem. Acta Trop 2000; 74: 115-119.
3. Piggott JA, Hochholzerl. Human melioidosis. A histologic study of acute and chronic melioidosis. Arch Path 1970; 90: 101-11.
4. Wong KT, Putchucharey SD, Vadivelu J. The histopathology of human melioidosis. Histopathology 1995; 26: 51-55.
5. Punyagupta S. Melioidosis : review of 686 cases and presentation of a new Clinical classification. In : Punyagupta S, Sirisanthana T, Stapatayayong B, editors. Melioidosis. Proceedings of national workshop on melioidosis. Bangkok : Bangkok medical publisher; 1989. p.217-29.
6. Brett PJ, Woods DE. Pathogenesis of and immunity to melioidosis. Acta Trop 2000; 74: 201-210.
7. Dance DAB. Ecology of *Burkholderia pseudomallei* and the interactions between environmental Burkholderia spp. and human-animal hosts. Acta Trop 2000; 74: 159-168.
8. Vuddhakul V, Tharavichitkul P, Na-ngam N, Jitsurong S, Kunthawa B, Noimay P, et al. Epidemiology of *Burkholderia pseudomallei* in Thailand. Am J Trop Med Hyg 1999; 60: 458-61.
9. Heithoff DM, Conner CP, Mahan MJ. Dissecting the biology of a pathogen during infection. Trends In Microbiol 1997; 5: 509-513.
10. Finlay BB, Falkow S. Common themes in microbial pathogenicity revisited. Microbiol Mol Biol Rev 1997; 61: 36-69.
11. Mahan MJ, Slauch JM, Mekalanos JJ. Selection of bacterial virulence genes that are specifically induced in host tissues. Science 1993; 259: 686-688.

12. Hensel M, Shea JE, Gleeson C, Jones MD, Dalton E, Holden DW. Simultaneous identification of bacterial virulence genes by negative selection. Science 1995; 269: 400-403.
13. Valdivia RH, Falkow S. Fluorescence-based isolation of bacterial genes expressed within host cells. Science 1997; 277: 2007-2011.
14. Mahan MJ, Pleithoff DM, Sinsheimer RL, Low DA. Assessment of bacterial pathogenesis by analysis of gene expression in the host Annu Rev Gent 2000; 34: 139-64.
15. Suk K, Das S, Sun W, et al. *Borrelia burgdorferi* genes selectively expressed in the infected host. Proc Natl Acad Sci USA 1995; 92: 4269-4273.
16. Handfield M, Brady LJ, Progulske-fox A, Hillman JD. IVIAT : a novel method to identify microbial genes expressed specifically during human infections. Trends In Microbiol 2000; 8: 336-339.
17. Whitmore A, Krishnaswami CS. An account of the discovery of a hitherto undescribed infective disease occurring among the population of Rangoon. Indian Med Gaz 1912; 47: 262-7.
18. Stanton AT, Fletcher W. Melioidosis, a new disease of the tropics. Trans Fourth Congr Far East Assoc Trop Med 1921; 2: 196-8.
19. Yabuuchi E, Kosako Y, Oyaizu H, Yano I, Hotta H, Hashimoto Y, et al. Proposal of Burkholderia gen. Nov. and transfer of seven species of the genus pseudomonas group II to the new genus, with the type species Burkholderia cepacia (Palleroni and Holmes 1981) comb nov. Microbiol Immunol 1992; 36: 1251-75.
20. Phansomboon S. A study on the pathogenicity of \*(*B. whitmori*) in animals. J Med Assoc Thai 1943; 29: 13-29
21. Dharakul T, Songvilai S. The many facets of melioidosis. Trends In Microbiol. 1999; 7: 138-140.

22. Chaowagul W, White NJ, Danced DAB, Wattanagoon Y, Naigowit P, Davis TME, et al. Melioidosis : a major cause of community acquired septicemia in northeastern Thailand. J Infect Dis 1989; 159: 890-8.
23. Gilligan PH, Whittier S. Burkholderia, stenotrophomonas, Ralstonia, Brevundimonas, C. mamonas and Acidovorax. In : Baron EJ, Pfaffer MA, Tenover FC, Yolken RH, editors. Manual of clinical Microbiology. 7<sup>th</sup> ed. Washington, DC : American society for microbiology; 1999. p. 526-38.
24. Ashdown LR. An improved screening technique for isolation of *Pseudomonas pseudomallei* from clinical specimens. Pathology 1979; 11: 293-7.
25. Wuthiekanun V, Smith MD, Dance DAB, Walsh AL, Pitt TL, White NJ. Biochemical characteristics of clinical and environmental isolates of *Burkholderia pseudomallei*. J Med Microbiol 1996; 45: 408-12.
26. Smith MD, Angus BJ, Wuthiekanun V, White NJ. Arabinose assimilation defines a nonvirulent biotype of *Burkholderia pseudomallei*. Infect Immun 1997; 65: 4319-21.
27. Brett PJ, Deshazer D, Woods DE. *Burkholderia thailandensis* sp. Nov., a *Burkholderia pseudomallei*-like species. Int J. Syst. Bacteriol 1998; 48: 317-320.
28. Inglis TJ, Mee BJ, Chang BJ. The environmental Microbiology of melioidosis. Rev Med Microbiol 2001; 12: 13-20.
29. Wuthieknun V, Smith MD, White NJ. Survival of *Burkholderia pseudomallei* in the absence of nutrients. Trans R Soc Trop Med Hyg 1995; 89-491.
30. Tong S, Yang S, Lu Z, H.W. Laboratory investigation of ecological factors influencing the environmental presence of *Burkholderia pseudomallei*. Microbiol Immunol 1996; 40: 451-453.

31. Inglis T, Robertson T, Wikstrom M, Chang B, Mee B. Does *Burkholderia pseudomallei* exist in a viable but non -culturable form? In Proceedings of the 9<sup>th</sup> international congress of bacteriology and applied microbiology. Sydney, August 1999; 127.
32. Songsivilai S, Dharakul T. Multiple replicons constitute the 6.5- megabase genome of *Burkholderia pseudomallei*. Acta Trop 2000; 74: 169-179.
33. Dance DAB. Melioidosis as an emerging global problem. Acta Trop 2000; 74: 115-119
34. Leelarasamee A. Melioidosis in Southeast Asia. Acta Trop 2000; 74: 129-132
35. Wuthiekanun V, Smith MD, Dance DAB, White NJ. Isolation of *Burkholderia pseudomallei* from soil in north –eastern Thailand. Tran R So Trop Med Hyg 1995; 89: 41-43.
36. Trakul somboon S, Vuddhakul V, Tharavichitkul P, Na-gnam N, Suputtamongkol Y, Thamlikitkul V. Epidemiology of arabinose assimilation in *Burkholderia pseudomallei* isolated from patients and soil in Thailand Southeast Asian. J Trop Med Publ Health 1990; 30: 756-759.
37. Wuthiekanun V, Suputtamongkol Y, Simpson AJH, Kanaphun P, White NJ. Value of throat swab in the diagnosis of melioidosis. J Clin Microbiol 2001; 39: 3801-3802.
- 38 Leelarasamee A. *Burkholderia pseudomallei* : the unbeatable foe ? Southeast Asian. J Trop Med Publ Health 1998; 29: 410-415.
39. Sirisinha S, Anuntagool N, Dharakul T, et al, Recent developments in laboratory diagnosis of melioidosis. Acta Trop 2000 ; 74 : 235-245.
40. Suputtamongkol Y, Hall AJ, Dance DAB, Chaowagul W, Rajchanuvong A, Smith MD, et al. The epidemiology of melioidosis in Ubon Ratchatani, northeast Thailand. Int J Epidemiol 1994; 23: 1082-1090.

41. Currie B, Howard D, Vinh N, Withnall K, Merianos A. The 1990-1991 outbreak of melioidosis in the northern territory of Australia : clinical aspects. Southeast Asian J Trop Med Public Health 1993; 24: 436-443.
42. Nachiangmai N, Patamasucon P, Tipayamonthien B, Kongpon A, Nakaviroj S *Burkholderia pseudomallei* in southern Thailand. Southen Thailand. Southeast Asian J Trop Med Public Health 1985; 16: 83-87.
43. Suputtamongkol Y, Chaowagul W, Chetchotisakd P, Lertpatanasuwan N, Intaranongpai S, Ruchutrakool T, et al. Risk factor for melioidosis and bacteremic melioidosis. Clin Infect Dis 1999; 29: 408-13.
44. Lertpatanasuwan N, Sermsri K, Petkaseam A, Trakulsomboon S, Thamlikitkul V, Suputtamongkol Y. Arabinose -positive *B. pseudomallei* infection in human : case report. Clin infect Dis 1999; 28: 927 –928.
45. Gauthier YP, Hagen RM, Brochier GS, et al. Study on the pathophysiology of experimental *Burkholderia pseudomallei* infection in mice. FEMS Immun Med Microbiol 2001; 30: 53-63.
46. Ulett GC, Currie BJ, Clair TW, et al. *Burkholderia pseudomallei* virulence : definition, stability and association with clonality. Microbes Infect 2001; 3: 621-631.
47. Reckseidler SL, Deshazer D, Sokol PA, Woods DE. Detection of bacterial virulence genes of subtractive hybridization : identification of capsular polysaccharide of *Burkholderia pseudomallei* as a major virulence determinant. Infect Immun 2001; 69: 34-44.
48. Vorachit M, Lam K, Jayanetra P, Costerton JW, Electron microscopy study of the mode of growth of *Burkholderia pseudomallei* *in vitro* and *in vivo*. Am J Trop Med Hyg 1995; 98: 379-391.
49. Wong KT, Puthucheary SD, Vadivelu J. The histopathology of human melioidosis. Histopathology 1995; 26: 51-55.

50. Egan AM, Gordan DL. *Burkholderia pseudomallei* activates complement and is ingested but not killed by polymorphonuclear leukocytes. Infect Immun 1996; 64: 4952-4959.
51. Kespichayawattana W, Rattanachetkul S, Wanun T, Utaisincharoen P, Sirisinha S. *Burkholderia pseudomallei* induces cell fusion and actin associated membrane protrusion : a possible mechanism for cell -to-cell spreading. Infect Immun 2000; 68: 5377-5384.
52. Utaisincharoen P, Tangthawornchaikul N, Kespichayawattana W, Chaisuriya P, Sirisinha S. Burkholderia pseudomallei interferes with inducible nitric oxide synthase (iNOS) production : a possible mechanism of evading macrophage killing. Microbiol Immunol 2001; 45: 307-313.
53. Stevens MP, Woods MW, Taylor LA, et al. An Inv /Mxi –Spa-like type III Protein secretion system in *Burkholderia pseudomallei* modulates intracellular behaviour of the pathogen. Mol Microbiol 2002; 46: 649-659.
54. Suputtamongkol Y, Kwiatkowski D, Dance DAB, Chaowagul W, White NJ. Tumor necrosis factor in septicemic melioidosis. J Infect Dis 1992; 165: 561-64.
55. Friendland JS, Suputtamongkol Y, Remick DGm et al, Prolonged elevation of interleukin – 8 and interleukin –6 concentrations in plasma and of leukocyte interleukin-8 mRNA levels during septicemic and localized *Burkholderia pseudomallei* infection. Infect Immun 1992; 60: 2402-08.
56. Simpson AJH, Smith MD, Weverling GJ, et al, Prognostic value of cytokine concentrations (tumor necrosis factor, interleukin 6 and interleukin 10) and clinical parameters in severe melioidosis. J Infect Dis 2000; 181: 621-25.

57. Lauw FN, Simpson AJ, Prins JM, et al. The CXC chemokines gamma interferon (IFN –gamma) – inducible protein 10 and monokine induced by IFN –gamma are released during severe melioidosis. Infect Immun 2000; 68: 3888-93.
58. Lauw FN, Simpson AJ, Hack CE, et al. Soluble granzymes are released during human endotoxemia and in patients with severe infection due to Gram –negative bacteria. J Infect Dis 2000; 182: 206-13.
59. Nuntayawanwat S, Dharakul T, Chaowagul W, Songsivilai S. Polymorphism in the promoter region of tumor necrosis factor –alpha gene is associated with severe melioidosis. Hum Immunol 1996 ; 60: 979-83
60. Dharakul T, Vejbaesya S, Chaowagul W, Luangtrakool P, Stephens HA, Songsivilai S. HLA-DR and DQ associations with melioidosis. Hum Immunol 1998; 59: 580-86.
61. Finlay BB, Falkow S. Common themes in microbial pathogenicity revisited. Microbiol Mol Biol Rev 1997; 61: 36-69.
62. Smith H. What happens to bacterial pathogens *in vivo*? Trends In Microbiol 1998; 6: 239-243.
63. Heithoff DM, Conner CP, Mahan MJ. Dissecting the biology of a pathogen during infection. Trends in Microbiol 1997; 509-513.
64. Smith H. State and future of studies on bacterial pathogenicity : impact of new methods of studying bacterial pathogens. Washington DC : ASM press, 2000 : 265-282.
65. Mahan MJ, Heithoff DM, Sinsheimer RL, Low DA, Assessment of bacterial pathogenesis by analysis of gene expression in the host. Annu Rev Genet 2000; 34: 139-164.
66. Chiang SL, Mekalanos JJ, Holden DW. *In vivo* genetic analysis of bacterial virulence. Ann Rev Microbiol 1999; 53: 129-154.

67. Mekalanos JJ. Environmental singals controlling expression of virulence determinants in bacteria. J Bacteriol 1992; 174: 1-7.
68. Angelichio MJ, Camilli A. *In vivo* expression technology. Infect Immun 2002; 70: 6518-6523.
69. Handifield M, Levesque RC. Strategies for isolation of *in vivo* expressed genes from bacteria. FEMS Microbiol Rev 1999; 23: 69-91.
70. Perry RD. Signature -tagged mutagenesis and the hunt for virulence factors. Trends in Microbiol 1999; 7: 385-388.
71. Shea JE, Santangelo JD, Feldman RG. Signature -tagged mutagenesis in the identification of virulence genes in pathogens. Curr Opin Microbiol.
72. Deb DK, Dahiya P, Srivastava KK, Srivastava R, Srivastava BS. Selective identification of new therapeutic targets of *Mycobacterium tuberculosis* by IVIAT approach. Tuberculosis 2000; 82: 175-182
73. Ebersole JL, Frey De, Taubman MA, Smith DJ. An ELISA for serum antibodies to *Actinobacillus actinomycetemcomitans*. J Periodont Res 1980; 15 : 621-632.
74. Sambrook J, Fritsch EF, Maniatis T. Molecular cloning : A laboratory manual. 2<sup>nd</sup> ed. 1989. pp 6.3 Cold Spring Harbor : Cold Spring Harbor laboratory press.
75. Drury L. Transformation of bacteria by electroporation in Harwood A (ed). Methods in Molecular Biology, Vol 58 : basic DNA and RNA protocols. 1996 : pp . 249-256. Totowa : humana press.
76. Robertson D, Shore S, Miller DM. Manipulation and expression of recombinant DNA : a laboratory manual. 1997 : pp . 107-132. USA : Academic press.
77. Marlhak DR. Strategies for protein purification and characterization : a laboratory course manual. 1996. pp. 205-218. USA : Cold Spring Harbor laboratory press.

78. Oefner PJ, Hunicke –smith SP, Chiang L, Dietrich F, Mulligan J, Davis RW  
Efficient random subcloning of DNA sheared in a recirculating point –  
sink flow system. Nucleic Acid Research 1996; 24 (20) : 3879-3886.
79. Thorstenson YR, Hunicke-Smith Sp, Oefner PJ, Davis RW. An automated  
hydrodynamic process for controlled, unbiased DNA shearing.  
Genome research 1998; 8 : 848-855.
80. Besemer J, Borodovsky M. Heuristic approach to deriving models for gene  
finding. Nucleic Acids Research. 1999; 27(19) : 3911-3920.
81. Brown SM. Sequence similarity searches on the world wide web.  
Biotechniques. 1998; 24 (2) : 248-249.
82. Alschul SF, Madden TL, Schaffer AA, and J, Zang Z, Miller W et al. Gapped  
BLAST and PSI –BLAST : a new generation of protein data base  
search programns. Nucleic Acid Reasearch 1997; 25 (17) : 3389-2402.
83. Anuntagool N, Sirisinha S. Antigenic relatedness between Burkholderia  
pseudomallei and Burkholderia mallei. Microbiol Immuno. 2002; 46  
(3) : 143-150.
84. Pallen MJ. Microbial genomes. Molecular Microbiol 1999; 5 : 907-912
85. Isenberg HD, Damato RF. Indigenous and pathogenic microorganisms of  
human In Balows A, Hausler WJ, Herrmann KL, Isenberg HD,  
Shadomy HJ (eds.) Manual of clinical microbiology . 5<sup>th</sup> ed. 1991.  
pp2-14. Washington : American Society for microbiology.
86. Smith H. The behavior of bacterial pathogens in vivo. In Busdy DJW,  
Thomas CM, Brown NL (eds) Molecular Microbiology. Nato ASI series,  
vol. H 103. 1998. pp 319-335. Heidelberg : Springer –verlag berlin.
87. Chodimella U, Hoppe WL, Whalen S, Ognibene AJ, Rutecki GW.

- Septicemia and suppuration in a Vietnam veteran. Hospital pract 1997; 32 : 219-221.
88. Rost B, FariselliP, Casadio R. Topology prediction for helical transmembrane proteins at 86% accuracy. Protein Science 1996; 7: 170-1718.
89. Roberts IS. The biochemistry and genetics of capsular polysaccharide production in bacteria. Annu Rev Microbiol 1996; 50 : 285-315.
90. Masoud H, Ho M, Schollaardt T, Perry MB. Characterization of the capsular polysaccharide of *Burkholderia (Pseudomonas) pseudomallei* 304b. J Bacteriol 1997; 179 : 5663-5669
91. Claus H, Maiden CJ, Maag R, Frosch M, Vogel u. Many carried meningococci lack the genes required for capsule synthesis and transport. Microbiology 2002; 148 : 1813-1819.
92. Kneidinger B, Graninger M, Puchberger M, Kosma P, Messner P, Biosynthesis of nucleotide activated D-glycero -d-mano-heptose. J Biol Chem 2001; 276 : 20935-20944.
93. Gryllos I, Cywes C, Shearer MH, et al. Regulation of capsule gene expression by group A streptococcus during pharyngeal colonization and invasive infection. Mol Microbiol 2001; 42 : 61-74.
94. Hueck CJ. Type III Protein secretion systems in bacterial pathogens of animals and plants. Microbiol Mol Biol Rev 1998; 62 : 379-433.
95. Attree O, Attree I. A second type III secretion system in *Burkholderia pseudomallei* : who is the real culprit ? Microbiology 2001; 147 : 3197-3199.
96. Rainbow L, Hart CA, Winstanley C. Distribution of type III secretion gene clusters in *Burkholderia pseudomallei*, *B. thailandensis* and *B. mallei*. J Med Microbiol 2002; 51 : 374-384.

97. Vig E, Green M, Liu Y, et al. SIMPL is a tumor necrosis factor specific regulator of nuclear factor -kB activity. J Biol Chem 2001; 276:7859-7866.
98. Pfeuffer T, Goebel W, Laubinger J, Bachmann M, Kuhn M. Laxp 180, a mammalian Act A-binding protein, identified with the yeast two – hybrid system, co –localizes with intracellular *Listeria monocytogenes*. Cell Microbiol 2000; 2 : 101-14.
99. Woo PC, Leung PKL, Wong Ssy, Ho PL, Yuen KY. *groEL* encodes a highly antigenic protein in *Burkholderia pseudomallei*. ClinDiagn Lab Immun 2001; 8 : 832-836.
100. Hennequin C, Collignon A, Karjalainen T. Analysis of expression of *groEL* (HSP 60) of *Clostridium difficile* in response to stress. Microbial Pathogenesis 2001; 31 : 255-260.
101. Woo PC, Woo GK, Lau SK, Wong SS, Yuen JK. Single gene target bacterial identification *groEL* gene sequencing for discriminating clinical isolates of *Burkholderia pseudomallei* and *Burkholderia thailandensis*. Diagn Microbial Infect Dis 2002; 44 : 143-149.
- ศูนย์วิทยทรรพยากร  
จุฬาลงกรณ์มหาวิทยาลัย
102. Kruger NJ. Detection of polypeptides on immunoblots using secondary antibodies of protein A. In : Walker JM, ed. Basic protein and peptide protocols. Methods in Molecular Biology. USA : Humana Press, 1994 : 215-216.
103. Olesen CEM, Mosier J, Voyta JC, Bronstein I. Chemiluminescent

immunodetection protocols with 1, 2-dioxetane substrates. Methods In Enzymology 2000; 305 : 417-429.

104. Wongprompitak P, Thapthai C, Songsivilai S, Dharakul T. *Burkholderia pseudomallei* – specific recombinant protein and its potential in the diagnosis of melioidosis. Asian Pacific J. Allergy Immun 2001; 19 : 37-41.
105. Atkins T, Prior R, Mack K, et al. Characterization of an acapsular mutant of *Burkholderia pseudomallei* identified by signature tagged mutagenesis. J Med Microbiol 2002; 51 : 539-547.

ศูนย์วิทยทรัพยากร  
จุฬาลงกรณ์มหาวิทยาลัย



## APPENDICES

# ศูนย์วิทยทรัพยากร จุฬาลงกรณ์มหาวิทยาลัย

## APPENDIX A

### Media and solutions

Media / solutions	Composition
PBS pH 7.0 for IVIAT	500 mM KH <sub>2</sub> PO <sub>4</sub> 500 mM K <sub>2</sub> HPO <sub>4</sub>
PBST for IVIAT	PBS containing 0.025% Tween
TAE buffer	40 mM Tris-HCl pH 8.5 5 mM NaOAc 1 mM EDTA
Electrophoresis loading buffer	50 mM EDTA pH 7.0 50% (w/v) sucrose 0.05% (w/v) bromphenol blue
12.5% Acrylamide gel for Protein electrophoresis	Acrylamide 30 : 0.8 1.5 M Tris pH 8.8 50% glycerol
Acrylamide stacking gel for protein electrophoresis	Acrylamide 30 : 0.8 0.5 M Tris pH 8.8
30% acrylamide stock solution	29.2% (w/v) acrylamide 0.8% (w/v) NN' – methylene- bis-acrylamide
Electrophoresis buffer	25 mM Tris 192 mM glycine 0.1% (w/v) SDS

Media / solutions	Composition
5X sample buffer	60 mM Tris-HCl pH 6.8 25% (w/v) glycerol 2% (w/v) SDS 14.4 mM 2-mercaptoethanol 0.1% (w/v) bromphenol blue
Coomassie blue gel stain	0.1% (w/v) coomassie blue R-250 45% (v/v) methanol 45% (v/v) H <sub>2</sub> O 10% (v/v) glacial acetic acid
Coomassie blue gel destain	10% (v/v) methanol 10% (v/v) glacial acetic acid 80% (v/v) H <sub>2</sub> O
IB buffer for Inclusion	100 mM NaCl
Body preparation	1 mM EDTA 50 mM Tris-HCl pH 8.0
Dialysis buffer for Inclusion	50 mM Tris-HCl pH 8.0
Body preparation	50 mM KCl 5 mM Mg Cl <sub>2</sub> 1 mM DTT 50% glycerol
Lysozyme for Inclusion	1% (w/v) lysozyme in
Body preparation	10 mM Tris-HCl pH 7.5

Media / solutions	Composition
PBS for IVIAT	500 mM KH <sub>2</sub> PO <sub>4</sub> 500 mM K <sub>2</sub> HPO <sub>4</sub> pH 7.5
PBST for IVIAT	PBS containing 0.025% Tween
Western blot transfer buffer	48 mM Tris 39 mM glycine 0.05% SDS
TBS	20 mM Tris-HCl pH 7.5 150 mM NaCl
TBST	TBS containing 0.05% Tween
10X Assay buffer	200 mM Tris-HCl pH 9.8 10 mM MgCl <sub>2</sub>
SM	0.58% (w/v) NaCl 0.2% (w/v) MgSO <sub>4</sub> .7H <sub>2</sub> O 50 mM Tris-HCl pH 7.5 0.01% (w/v) gelatin
LB broth	1% (w/v) NaCl 1% (w/v) tryptone 0.5% (w/v) yeast extract pH 7.0
NZY broth	0.5% (w/v) NaCl 0.2% (w/v) MgSO <sub>4</sub> .7H <sub>2</sub> O 0.5% (w/v) yeast extract 1% (w/v) NZ amine, pH 7.5

Media / solutions	Composition
NZY agar	NZY broth containing 1% (w/v) agar
SOC	2% (w/v) tryptone 0.5% (w/v) yeast extract 0.05% (w/v) NaCl 2.5 mM KCl 20 mM MgCl <sub>2</sub> 20 mM glucose
2YT	1.6% (w/v) tryptone 1% (w/v) yeast extract 0.5% (w/v) NaCl, pH 7.0
Top agar	NZY broth containing 0.7% agarose

ศูนย์วิทยทรัพยากร  
จุฬาลงกรณ์มหาวิทยาลัย

## APPENDIX B

### DNA Sequence of isolated clones and genes

#### 1. Clone Bp1 whole inserted DNA sequence (1378 bp).

1 ACGATGCTCA GATATCCAA CGCCTGGCG GCTCCGGCAA GCTCGAAAAT  
51 TCCGTCCAGA AAGACAAATT TCTCGATCTC GAAAACATAG CCGTCGTCGC  
101 GATTGATCAC CCCATCGCGA TCCAGGAACA ACGCGCGATT CTTCACTTGC  
151 AGATTCCCGA AAGCTCGTCT TGAGCGCGGT AAAAGTCCTC GGGAACGCCG  
201 ATATCGATAA ACTGCCACG CGTCACGAAA ACGTCGAACC TGCGAGACTG  
251 GACTGCGGAC GACATGAAAT CGGTTTCGAA CGAGAACGCC TCCGCCGTCT  
301 CGCCGGCAAG AATGTCCTTC GGCAAGACAT AGCAACCTGC GTTGATCAAG  
351 CCCGGTCCCC AAACGCCCTT TTCGGCAAAA CCCGTCACCC GACCGCCGTC  
401 GACGACGAGG CGGCCGTATC GCCCCGTATC CGGCACCTGC CGGGCCACGA  
451 TCGTCGGGAA TCCACCCGTC TGCCAACCGT CGTCCAGTTC GTCGACTTCG  
501 AGATCCAGAT ACGTATCGCC GTTGAACACG AACGCGTGGT CGCCCTCGCA  
551 GTACGGCAAC GTTGCCTTC GTGCGCCTCC CGTACCCAAC GGATCGCTCT  
601 CGACTGAATA CGCGAGGTCG ATTCCGGCGA AGCGATCACC GAAATGGCTC  
651 ATGATTTCT CGGCCATGAA GCCAACGGAC AGGACCACCC GAGAAAACCTT  
701 CTTTCGCTC AAACGCGTCA AGAGGATTTC AAGAAATGGC CTGCCGGCAA  
751 TAGGCGCCAT CGGCTGGGA ACGTCGGAGA CGACCGTGCG CAAGCGCGTA  
801 CCGAACCCCTC CGGCCAAGAT GATCGCTTCT CTCATTCAC TGCTTCCCGA  
851 AAATGGAGTG CTCGACCAGC CCGCAAACGA TATGTCCGAG AACCAGGTGC  
901 CCCTCCTGGA TCTTCGGGGT GTCGGCCGAC GGGACTTCGA GCAGCAAGTC  
951 GCACAGCTCG CGCATTCTC CGCCCGATT GCCGGTGAAG CCGACGCACG  
1001 TCATCCCCTT TGCCTTCGCT TCGCGAAAAG CCGCCAGTAT GTTCGGGGAC  
1051 TTCCCGGAAG TCGAATAACC GATCAGCACG TCGCCCTCGT TGCCGAGCGC  
1101 CTGAACCTGA CGGGAAAACA GTTTCTCGTA GCCGTAATCG TTTCCGATCG  
1151 CCGTGAGAAT GGACGTGTCC GTCGTCAACG CCACTGCCGG CAGCCGGGGC  
1201 CGATCGAAAG CAAACCGGCT GACGAACTCC CCTGCGATAT GCTGCGCATC  
1251 AGCCCGCGTT CCTCCGTTAC CGGCGAGCAA CACCTTGCCG CCTTGAGCGA  
1301 TGGATGCAAT GCAGGCATCG GCAACTTCC GAACCGTGC AAGCAACCGT  
1351 TCGTCTGCCA GCATCGCGGC CATTACCC

2. *gmhA* gene (594 bp).

```

1 ATGGAGAAC GCGAATTGAC GTACATCACG AACAGCATCG CCGAGGCCA GCGGGTAATG
61 GCCCGATGC TGGCAGACGA ACGGTTGCTT GCGACGGTC AGAAAGTTGC CGATGCCTGC
121 ATTGCGTCCA TCGCTCAAGG CGGCAAGGTA TTGCTGCCG GTAACGGCGG AAGCGCGGCT
181 GATGCGCAGC ATATCGCAGG GGAGTTGTC AGCCGGTTG CTTCGATCG GCCCGGGCTG
241 CCGGCAGTGG CGTTGACGAC GGACACGTCC ATTCTCACGG CGATCGGAAA CGATTACGGC
301 TACGAGAAC TGTTTCCCG TCAGGTTAG GCGCTCGCA ACAAGGGCGA CGTGCTGATC
361 GGTATTGCA CTCCGGAAA GTCCCGAAC ATACTGGCGG CTTTCGCGA AGCCAAGGCA
421 AAGGGGATGA CGTGCCTCGG CTTCACCGGC AATCGCGCG GAGAAATGCG CGAGCTGTGC
481 GACTTGCTGC TCGAAGTCCC GTCGGCCGAC ACCCCGAAGA TCCAGGAGGG GCACCTGGTT
541 CTCGGACATA TCGTTGCGG GCTGGTCGAG CACTCCATT TCGGGAAGCA GTGA

```

3. *wcbM* gene (693 bp).

```

1 ATGAGAGAAC CGATCATCTT GGCCGGAGGG TTCGGTACGC GCTTGCAC
51 GGTCGTCTCC GACGTTCCA AGCCGATGGC GCCTATTGCC GGCAGGCCAT
101 TTCTGAAAT CCTCTTGACG CGTTGAGCG AAAAGAAGTT TTCTCGGGTG
151 GTCCTGTCGG TTGGCTTCAT GGCCGAGAAA ATCATGAGCC ATTCGGTG
201 TCGCTTCGCC GGAATCGACC TCGCGTATTG AGTCGAGAGC GATCCGTTGG
251 GTACGGGAGG CGCACTGAAG GCAACGTTGC CGTACTGCGA GGGCGACAC
301 GCGTTCGTGT TCAACGGCGA TACGTATCTG GATCTCGAAG TCGACGAAC
351 GGACGACGGT TGGCAGACGG GTGGATTCCC GACGATCGTG GCCCGGCAGG
401 TGCCGGATAC GGGCGATAC GGCGCCTCG TCGTCGACGG CGGTCGGGTG
451 ACGGGTTTG CCGAAAAGGG CGTTCGGGA CCAGGCTTGA TCAACGCAGG
501 TTGCTATGTC CTGCCGAAGG ACATTCTTGC CGGCGAGACG GCGGAGGCGT
551 TCTCGTTCGA AACCGATTTC ATGTCGTCCG CAGTCCAGTC TCGCAGGTTG
601 GACGTTTCG TGACGCGTGG GCAGTTATC GATATCGCG TTCCCGAGGA
651 CTTTACCGC GCTCAAGACG AGCTTCCGG AATCTGCAAG TGA

```

#### 4. Clone Bp3 whole inserted DNA sequence (1738 bp).

1 CGACGGCGTG CTGTTCTACC ATTCGAGCTG CCCCAGAACCC GGTATGCCCG  
51 GGCTCGCGCG AGTCTCGTCG ACGCCCTACC CCGACCCCCAC GCAGTTCGAT  
101 TCGCGCAGTC CGTACCACGA TCCGAAGTCG ACGCGGGAAAG CGCCGCGCTG  
151 GGTACTCGTC GACGTGCGCT TCGTCAGGAA ATCGCCGCTC GTTCCCCTCG  
201 CCGCGCTGCG CGAGCACGAG GCGCTCGCA ACATGCGCGT GCTCGCGAAG  
251 GGCAACCGGC TGTCGATCAC GCCCGTCACG CCAAACGAGT GGCGCTTCAT  
301 CACGCAGCGC CTGATGAAGT GACGCGGCCGCC ACACGTCAA ACATTGTCAG  
351 AGCCGGCGCG GCAAGCCGCG TCGGCCGGAA CCTCCGCCCCG CTTTCGGCGG  
401 CCTAACGGGC GCTCGACGAA TGCCGTGCCG CGTTGCCGG GCGTTTCGAG  
451 GAAAGCGCGG CGTCCGTCCG CGCGCCGCAC GCCCTCAAGC GCGTGCAGCGC  
501 TTACGCTCAA GGAGTCAACA ATGACGAAAA AATCCGCTCT TGCCGTTGCC  
551 GTCACGCTCG CGGCGGCCGT GCCGATCGCG CTCGCGCTCG CCCCCGTCGGC  
601 CGCGCGCGCG CAAAGCATGG GGCAGATGCA GCCGCCCCGCG GGCGTGCTGT  
651 CGCTGTCCGC GCAGGGCGAGC ACCGACGTCC CGCAGGACGT CGTCGATATC  
701 ACGCTGTTCT ACGAACAAACA GGCAGAAGGAC CGGGGCACGC TGACCGCGGA  
751 GCTGAACAAG CGCGCGGATA CGGCGCTCGC GCAGGGCGCGC GGGGTACCGG  
801 GCGTCACGGC CGCACGGGC GAATTCTCGG TGTCGCCGAG CGTCGATCGC  
851 GACGGCAAGA TCTCCGCGTG GCGCGGCCGC ACCGAGGTG TGCTCGAGTC  
901 GCACGACTTC GCGGCCGCAT CGAACGCTCGC CGGCCAGTTG AGCCCGATGA  
951 TGCAGGTGGG CAACGTGTCG TTCTCGCTGT CGCCCGAGGC GCAGGGCGCC  
1001 GCCGAGCAGA AACTCACGTC GGAGGGCGATC AAGGCGTTCC GCGCGCGCGC  
1051 CGAGGAAGCG ACGCGCGCGT TCGGCTACAG CAACTACTCG ATCCGCGAGG  
1101 TGAACGTGGG CAGCGGCCGC AACGTGCAGC CGTACCCGCG GATGTTCGCG  
1151 ATGGCCGCGC CCGCGATGGA CAGTGCAGAAG ATGAGGGCGC CGATGCCGT  
1201 CGAAGGGCGC AAGACGACCG TGACCGTCAA TGTGAACGGC TCGGTGCAGA  
1251 TGAAGTGACA CACGTGGCG CGCGCGTGGT GCGCCTCGCG ATCGATGCAA  
1301 AACGCCGGCG ATGCCGCCGC GTTTTTCTT GGGGGCCGTG TCGTCGGGA  
1351 TTCGCGGCCCG TGTGTCGCAC GGAACCGGCC AAGCGCGGTT GCGGGCGCGT  
1401 GGCCGCTCAT CGCACGGCG ATCGCGCCAG CGCAGCGTCC TGCGTGCCCC  
1451 CGTGTCCGTG TCCGTGTCG TGTCCGTGTC CGTGTCCGTG TCCGTGCTCC  
1501 GTGCTCCGTG CTCCGTGCTC CGCAGCCAGT ATGCAGGCAC ACGCGGGACA  
1551 CGTTTGACCG CACGCCGCCGC ACCGCCCCGCT CCACCGCCCCG CGGGCGGAGC  
1601 AACCGTGCAC AATCGCGCCG AACGATCAGG CCTGCTTCGC GACGCCGCCG  
1651 CGCCGATACG CCCAGATCAT CATCAGCACCG CGGGCGACGA TCATCGGCAG  
1701 CGACAGCCAC TGCCCCATCG ACAGGCCGAA CGTCAGCA

## 5. Conserved hypothetical protein gene (462 bp).

```

1 ATGCGCTACT GGCTGATGAA ATCCGAACCG GACGAGGCAA GCATCGACGA
51 CCTCGCCGCC GCCCCCCGACC AGACCTTGCC GTGGACCGGC GTGCGCAACT
101 ATCAGGCGCG CAACTTCATG CGCGACACGA TGAGATCGG CGACGGCGTG
151 CTGTTCTACC ATTCAAGCTG CCCCGAACCC GGTATCGCCG GGCTCGCGCG
201 AGTCTCGTCG ACGCCCTACC CCGACCCCCAC GCAGTTCGAT TCGCGCAGTC
251 CGTACACAGA TCCGAAGTCG ACGCGGGAAAG CGCCGCGCTG GGTACTCGTC
301 GACGTGCGCT TCGTCAGGAA ATCGCCGCTC GTTCCCCTCG CCGCGCTGCG
351 CGAGCACGAG GCGCTCGCGA ACATGCGCGT GCTCGCGAAG GGCAACCGGC
401 TGTCGATCAC GCCCGTCACG CCAAACGAGT GGCGCTTCAT CACGCAGCGC
451 CTGATGAAGT GA

```

## 6. Hypothetical signal peptide protein gene (738 bp).

```

1 ATGACGAAAA AATCCGCTCT TGCCGTTGCC GTCACGCTCG CGGCGGCCGCT
51 GCCGATCGCG CTCGCGCTCG CCCCGTCCGC CGCGCGCGCG CAAAGCATGG
101 GGCAGATGCA GCCGCCCCGCG GCGTGCTGT CGCTGTCCGC GCAGGGGAGC
151 ACCGACGTCC CGCAGGACGT CGTCGATATC ACGCTGTTCT ACGAACAAACA
201 GGCAGAAGGAC CCGGGCACGC TGACCGCGGA GCTGAACAAG CGCGCGGATA
251 CGGCGCTCGC GCAGGGCGCGC GGGGTACCGG GCGTCACGGC CCGCACGGC
301 GAATTCTCGG TGTCGCGAG CGTCGATCGC GACGGCAAGA TCTCCGCGTG
351 GCGCGGCCGC ACCGAGGTCG TGCTCGAGTC GCACGACTTC GCGGCCGCAT
401 CGAAGCTCGC CGGCCAGTTG AGCCCGATGA TGCAGGTGGG CAACGTGTCG
451 TTCTCGCTGT CGCCCGAGGC GCAGCGCGCC GCGAGCAGA AACTCACGTC
501 GGAGGGGATC AAGGCCGTTCC GCGCGCGCGC CGAGGAAGCG ACGCGCGCGT
551 TCGGCTACAG CAACTACTCG ATCCGCGAGG TGAACGTCGG CAGCGGCCGC
601 AACGTGCAGC CGTACCCGCG GATGTTCGCG ATGGCCGCGC CCGCGATGGA
651 CAGTGCAGAAG ATGAGCGCGC CGATGCCGT CGAAGGGCGC AAGACGACCG
701 TGACCGTCAA TGTGAACGGC TCGGTGCAGA TGAAGTGA

```

## 7. Clone Bp5 whole inserted DNA sequence (1,427 bp).

1 TTGATCCATG TATCCGTATC TTTCGGAGC GAAGACGGGG CCGCCTTCG  
51 CGTGGCGGCC GCTCGCGCA CGAACGTGC GATTCTAAC A TGCGCCCCAA  
101 CCCGCCCCAA CGCGGGCCCC TCTATATAAG AATGGCGAAA CGCGCGCGCA  
151 TGC GGCGCAC GCCCGCCGAT CGACGCGCGC ACAAAAAAAC ACAAAATGGC  
201 CGATGAAAGG TTCGCGCCTC AAAAAATCTG CAATTCCCG CCAACGGACT  
251 TGACAATTT CGAAACATGC GGCAAGCGGT GTTGCAC TCG CGCCACATTC  
301 GCTAGACCGA TGTTAGGGTG GGGAGGATT TTTTCGTGG GAAGGGAACG  
351 ATATGGCAAA ATGCCAACGC ACTGAAAGTT AGACGCTGCT CGAAGTCTAC  
401 ACAGAGCGGT GCCGCGTTT TTGTGCCGCA CCAATGTTAT ACTTCGAGCA  
451 ATGTATGACT TGT CATCGTC AACAGGCTCG CAGTAACCT GCGGTAATCT  
501 CAATTCGAG AGGAGAAATA TGAATAAACT TTCAAAGCTC GCGTTCAT TG  
551 CAGCTACCGC AGTTATGGCT GCATCCGCTT CGGCACAGTC GGTGCCGGCG  
601 TCGCGACAAG CCGTGAATGA CAACTGGGTG AATGGCACGG GCGAATGGGT  
651 GTGGATGAAC GGCACGAACG AGCTCTGCTG GCGCGATGCG TTCTGGACGC  
701 CGGCCACCGC CAACGCCAAG TGCGATGGCG CACTGGTCGC CCAGGCACCG  
751 GCACCGGCGC CGGTCGCACC GGTTGCTCCG GCCATCACGA GCCAGAAAGAT  
801 TACGTACCAA GCCGATA CGC TGTTGACTT CGACAAGGCC GTCCTGAAGC  
851 CGGCCGGCAA GCAGAAGCTT GACGA ACTGG CCGCGAAGAT CCAGGGCATG  
901 AACGTCGAAG TGGTCGTGGC CACGGGCTAC ACGGACCGCA TCGGTTCGGA  
951 CAAGTACAAC GACCGTCTGT CGCTGC CGCCGCAAGCC GTCAAGTCGT  
1001 ACCTCGTCAG CAAGGGTGTC CCGGCGAAC AAGTCTACAC GGAAGGCAAG  
1051 GCGAAGCGCA ACCCGGTAC GGGCAACACC TGCAAGCAGA AGAACCGCAA  
1101 GCAGCTCATC GCCTGCC TCG CACCGGACCG CGCGTGGAA GTCGAAGTGG  
1151 TCGGCACGCA GGAAGTGCAG AAGACGACCG TTCCGGCGCA GTAAGCCCG  
1201 AATCGATCGC ATCTGCTTCA AAAGCCCCCGC TCCGGCGGGG CTTTTTCA  
1251 GATGCGGGCCC GCCTCACCCG CGGCCGCGTT CCTGGCGACC TCGCCCGCG  
1301 GCTTATATAAC TCGTGGCTTG GCGGGCGCGC CGCCGCCCTC TTTCCCGCAT  
1351 CCGCTTGCGC ACATGACGAA CGCCGATCCG CACGA ACTCC AGAAATTCA  
1401 CGACCTCGCT CACAAATGGT GGGATCC

8. *ompA* gene, clone Bp5 (675 bp).

```

1 ATGAATAAAC TTTCAAAGCT CGCGTTCATT GCAGCTACCG CAGTTATGGC
51 TGCATCCGCT TCGGCACAGT CGGTGCCGGC GTCGCGACAA GCCGTGAATG
101 ACAACTGGGT GAATGGCACG GGCGAATGGG TGTGGATGAA CGGCACGAAC
151 GAGCTCTGCT GGCGCGATGC GTTCTGGACG CCGGCCACCG CCAACGCCAA
201 GTGCGATGGC GCACTGGTCG CCCAGGCACC GGCACCGGGCG CCGGTCGCAC
251 CGGTTGCTCC GGCCATCACG AGCCAGAAGA TTACGTACCA AGCCGATACG
301 CTGTTCGACT TCGACAAGGC CGTCCTGAAG CCGGCCGGCA AGCAGAAGCT
351 TGACGAACTG GCCGCGAAGA TCCAGGGCAT GAACGTCGAA GTGGTCTG
401 CCACGGGCTA CACGGACCGC ATCGGTTCGG ACAAGTACAA CGACCGTCTG
451 TCGCTGCGCC GCGCGCAAGC CGTCAAGTCG TACCTCGTCA GCAAGGGTGT
501 CCCGGCGAAC AAGGTCTACA CGGAAGGCAA GGGCAAGCGC AACCCGGTCA
551 CGGGCAACAC CTGCAAGCAG AAGAACCGCA AGCAGCTCAT CGCCTGCCCTC
601 GCACCGGACC GCCGCGTGG AAGTCGAAGTG GTCGGCACGC AGGAAGTGCA
651 AGACGACC GTTCCGGCGC AGTAAGA

```

9. Clone Bp6 whole inserted DNA sequence (2,500 bp).

```

1 TCGTCTCTAA GATCGGAATT CCGATGGTTG AACGCATGGC GACATACAAG
51 GAACTGAAGG CTCGGGCCGA GCGCCTGAGC GCGCAGGCCG AAGCGGCCG
101 GCAGGCTGAA TTGCAGGCCG CGATCGAAGA CGTGCAGCGCG AAGGTGCGCG
151 AGTACGGTCT CACGGCGTAC GACGTGTTCG GACACCGAAA GAAGCCGGGC
201 GAGCGCCATC GCAGGGCGGT CAGGCCGAAG TACCGCGATC CGGCGACGGG
251 CGCGACCTGG ACCGGACCGC GCATCGAGCC GAAATGGATC CGGGGCCGCA
301 ATCGGGACGA ATTCCCTGATC GAATGAAGGC GGGCGCCGCG CATCGCGCGC
351 GCCCTTGGCG AATCCGATAG AACGAAGGTG AACCACGATG AACATGCATG
401 TCGACATGGG ACGCGCGCTG ACCGTGCGCG ATTGGCCGGC GCTCGAGGCG
451 CTCGCGAAGA CGATGCCGGC CGATGCCGGC GCGCGGGCGA TGACCGACGA
501 CGATCTGCGC GCAGCGGGCG TCGATGCCCG CGTGCCGGAG CAAAGCTCG
551 GCGCGGCGAT CGACGAATTG GCGTCGCTCC GGCTGCCGA TCGGATCGAC
601 GGGCGCTTCG TCGATGCCCG CCGCGCGAAC CTCACGGTGT TCGACGATGC
651 ACGCGTGGCG GTGCGCGGCC ACGCGCGCGC GCAGCGAAC CTGCTCGAGC
701 GCCTGGAAAC CGAGCTCCTG GGCAGCACGC TGGACACCGC GGGCGACGAA
751 GGCAGGCGATCC AGCCGGACCC GATCCTTCAG GGGCTCGTCG ACGTGATCGG
801 CCAGGGCAAA TCCGATATCG ATGCGTACGC AACGATCGTC GAGGGGCTGA

```

851 CGAAGTACTT CCAGAGCGTC GCCGACGTGA TGAGCAAGCT GCAGGACTAC  
901 ATCTCGGCCA AAGACGACAA GAACATGAAG ATCGACGGCG GCAAGATCAA  
951 GGC GTT GATC CAG CAG GTCA TCG ACC ATCT GCCGACGATG CAG TTGCCGA  
1001 AGGGGGCCGA CAT CGC GCGC TGG CGCAAGG AGCT CGG CGA TGCCG TCTCG  
1051 ATCAGCGATT CGGGCGTCGT GACGATCAAT CCGGACAAGC TGATCAAGAT  
1101 GCGCGATT CGT CTG CCCCCTG ACGGCACGGT GTGGGACACC GCGCGCTACC  
1151 AGGCCTGGAA CACCGCGTTC TCCGGCCAGA AGGACAACAT CCAGAACGAC  
1201 GTGCAGACGC TCGTCAAGGAA ATACTCGCAC CAGAACTCGA ACTTCGACAA  
1251 TCTGGTCAAG GTGCTGAGCG GCGCGATCTC GACGCTCACG GACACCGCCA  
1301 AGAGCTATCT GCAGATCTGA ACGAGGGGCC GCGCCCACGC CGCCGCGCAT  
1351 CGCCGCGGCCG ATGCGCGCGG GCGCGGCCCT TGCCCCGATC CGAAGCAACC  
1401 GACAAGAGGA AAAGATT CGC CATGCCGCCG TCCATT CACC GAACTTCGTC  
1451 CATCAACAGC ACGCCGGCCG TGACCGCGGC CGCGCACGG CGCGCGTCCG  
1501 GGCCCGGCCG TGTTCTGTCG GCCGACGTCG CGCGCGTGGC GAGCGCGCGC  
1551 CGGCACACGA TGCCGAAAT CGCGCGCGG CGCGCGTCG ACGGCGATCG  
1601 CGCCGCGCCC GCGCCGCGCG AATCGTTCCG CGGGCGGCTC GAGACCGTAT  
1651 CGTCGCGCGC GCCGACGCCG CCGCCCGAGG CGGACTCGCG CGCCGCGAAT  
1701 GCGCGCACGA CGGGCGCCGC GGATGGGGCG GGGCGACGG GCGCGCGCG  
1751 CGGCCCATCG CACGGCGCGA CGTCGCTCGA TCGCGCGCG GCGTATCTGG  
1801 CCGAACTCGC CGGCGACCGT GGCGCGCGC TGACGACGCT CGTCGCGCAA  
1851 CTGAACGGCC ATGATCGCCG CGCGCTCGAC CATCTCGCGC TGACGGTGCA  
1901 TCGCCTGCAC CTGAGCGTCG ACGACGCGAG CCACCGGAAG ACGTTCGCCG  
1951 GCATCGCGA CGCGCTCGCG TCGTTCTCG CGGCCGCCGC GCGCGCGAAC  
2001 GCGAAACCCG GCTCGACGCC GCGCGATCTG TCGCTCGAGG GCAACAAGGG  
2051 TTACCGCAAG CTTGCGAGT GCGTCGGCGC GTTGCTGAGG TCCGACGGCA  
2101 TCGCGAACGC GCTGTTCGGC CGCGCGCGCG AGCGCGCGA CGCGCGCGCG  
2151 GACACGCTCG TGTCCGAGCG TCTCGCGCGC CGGATGGACG GGCACGTGCG  
2201 CACGAACGGG ATGGTGAACC GGCACGGCGG CGACCGCGAG CGCCGGATGG  
2251 CGGACGCCGC GCAGCGCTTC GCGCTCGCG CGCGCCACGC GACGAATCTC  
2301 GTCGACATGG TCGGCACCGC GCTGGAGCTG CTCGGCGCA CCGATCAGTT  
2351 GCTCAACGAC GTCTCGCTGA AGCGGCCGGC CGCCGAGCCG GGCGGCCGC  
2401 GCGCGCCGGG CGGCGGCCGC CGCGCGCCCG CGGGGCCGC GTCGGCGCG  
2451 GCGCAGCAGC CGGCCGCGCC GGCGCTCGTC AACAAATCATA ACGAGAACAA  
2501 CGTGAACGTG AATCTCGACG GGCTGGAGAG GCTCGCGC GATCTCGGCA  
2551 AGATGATGGG CGCGCTGCTC GAGCGCATGG

10. *bpH3* gene (291 bp), clone Bp6.

```

1 ATGGCGACAT ACAAGGAAC T G A A G G C T C G G G C C G A G G C G C TGAGCGCGCA
51 GGCGAAGCG GCGCGG CAGG CTGAATTGCA GGCCGCGATC GAAGACGTGC
101 GCGCGAAGGT GCGCGAGTAC GGTCTCACGG CGTACGACGT GTTCGGACAC
151 CGAAAGAACG CGGGCGAGCG CCATCGCGGG GCGGTCAAGGC CGAAGTACCG
201 CGATCCGGCG ACGGGCGCGA CCTGGACCGG ACGCGGCATC GAGCCGAAAT
251 GGATCCGGGG CCGCAATCGG GACGAATTCC TGATCGAATG A

```

11. *bipD* gene( 933 bp), clone Bp6.

```

1 ATGAACATGC ATGTCGACAT GGGACGCGCG CTGACCGTGC GCGATTGCC
51 GCGCTCGAG GCGCTCGCGA AGACGATGCC GGCGATGCC GGCGCGCGGG
101 CGATGACCGA CGACGATCTG CGCGCAGCGG GCGTCGATCG CCGCGTGCG
151 GAGCAAAAGC TCGGCGCGGC GATCGACGAA TTCGCGTCGC TCCGGCTGCC
201 CGATCGGATC GACGGCGCT TCGTCGATGG CCGCCGCGCG AACCTCACGG
251 TGTTCGACGA TGCACCGCGT GCGGTGCGCG GCCACGCGCG CGCGCAGCGC
301 AACCTGCTCG AGCGCCTGGA AACCGAGCTC CTGGCGGCA CGCTGGACAC
351 CGCGGGCGAC GAAGGCGGCA TCCAGCCGGA CCCGATCCTT CAGGGGCTCG
401 TCGACGTGAT CGGCCAGGGC AAATCCGATA TCGATGCGTA CGAACGATC
451 GTCGAGGGGC TGACGAAGTA CTTCCAGAGC GTCGCCGACG TGATGAGCAA
501 GCTGCAGGAC TACATCTCGG CCAAAGACGA CAAGAACATG AAGATCGACG
551 GCGGCAAGAT CAAGGCCTTG ATCCAGCAGG TCATCGACCA TCTGCCGACG
601 ATGCAGTTGC CGAAGGGGGC CGACATCGCG CGCTGGCGCA AGGAGCTCGG
651 CGATGCCGTC TCGATCAGCG ATTGGCGT CGTGACGATC AATCCGGACA
701 AGCTGATCAA GATGCGCGAT TCGCTGCCCT CTGACGGCAC GGTGTGGAC
751 ACCGCGCGCT ACCAGGCCTG GAACACCGCG TTCTCCGGCC AGAAGGACAA
801 CATCCAGAAC GACGTGCAGA CGCTCGTCA AAAATACTCG CACCAGAACT
851 CGAAACTCGA CAATCTGGTC AAGGTGCTGA CGGGCGCGAT CTCGACGCTC
901 ACGGACACCG CCAAGAGCTA TCTGCAGATC TGA

```

12. Clone Bp7 whole inserted DNA sequence (2,301 bp).

1 GGATCGTCGG CAGCACCTGC TTGCCGGCGA TCGTGTGCGC GCAGGCCGCC  
 51 ATGAAGCCGA CCGCGATTCC GGGATGAGC AGCAGCGGCA GCGCCGCGCC  
 101 GATCACGGGC ACGAGCGACA CCAGCATCAT CGCGAGCAGG TACGTGAAGA  
 151 ACAGTGTGAC GAATGCGAGC GGATTGCGCC GGAACAGCCA GATGCCTTGG  
 201 CGGAACCACA CGTAGCCGGT CTTGGCGGAG ACTTCGATCA GTTGCATGCG  
 251 TGGGTCTCGG GGAGCGCGCC CGCGTGCAGC ATGCGTTGCG GCAGGATGCG  
 301 TTCGAAATGG CCCGGGTCGT GCGGCTTGAG CATCTGCGCG GCGCGGGGAA  
 351 GGTGGAAATC ATACAGGCAGC GATACCCAGA AGCGGTACGC GCCCGCGCGC  
 401 AGCATGTGCG CCCAGTGGCG GCGCTCGCCC GCGGTGAACG GGCGCACCGT  
 451 CTGGTACGCG CGCAGCAGCG CGTCGGCGCG CGCGCGCTCG AGCGCGCCCG  
 501 TCGGCAGATC GACGCACCAAG TCGTTGACCG TCACCGCGAC GTCGAACAGC  
 551 CATTGTCGC AGCCGGCGAA GTAGAAATCG AAGAAGCCGC CGAGCCGCAC  
 601 CGAATGGCCG GTGTGGGCT CCGCGTGCAGC GAAGAGCGCA TTGTCGCGAA  
 651 ACAGGTCGCA ATGGCACGGG CCTTCCGGCA GCGCCGCGTA ATCGTCCGAT  
 701 GCGAAGAACG CGGCCCTGGT CGCGAGCTCG CCTTCCAGCA GCGCGCGCTG  
 751 CTCGCCCCGTG ACGAACGGCG CGATCGCGGG CACCGTGTG CGCCACCACG  
 801 GCAGGCTGCG CAAGTTGGC TGATGCCGCG GATAGTCGCG GCCCGCGAGG  
 851 TGCATGCGCG CGAGCATCTG CCCGACTTCG ACGCAGTGCT CGACGCCCGG  
 901 CGCGAGCTGC GCCGCGCCCT CGAGCTTGGT GACCGATCGCG GCCGGCTTGC  
 951 CGTGCAACTC GCCGAACAGC GTGCCGTCGT CGCGCGCGAC GGGCGCGGGC  
 1001 ACGGGCACCC CGTGTTCGC GAGATGGCTC ATCAGATCGA CGTAGAACGG  
 1051 CAATTGCCCG GCCGTCAGGT TCTCGAAGAT CGTGAGCACG TATTGCGCCG  
 1101 GCGTCGTCGT CAGGAAGAAG TTGCTGTTCT CGATGCCGGA CGGAATGCCG  
 1151 CGGAACGCGA CAACGTCGCC GAGATCGTAG TGGCGCATCC AGAGTGCAGG  
 1201 GTCAGCGTCG GAAACTGCGG TGAAAACGGC CATGCAGGAA ACGTCGGTTC  
 1251 GGGTTGGCGC GCCGGCGCG TGCCGGTGTG GCAAAAGGGC GGAAGGGGGC  
 1301 CGGCGCCGCG CGGCGCCGGC GGTGCGTCAA TAGTGCAGGT TGACCGACGG  
 1351 CAGGCGCGTG ACAGGCACGC CGGCGTCGTG CGGGCGCGGC GACGTATCGG  
 1401 GCGACGCGCT CATCTGATAG CGGGTGCCGA AGTCGATT CACGTTGATC  
 1451 TCGACCGGCT TGCCGCGATC CCGGAATTG GTGACTTCGG TGCCGTTCTT  
 1501 GCTTTTTTCG TGAAAGCTCG GCGTGCAGCG CACGTCGTTG AAATCGACCT  
 1551 TCGAAGTCAC TTGGCGCCG GGGCGGTTGA TCTTCGTGAG ATCGGGCAGC  
 1601 CGGGCCGCCT CGTTGGCCGC GGCGTGGGCC TTTGCGTCGG CGGCCGCTTG  
 1651 TCGGGCGTCG CGGGCGCGGG CGGGGCCGAC GAGGGCGAGT GCCGTCAGGG

1701 CGGCAGCGAA AAGGAGCGGC TTCATCGTGT TTCTCCATT GAACCGTTCG  
 1751 ATTTAGCAA ATACCGGCCT CGTGCGATG CCGTCGACG ATGCGCGAGC  
 1801 GGCCTGCCG CCGCGATGGG CATCGGCTGC GCGAGCGGGT TCCGTATAA  
 1851 TGTCGAAACG ATCTGAAGAG GCACTGCTAG ATGAAGAACG ATCCAACCG  
 1901 CCGTTCCCGG ACGCGCACGC CGGGTAGCCC GTGCGTCGAA GCGTCGACG  
 1951 ACCCGATCGC CGCCGTCGCG CGGCTCTCCG AGATCTACGA GACGAACACC  
 2001 GCGTTCTGC GCGACGCGTT CGCGCGCTAT CGCGCAACG AAGCGTTCGA  
 2051 CGAGCACGTG CGCGCGTGCT ATCCGTTCGT GCGCATCCGC ACCGACGTCA  
 2101 ACACGCACAT CGATTCGCGC CGCTCGTACG GCTTCGTCGC CGGCCCCGGC  
 2151 GTGTCGAGA CGACCGTCAC GCGCCCGGAC CTGTCGCGA ACTACTACCG  
 2201 CGAGCAATTG CGCCTGCTCG CGAAGAACCA TCACGTTGG ATCGAAGTCG  
 2251 GCGTATCGGC GCAGCCGATC CCGGTTCACT TCGCGTTCTC CGAAGGCATT  
 2301 C

### 13. Transmembrane protein gene (813 bp), clone Bp7.

1 ATGCAACTGA TCGAAGTCTC CGCCAAGACC GGCTACGTGT GGTTCCGCCA  
 51 AGGCATCTGG CTGTTCCGGC GCAATCCGCT CGCATTGTC AACTGTTCT  
 101 TCACGTACCT GCTCGCGATG ATGCTGGTGT CGCTCGTGC CGTGATCGGC  
 151 CGGGCGCTGC CGCTGCTGCT CATTCCCGGA ATCGCGGTGCG GCTTCATGGC  
 201 GGCTGCCGC GACACGATCG CCGGCAAGCA GGTGCTGCCG ACGATCCTGA  
 251 TCGACGGCTT CCGCTCGTAC GGCCCACCG TCACCGAGCG GCTGCTCGCG  
 301 CTCGGCGGGC TCTACATCGT TTCGATGGCG GCCGTGTTCG CGTGCTCGGC  
 351 GCTCGGCGAC GGCGGCACGC TGCTGAAGAT CATGTTCGGT CTCGGCGCCG  
 401 AGAACCTCGG GCCGGAGGCG CTCGATTCCG CGGGCGTCAG GATCGCGGTA  
 451 CTGATCGCGG CCGCGCTGTA CGCGCCGGTC GCGATGATGT TCTGGTTCGC  
 501 GCCGGTGTG ACCCGCTGGC ACGACGTGCC GCCCGTAAA GCGCTGTTCT  
 551 TCAGCGTCGT GAGCTGCTGG CGCAACAAGG GCGCGTTCAC CGTCTATGGA  
 601 CTGCTGTGGT TCGCGCTAGC GCTCGCGTG TCGTCGGGC TCGCCGCGCT  
 651 GATGCAGGCG CTCGGCGCCA GCGCCTACGC GCTCACGGTG ATGATGCCGG  
 701 CCTCGATCGT CATCACCGCG ATGCTCTACT GCTCCTCTA TGCAACCTAT  
 751 CGCGGCTGCT TCGCGTGCA GGAGCCGGGG GCGCAGAACATC CGCCGAACGC  
 801 ATCCGGCCGT TGA

14. Homoserine kinase protein gene (996 bp), clone Bp 7.

```

1 ATGGCTGTT TCACCCGGT CACCAACGCC GAGATGCCGC TCTGGCTGGA GCAATACGAC
61 GTGGGCACGG TCCGCGCGCT GCGCGGCATT CCCTCGGGGA TCGAAAACAC CAACTTCTC
121 CTGACCACGG AGAAGGACGG CGCCACGCAC GAGTACGTCG TCACGCTGTT CGAGCGGCTG
181 ACCAGCGAGC AACTGCCGTT CTACCTGTAC CTGATGCAGC ATCTGGCGCA GCACGGCATC
241 TCGTGCCGG CGCCGATTCC CGGCCGCGAC GGCGCGATCC TGCGCCCGCT CAAGGGCAAG
301 CCGCGACCA TCGTACGCG CCTGCCCCGA CGCTGAACC TGGCGCCCAC GACGAGCGAA
361 TGCGCCATCG TCGGCACAT GCTGGCGCGC ATGCACCTGG CCGGCCGCGA CTACCCGCG
421 CACCAGCCCA ACCTGCGCAG CCTGCCGTGG TGGAACGAAG TGGTGCCCGA CATCCAGCCC
481 TTCGTGCAGG GCGCCACGCG CGAGCTGCTG GTCGCCGAGC TGGCCCACCA GCAGCGCTC
541 TTGGCAGCG CCGACTATGC CGCCCTGCCC GAGGGCCCGT GCCACTGCGA CCTGTTCCGC
601 GACAACGTGC TGTCGAGCC GGCCACTGAC AGCCAGCCCC AGCGCCTGGG CGGGTTCTT
661 GATTCTATT TCGCCGGCGT CGACAAATGG CTGTCGACG TGGCCGTGAC CGTCAACGAC
721 TGGTGCCTCG ACCTGCCAC GGGTGCCTCG GATGCCAAC GGATGCGCGC CATGCTGCGC
781 GCCTATCACG CGGTGCGGCC TTTCACCGAC GCGGAGGCC GTCACTGGCG GGACATGCTG
841 CGCGCCGCGG CCTATCGCTT CTGGGTATCG CGCCTGTGGG ACTTCCACCT GCCGCGCGAC
901 GCCGAACTGC TGCAGCCGCA TGATCCGACC CACTTCGAGC GCGTGCTGCG CGAACGGGTG
961 CGCGCCGAGG GGCTGACATT GGATATTCCC GAACCATGCA ACTGA

```

15. Transmembrane protein gene 2 (399 bp), clone Bp7.

```

1 ATGAAGCCGC TCCTTTCGC CGCCGCCCTG ACGGCACCTCG CCCTCGTCGG
51 CCCCGCCCGC GCCGCCGACG CCGCACAAAGC GGCCGCCGAC GCAAAGGCC
101 AGGCCGCGGC CAACGAGGCG GCCGGGCTGC CCGATCTCAC GAAGATCAAC
151 CGCCCCGGCG CGGAAGTGAC TTCGAAGGTC GATTCAACG ACGTGCCTCG
201 CACGCCGAGC TTTCACGAAA AAAGCAAGAA CGGCACCGAA GTCACCGAAT
251 TCCGGGATCG CGGCAAGCCG GTCGAGATCA ACGTGAAATC GAACTTCGGC
301 ACCCGCTATC AGATGAGCGC GTCGCCGAT ACGTCGCCGC GCCCGCACGA
351 CGCCGGCGTG CCTGTCACGC GCCTGCCGTC GGTCAACCTG CACTATTGA

```

## 16. Clone Bp9 whole inserted DNA sequence (1300 bp).

1 AGCGCAGCTT CGGCAGCCCG ACGGTACGA AGGACGGTGT GTCGGTCGCG  
51 AAGGAAATCG AGCTGAAGGA CAAGCTCCAG AACATGGCG CGCAGATGGT  
101 CAAGGAAGTC GCTTCCAAGA CCAGCGACAA CGCCGGCGAC GGCACGACGA  
151 CGGCCACCCT CTCGCGCAA TCGATCGTCC GCGAAGGCAT GAAGTACGTC  
201 GCATCGGGCA TGAACCCGAT GGACCTGAAG CGCGGCATCG ACAAGGCAGT  
251 CGCCGCGGCA GTCGAAGAGC TGAAGAAGAT CAGCAAGCCG TGCACGACGA  
301 ACAAGGAAAT CGCGCAAGTC GGCGCGATCT CGCGAACAG CGATTCTCG  
351 ATCGCGATC GCATCGCTGA AGCGATGGAC AAGGTCGGCA AGGAAGGCCTG  
401 GATCACCGTC GAAGACGGCA AGTCGCTCGC CGACGAGCTC GACGTCGTCG  
451 AAGGCATGCA GTTCGACCGC GGCTACCTGT CGCCGTACTT CATCAACAAC  
501 CCGGACAAGC AAGTCGCCGT CCTCGAGAAC CCGTTCGTGC TGCTGCACGA  
551 CAAGAAGGTG TCGAACATCC GCGACCTGTT GCCGGTGCTC GAGCAAGTCG  
601 CGAAGGCTGG CGTCCGCTG CTGATCATCG CCGAAGACGT CGAAGGCAGA  
651 GCGCTCGCAA CGCTGGCGT CAACAACATC CGCGGCATCC TGAAGACCGT  
701 TGCAGTCAAG GCGCCGGCT TCGCGATCG TCGCAAGGCG ATGCTGGAAG  
751 ACATCGCGAT CCTGACGGGC GGCCAGGTCA TCGCGGAAGA AACCGGCCTC  
801 ACGCTCGAGA AGGCAACGCT GGCAGAACTG GGCCAGGCGA AGCGCATCGA  
851 AGTGGCAAG GAAAACACGA CGATCATCGA CGCGCGGGC GAAGCCGTGA  
901 ACATCGAAGC GCGCGTCAAG CAAATCCGCA CGCAAATCGA AGAAGCGACA  
951 TCGGACTACG ACCGTAAAAA GCTGCAAGAG CGCGTGGCCA AGCTGGCAGG  
1001 CGCGTGGCG GTGATCAAGG TTGGCGCTGC GACCGAAGTC GAAATGAAGG  
1051 AAAAGAAGGC ACGTGTGAG GACGCGCTGC ACGCCACCCG CGCTGCCGTT  
1101 GAAGAAGGCA TCGTCCCAGG CGCGCGCGTC GCGCTGATCC GCGCACGCAC  
1151 CGCGATCGCG GGCCTGACCG GCGTGAACGC CGACCAAGAAC GCCGGCATCA  
1201 AGATCGTGCT GCGCGCGATG GAAGAGCCGC TGCGCCAGAT CGTCACGAAC  
1251 GGCGGCGAAG AAGCGAGCGT CGTGGTGGCG GCAGTTGCTG CGGGCAAGGG

17. *groEL* gene(1596 bp), clone Bp9.

1 ATGGTCGAAG GCGTGAACAT TCTGCCAAC GCTGTGAAGG TCACGCTGGG  
 51 TCCGAAGGGC CGCAACGTGG TGCTCGAGCG CAGCTCGGC GGCCCACGG  
 101 TCACGAAGGA CGGTGTGTCG GTCGCGAAGG AAATCGAGCT GAAGGACAAG  
 151 CTCCAGAACAA TGGGCGCGCA GATGGTCAAG GAAGTCGCTT CCAAGACCAG  
 201 CGACAACGCC GGCGACGGCA CGACGACGGC CACCGTCCTC GCGCAATCGA  
 251 TCGTCCGCCA AGGCATGAAG TACGTCGCAT CGGGCATGAA CCCGATGGAC  
 301 CTGAAGCGCG GCATCGACAA GGCAGTCGCC GCGGCAGTCG AAGAGCTGAA  
 351 GAAGATCAGC AAGCCGTGCA CGACGAACAA GGAAATCGCG CAAGTCGGCG  
 401 CGATCTCGGC GAACAGCGAT TCGTCGATCG GCGATCGCAT CGCTGAAGCG  
 451 ATGGACAAGG TCGGCAAGGA AGGCGTGATC ACCGTCGAAG ACGGCAAGTC  
 501 GCTCGCCGAC GAGCTCGACG TCGTCGAAGG CATGCAGTTC GACCGCGGCT  
 551 ACCTGTCGCC GTACTTCATC ACAAACCCGG ACAAGCAAGT CGCCGTCTC  
 601 GAGAACCCGT TCGTGCTGCT GCACGACAAG AAGGTGTCGA ACATCCGCGA  
 651 CCTGTTGCCG GTGCTCGAGC AAGTCGCGAA GGCTGGCCGT CCGCTGCTGA  
 701 TCATCGCCGA AGACGTCGAA GGCGAAGCGC TCGCAACGCT GGTCGTCAAC  
 751 AACATCCGCG GCATCCTGAA GACCGTTGCG GTCAAGGCGC CGGGCTTCGG  
 801 CGATCGTCGC AAGGGGATGC TGGAAGACAT CGCGATCCTG ACGGGCGGCC  
 851 AGGTCATCGC GGAAGAAACC GGCCTCACGC TCGAGAAGGC AACGCTGGCA  
 901 GAACTGGGCC AGGCGAAGCG CATCGAAGTG GGCAAGGAAA ACACGACGAT  
 951 CATCGACGGC GCGGGCGAAG CCGTGAACAT CGAAGCGCGC GTCAAGCAAA  
 1001 TCCGCACGCA AATCGAAGAA GCGACATCGG ACTACGACCG TGAAAAGCTG  
 1051 CAAGAGCGCG TGGCCAAGCT GGCAGGGCGG GTGGCGGTGA TCAAGGTTGG  
 1101 CGCTGCGACC GAAGTCGAAA TGAAGGAAAA GAAGGCACGT GTCGAGGACG  
 1151 CGCTGCACGC CACCCCGCCT GCCGTTGAAG AAGGCATCGT CCCGGGCGGC  
 1201 GGCCTCGCGC TGATCCGCGC ACGCACCGCG ATCGCGGGCC TGACCGGCGT  
 1251 GAACGCCGAC CAGAACGCCG GCATCAAGAT CGTGTGCGC GCGATGGAAG  
 1301 AGCCGCTGCG CCAGATCGTC ACGAACGGCG GCGAAGAAGC GAGCGTCGTG  
 1351 GTGGCGGCAG TTGCTCGGG CAAGGGCAAC TACGGCTACA ACGCGGCGAC  
 1401 GGGCGAGTAC GTCGACATGG TCGAAGCCGG CGTCGTCGAT CCGACGAAGG  
 1451 TCACCCGTAC CGCGCTGCGAG AACGCGGCTT CGGTGCGCCGG CCTGCTGCTG  
 1501 ACGACGGACG CAGCCGTTGC CGAACTGCCG AAGGAAGACG CTCCGATGCC  
 1551 GGGCGGCATG CCGGGCGGCA TGGGCGGCAT GGGCATGGAC ATGTAA

## BIOGRAPHY

Mr.Siroj Jitsurong was born on September 21, 1957 in Lumpang, Thailand. He received his Bachelor degree of Science (Med Tech) in 1980, Chiangmai University and Master's degree of Tropical Medicine in 1985 from Faculty of Tropical Medicine, Mahidol University. His current position is Assistant Professor, Department of Pathology, Faculty of medicine, Prince of Songkhla university. He has enrolled the Royal Golden Jubilee Scholar (RGJ) in graduate program for the Degree of Doctor of Philosophy in Medical Microbiology, Inter-departmental program in Medical Microbiology, Chulalongkorn University since 1998.

ศูนย์วิทยทรัพยากร  
จุฬาลงกรณ์มหาวิทยาลัย