

## REFERENCES

1. Doerfler,W. (1983). DNA methylation and gene activity. *Annu. Rev. Biochem.*, 52, 93–124.
2. Riggs,A.D. and Jones,P.A. (1983). 5-Methylcytosine, gene regulation, and cancer. *Adv. Cancer Res.*, 40,1–30.
3. Gardiner-Garden,M. and Frommer,M. (1987). CpG islands in vertebrate genomes. *J. Mol. Biol.*, 196, 261–282.
4. International Human Genome Sequencing Consortium. (2001). Initial sequencing and analysis of the human genome. *Nature*, 409, 860–921.
5. Bird,A., Taggart,M., Frommer,M., Miller,O.J. and Macleod,D. (1985). A fraction of the mouse genome that is derived from islands of nonmethylated, CpG-rich DNA. *Cell*, 40,91–99.
6. Goto,T. and Monk,M. (1998). Regulation of X-chromosome inactivation in development in mice and humans. *Microbiol. Mol. Biol. Rev.*,62,362-378.
7. Tycko,B. (1997). DNA methylation in genomic imprinting [see comments]. *Mutat. Res.*,386, 131–140.
8. Issa,J.P.(2000). CpG-island methylation in aging and cancer. *Curr. Top.Microbiol. Immunol.*, 249, 101–118.
9. Bestor,T., Laudano,A., Mattaliano,R.and Ingram,V. (1988). Cloning and sequencing of cDNA encoding DNA methyltransferase of mouse cells. The carboxyl-terminal domain of the mammalian enzymes is related to bacterial restriction methyltransferases. *JMol. Biol.* 203, 971–983.
- 10.Okano,M., Bell,D.W., Haber,D.A. and Li,E. (1999). DNA methyltransferases Dnmt3a and Dnmt3b are essential for de novo methylation and mammalian development. *Cell* 99, 247–257.
- 11.Hansen,R.S., Wijmenga,C., Luo,P., Stanek,A.M., Canfield,T.K. Weemaes,C.M. and Gartler,S.M. (1999). The DNMT3B methyltransferase gene is mutated in

12. Wolffe,A.P., Jones,P.L. and Wade,P.A. (1999). DNA demethylation. *Proc. Natl. Acad. Sci. U.S.A.* 96, 5894–5896.
13. Jost,J.P., Schwarz,S., Hess,D., Anglker,H., Fuller-Pace,F.V., Stahl,H., Thiry,S. and Siegmann,M. (1999). A chicken embryo protein related to the mammalian DEAD box protein p68 is tightly associated with the highly purified protein-RNA complex of 5-MeC-DNA glycosylase. *Nucleic Acids Res.* 27, 3245–3252.
14. endrich,B., Hardeland,U., Ng,H.H., Jiricny,J. and Bird, A. (1999). The thymine glycosylase MBD4 can bind to the product of deamination at methylated CpG sites. *Nature* 401, 301–304.
15. Bhattacharya,S.K., Ramchandani,S., Cervoni,N.and Szyf,M. (1999). A mammalian protein with specific demethylase activity for mCpG DNA. *Nature* 397, 579–583.
16. Jones, P.A.and Takai,D. (2001). The role of DNA methylation in epigenetics. *Science* 293, 1068–1070.
17. Comb, M.and Goodman, H.M. (1990). CpG methylation inhibits proenkephalin gene expression and binding of the transcription factor AP-2. *Nucleic Acids Res.* 18, 3975 – 3982.
18. Baylin,S. (2002). Mechanisms underlying epigenetically mediated gene silencing in cancer. *Semin. Cancer Biol.* 12, 331.
19. Nan, X., Ng, H.H., Johnson, C.A., Laherty,C.D., TurnerB.M., Eisenman, R.N.and Bird A. (1998). Transcriptional repression by the methyl-CpG-binding protein MeCP2 involves a histone deacetylase
20. Tissenbaum,H.A. and Guarente,L. (2001). Increased dosage of a *sir-2* gene extends lifespan in *Caenorhabditis elegans*. *Nature* .410, 227- 230.complex. *Nature* 393, 386–389.
21. Spotswood,H.T. and Turner, B.M. (2002). An increasingly complex code. *J. Clin. Invest.*110, 577–582.
22. Manel Estelle. (2002). Cancer as an epigenetic disease: DNA methylation and chromatin alterations in human tumors. *journal of pathology*.196, 1-7.

23. Pallis,M., Robins,A. and Powell,R. (1993). Quantitative analysis of lymphocyte CD11a using standardized flow cytometry. *Scand. J. Immunol.* 38, 559–564.
24. Yoder,J.A., Walsh,C.P., and Bestor,T.H. (1997). Cytosine methylation and the ecology of intragenomic parasites. *Trends Genet.* 13: 335–340.
25. Bird, A.P. (1995). Gene number, noise reduction and biological complexity. *Trends Genet.* 11: 94–100.
26. Razin, A. and Shemer,R. (1995) DNA methylation in early development. *Hum Mol Genet.Spec* 4:1751-5.
27. Li, E., Bestor, T.H., and Jaenisch, R. (1992). Targeted mutation of the DNA methyltransferase gene results in embryonic lethality. *Cell* 69, 915–926.
28. Okano, M., Bell, D.W. and Haber, D.A., Li, E. (1999). DNA methyltransferases Dnmt3a and Dnmt3b are essential for de novo methylation and mammalian development. *Cell* 99, 247–257.
29. Wilson, V.L., Smith, R.A., Ma, S.and Cutler, R.G. (1987). Genomic 5-methyldeoxycytidine decreases with age. *J.Biol. Chem.* 262, 9948–9951.
30. Vanyushin,B.F., Nemirovsky,L.E., Klimenko,V.V., Vasiliev,V.K. and Belozersky, A.N. (1973). The 5-methylcytosine in DNA of rats. Tissue and age specificity and the changes induced by hydrocortisone and other agents. *Gerontologia* 19, 138–152.
31. Rath,P.C. and Kanungo,M.S. (1989). Methylation of repetitive DNA sequences in the brain during aging of the rat. *FEBS Lett.* 244, 193–198.
32. Romanov,G.A. and Vanyushin,B.F. (1981). Methylation of reiterated sequences in mammalian DNAs. Effects of the tissue type, age, malignancy and hormonal induction. *Biochim. Biophys. Acta* 653, 204–218.
33. Lengauer, C., Kinzler, K.W. and Vogelstein,B. (1997). DNA methylation and genetic instability in colorectal cancer cells. *Proc. Natl. Acad. Sci. U.S.A.* 94, 2545–2550.
34. Barbot, W., Dupressoir, A., Lazar, V. and Heidmann, T. (2002). Epigenetic regulation of an IAP retrotransposon in the aging mouse: progressive demethylation

- and de-silencing of the element by its repetitive induction. Nucleic Acids Res. 30, 2365-73.
35. Ono,T., Shinya, K., Uehara, Y. and Okada, S. (1989). Endogenous virus genomes become hypomethylated tissue specifically during aging process of C57BL mice. Mech. Ageing Dev. 50,27-36.
36. Christoph Plass. (2002). Cancer epigenomics. Human Molecular Genetic. 20,2479-2488
37. Gama-Sosa MA, Midgett RM, Slagel VA, Githens S, Kuo KC, Gehrke CW and Ehrlich M. (1983a). Biochim. Biophys. Acta, 740, 212-219.
38. Melanie Ehrlich. (2002). DNA methylation in cancer:too much, but also too little. Oncogene.21, 5400-5413.
- 39.Chen RZ, Pettersson U, et al. (1998). DNA hypomethylation leads to elevated mutation rates. Nature. 395,89-93.
- 40.Xu GI, Bestor. (1999)Chromosome instability and immunodeficiency syndrome caused by mutation in a DNA methyltransferase gene. Nature.402, 187-191
41. Thayer RE,Singer MF. (1993). Undermethylation of specific LINE-1 sequences in human cell producing a LINE-1 encoding protein. Gene.133, 273-277.
42. Garrick D, Fiering S, Martin D and Whitelaw E. (1998). Repeat-induced gene silencing in mammals. Nat.Genet., 18, 56-59.
43. Morgan HD, Sutherland HG, Martin DI and Whitelaw E.(1999). Epigenetic inheritance at the agouti locus in the mouse. Nat. Genet., 23, 314- 318.
44. Dupressoir A and Heidmann T. (1997). Expression of intracisternal A-particle retrotransposons in primary tumors of oncogene-expressing transgenic mice. Oncogene, 14, 2951-2958.
45. Yates, P. A., Burman, R. W., Mummaneni, P., Krussel, S., and Turker, M. S. (1999). Tandem B1 elements located in a mouse methylation center provide a target for de Novo DNA methylation. J. Biol. Chem. 274, 36357–36361
46. Smit, A. F. A., and Riggs, A. D. (1996). Tiggers and other DNA transposons fossils in the human genome. Proc. Natl. Acad. Sci. U.S.A. 93, 1443–1448

47. Ostertag, E. M., and Kazazian, H. H., Jr. (2001). Biology of mammalian L1 retrotransposons. *Annu. Rev. Genet.* 35, 501–538.
48. Robertson, K. D., and Wolffe, A. P. (2000). DNA methylation in health and disease. *Nature Rev.* 1, 11–19.
49. Eickbush, T. H., and Malik, H. S. (2002). Origins and evolution of retrotransposons. In *Mobile DNA II* (N. L. Craig, R. Craigie, M. Gellert, and A. M. Lambowitz, Eds.), pp. 1111–1144. ASM Press, Washington.
50. Loeb, D. D., Padgett, R. W., Hardies, S. C., Shehee, W. R., Comer, M.B., Edgell, M. H., and Hutchison, C. A., III. (1986). The sequence of a large L1Md element reveals a tandemly repeated 5' end and several features found in retrotransposons. *Mol. Cell.Biol.* 6, 168–182.
51. Kolosha, V. O., and Martin, S. L. (1997). In vitro properties of the first ORF protein from mouse LINE-1 support its role in ribonucleoprotein particle formation during retrotransposition. *Proc. Natl. Acad. Sci. U.S.A.* 94, 10155–10160.
52. Weiner, A. M. (2002). SINEs and LINEs: The art of biting the hand that feeds you. *Curr. Opin. Cell Biol.* 14, 343–350.
53. Moran, J. V., DeBerardinis, R. J., and Kazazian, H. H., Jr. (1999). Exon shuffling by L1 retrotransposition. *Science* 283, 1530–1534.
54. Symer, D. E., Connelly, C., Szak, S. T., Caputo, E. M., Cost, G. J., Parmigiani, G., and Boeke, J. D. (2002). Human L1 retrotransposition is associated with genetic instability in vivo. *Cell* 110, 327–338.
55. Holmes, S. E., Dombroski, B. A., Krebs, C. M., Boehm, C. D., and Kazazian, H. H., Jr. (1994). A new retrotransposable human L1 element from the LRE2 locus on chromosome 1q produces a chimaeric insertion. *Nat. Genet.* 7, 143–148.
56. Kazazian, H. H., Jr., and Goodier, J. L. (2002). LINE Drive: Retrotransposition and genome instability. *Cell* 110, 277–280.
57. Gilbert, N., Lutz-Prigge, S., and Moran, J. V. (2002). Genomic deletions created upon LINE-1 retrotransposition. *Cell* 110, 315–325.

58. Viel, A., Petronzelli, F., Della Puppa, L., Lucci-Cordisco, E., Fornasarig, M., Pucciarelli, S., Rovella, V., Quaia, M., Ponz de Leon, M., and Boiocchi, M., Genuardi, M. (2002). Different molecular mechanisms underlie genomic deletions in the MLH1 gene. *Hum. Mutat.* 20, 368–374.
59. Loeb, L. A. (2001). A mutator phenotype in cancer. *Cancer Res.* 61, 3230–3239.
60. Miki Y, Nishisho I, Horii A, Miyoshi Y, Utsunomiya J, Kinzler KW, Vogelstein B and Nakamura Y. (1992). Disruption of the APC gene by a retrotransposal insertion of L1 sequence in a colon cancer *Cancer Res.*, 52, 643 -645.
61. Wei W, Gilbert N, OoiSL, LawlerJF,OstertagEM,Kazazian HH, Boeke JD and Moran JV. (2001). Human L1 retrotransposition: cis preference versus trans complementation. *Mol.Cell. Biol.*, 21, 1429 –1439.
62. Whitelaw, E. and Martin, DI. (2001). Retrotransposons as epigenetic mediators of phenotypic variation in mammals. *Nat. Genet.*, 27, 361-365.
63. Kazazian Jr HH and Moran JV. (1998). The impact of L1 retrotransposons on the human genome. *Nat. Genet.*, 19, 19 -24.
64. Chen, R. Z., Pettersson, U., Beard, C., Jackson-Grusby, L., and Jaenisch, R. (1998). DNA hypomethylation leads to elevated mutation rates. *Nature* 395, 89-93.
65. Takai, D., Yagi, Y., Habib, N., Sugimura, T., and Ushijima, T. (2000). Hypomethylation of LINE1 retrotransposon in human hepatocellular carcinomas, but not in surrounding liver cirrhosis. *Jpn. J. Clin. Oncol.* 30, 306–309.
66. Dante, R., Dante-Paire, J., Dominique, R., and Roizes, G. (1992). Methylation patterns of long interspersed repeated DNA and alphoid repetitive DNA from human cell lines and tumors. *Anticancer Res.* 12, 559–564.
67. Jürgens, B., Schmitz-Dräger, B. J., and Schulz, W. A. (1996). Hypomethylation of L1 LINE Sequences prevailing in human urothelial carcinoma. *Cancer Res.* 56, 5698–5703.

68. Santourlidis S, Florl A, Ackermann R, Wirtz HC and Schulz WA. (1999). High frequency of alterations in DNA methylation in adenocarcinoma of the prostate. *Prostate*, 39, 166-174.
69. Ushijima T, Morimura K, Hosoya Y, Okonogi H, Tatematsu M, Sugimura T and Nagao M. (1997). Establishment of methylation-sensitive-representational difference analysis and isolation of hypo- and hypermethylated genomic fragments in mouse liver tumors. *Proc. Natl. Acad. Sci. USA*, 94, 2284 - 9
70. Zhenggang Xiong and Peter W.Laird. (1997). COBRA: a sensitive and quantitative DNA methylation assay. *Nucleic Acids Research* 25, 2532-2534.
72. Blenda CK Wong and YM Dennis LO. (2003). Cell-free DNA and RNA in plasma as new tools for molecular diagnostic. *Mol. Diagn.* 3(6), 785-797.
73. Lyon (1994). The X inactivation and X chromosome imprinting. *Eur j Human Genet.* 2 (4), 255-61.
74. Wolf Reik, Wendy, and Jorn Walter (2001). Epigenetic reprogramming in mammalian development. *Science*. 293, 1089-1093.
75. Vera Schramke and Robin Allshire (2003). Hairpin RNAs and retrotransposon LTRs effect RNAi and chromatin-based gene silencing. *Science*. 301, 1069-1074.
76. Amir Eden, Francois Gaudet and Alpana Waghmare (2003). Chromosomal instability and tumors promoted by DNA hypomethylation. *Science*. 30,455.
77. Florl AR, Lower R and Schulz WA(1999). DNA methylation and expression of LINE-1 and HERV-K provirus sequences in urothelial and renal cell carcinomas. *Br J Cancer*. 80,1312-1321.
78. Goelz SE, Vogelstein B and Feinberg AP (1985). Hypomethylation of DNA from benign and malignant human colon neoplasm. *Science*. 228,187-189.
79. Suter CM, Martin DI and Ward RL (2003). Hypomethylation of L1 retrprtransposons in colorect cancer and adjacent normal tissue. *Int j Colorectal Dis.*

## APPENDIX A

### BUFFERS AND REAGENT

#### 1. Lysis Buffer 1

Sucrose	109.54	g
1.0 M Tris – HCl (pH 7.5)	10	ml
1.0 M MgCl <sub>2</sub>	5	ml
Triton X – 100 (pure)	10	ml
Distilled water to	1,000	ml

Sterilize the solution by autoclaving and store in a refrigerator (at 4<sup>o</sup>C).

#### 2. Lysis Buffer 2

5.0 M NaCl	15	ml
0.5 M EDTA (pH 8.0)	48	ml
Distilled water to	1,000	ml

Sterilize the solution by autoclaving and store at room temperature.

#### 3. 10% SDS solution

Sodium dodecyl sulfate	10	g
Distilled water to	100	ml

Mix the solution and store at room temperature.

#### 4. 20 mg/ml Proteinase K

Proteinase K 2 mg

Distilled water to 1 ml

Mix the solution and store in a refrigerator (at -20°C).

#### 5. 1.0 M Tris – HCl

Tris base 12.11 g

Dissolve in distilled water and adjusted pH to 7.5 with HCl

Distilled water to 100 ml

Sterilize the solution by autoclaving and store at room temperature.

#### 6. 0.5 M EDTA (pH 8.0)

Disodium ethylenediamine tetraacetate.2H<sub>2</sub>O 186.6 g

Dissolve in distilled water and adjusted pH to 8.0 with NaOH

Distilled water to 1,000 ml

Sterilize the solution by autoclaving and store at room temperature.

7. 1.0 M  $\text{MgCl}_2$  solution

Magnesium chloride.6H<sub>2</sub>O 20.33 g

Distilled water to 100 ml

Dispense the solution into aliquots and sterilize by autoclaving.

#### 8. 5 M NaCl solution

Sodium chloride 29.25 g

Distilled water to	100	ml
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Dispense the solution into aliquot and sterilize by autoclaving.

9. 10X Tris borate buffer (10X TBE buffer)

Tris – base	100	g
Boric acid	55	g
0.5 M EDTA (pH 8.0)	40	ml

Adjust volume to 1,000 ml with distilled water. The solution was mixed and store at room temperature.

10. 6X loading dye

Bromphenol blue	0.25	g
Xylene cyanol	0.25	g
Glycerol	50	ml
1M Tris (pH 8.0)	1	ml
Distilled water until	100	ml

Mixed and stored at 4°C

11. 7.5 M Ammonium acetate ( $\text{CH}_3\text{COONH}_4$ )

Ammonium acetate	57.81	g
Distilled water	80	ml

Adjust volume to 100 ml with distilled water and sterilize by autoclaving.

**12. 25:24:1 (v/v) Phenol-chloroform-isoamyl alcohol**

Phenol	25	volume
Chloroform	24	volume
Isoamyl alcohol	1	volume

Mix the reagent and store in a sterile bottle kept in a refrigerator.

**13. 12% Non-denature acrylamide gel (w/v)**

40%acrylamide: Bis (19:1)	3	ml
5X TBE	1	ml
10% ammoniumpersulfate	105	μl
TEMED	8	μl
H <sub>2</sub> O	6	ml

Dissolve by heating in microwave oven and occasional mix.

**14. TE buffer**

Tris base	1.21	g
5M EDTA	200	μl

Adjust pH to 7.5 with conc.HCL and adjust volume to 1.0 litre with H<sub>2</sub>O.

## APPENDIX B

### Sequence of LINE-1 and primer

Human transposon L1.2 ACCESSION M80343 (5'UTR)

#### BEFORE BISULFITE

Ggggggaggagccaagatggccgaataggaacagctccggctcacagctcccgatcggtgacgcacgcagaa  
 gacggtgattctgcattccatctgaggtaccgggtcatctcacttagggagtgccagacagtggcgaggcca  
 gtgtgtgcgcaccgtgcgcagccgaagcagggcgaggcattgcctcacctggaaagcgcaaggggtcag  
 ggagttcccttctgagtcaaagaaagggtgacggtcgcacctggaaaatcggtcactcccacccgaatattgc  
 gctttcagaccggctaagaaacggcgaccacgagactatacccacacctggctcgagggctacgccc  
 acggaatctcgctgattgctgacacgactgtagatcaaactgcaaggcggcaacgaggctggggagggg  
 cggccgcattgccaggctgcttagtaaacaaggcagc

#### AFTER BISULFITE

GgggggaggagTTaagatggTCGaataggaATagTtTCGgtTtaTagTtTTTagCGtagCGaCG  
 TagaagaCGgtgattTtgTatTTatTtgaggtTCGggtTatTtTaTtagggagtTTagaTagtgggC  
 GTaggTTagtgtgtgCGTaTCGtgCGCGagTCGaagTagg

#### FCORBA LINE-1

gCGaggTattgTTtTaTTtggaaCGTaagggtTagggagtTTTttTtgagtTaaagaa

TasI

TaqI

AggggtgaCGgtCGTaTTtggaaaatCGggtTaTtTTTaTTCGaaatattgCGTtttTga

#### RCORBA LINE-1

TCGgTtaagaaaCGgCGTaTTaCGagaTtatTTTaTaTTtgTtCGgagggtTTtaCGTTTaC  
 GgaatTtCGTgattgTtagTaTagTagtTtgagatTaaaTtgTaaggCGgTaaCGaggTtggggagg  
 ggCGTTCGTTattgTTTtaggTttgTttagtaaaTaaagTagT

F COBRALINE-1 = ccgTaagggtTagggagtTTT

R COBRALINE-1 = RTAAAACCCTCCRAACCAAATATAAA

Total amplicon = 160 bp and TasI = 63+ 97 and TagI = 80+80

## BIOGRAPHY

Miss Krisanee Chalitchagorn was born in Nakornsawan in 1979. She graduated from Faculty of Science, Chulalongkorn University in Biochemistry program and then attended to participate in Medical Science program in Faculty of Medicine for her master degree.

