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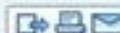
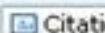
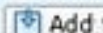
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- British Journal of Industrial Medicine (80)
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- Preventive Medicine (66)
- Lancet (62)

- Hong, W.K. (35)
- Wichmann, H.E. (35)
- Albanes, D. (33)
- Samet, J.M. (33)
- Benhamou, S. (32)
- 1999 (286)
- 1998 (294)
- 1997 (267)
- 1996 (269)
- 1995 (226)

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<input type="checkbox"/> International Journal of	<input type="checkbox"/> Jockel, K.H. (36)	<input type="checkbox"/> 2000 (306)	Less...	<input type="checkbox"/> Physics and Astronomy (104)
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An important method for silencing tumor suppressor genes in cancers is by aberrant methylation (referred to as methylation) of CpG islands in gene promoter regions. In lung cancer, methylation of the genes retinoic acid receptor β -2 (RAR β -2), CDH13 (H-cadherin), p16INK4a (p16), RASSF1A (RAS association domain family 1) is frequent. Thus, we investigated methylation of these genes in 4 different types of specimens (oropharyngeal brushes, sputum samples, bronchial brushes and bronchioalveolar lavage [BAL] samples) of the upper aerodigestive tract epithelium from heavy smokers without evidence of cancer but with morphometric evidence of sputum atypia and compared the frequencies of methylation in the different types of specimens. In addition, we also analyzed sputum samples from 30 never smokers for methylation of these genes. Our major findings are: (i) At least one gene was methylated in one or more specimens from 48% of the smokers. However, methylation was statistically significant less frequently in never smokers compared to smokers. (ii) In general, methylation occurred more frequently in samples from the central

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Volume 107, Issue 4, 20 November 2003, Pages 612-616

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CD133 positive endothelial progenitor cells contribute to the tumour vasculature in non-small cell lung^a Hamon Ctr. Therapeut. Oncol. Res., Univ. of TX Southwestern Medical Center, Dallas, TX, USA^b Clinical Division of Oncology, Department of Medicine 1, University of Texas Southwestern Medical Center, Dallas, TX, USA^c British Columbia Cancer Agency, Vancouver, BC, Canada^d Department of Pathology, Univ. of TX Southwestern Medical Center, Dallas, TX, USA^e Department of Internal Medicine, Univ. of TX Southwestern Medical Center, Dallas, TX, USA^f Department of Pharmacology, University of Texas Southwestern Medical Center, Dallas, TX, USA

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High-throughput oncogene mutation profiling in human cancer
(2007) *Nature Genetics*
[Abstract + Refs](#)

- [Sato, N., Fukui, T., Higuchi, T., Yokoyama, K., Kondo, M., Nagasaka, G., Y. \(...\), Sekine](#)

Personal

Name	Minna, John D.
Other formats	Minna, J. D. Minna, J. Minna, John
Author ID	7101758359
Affiliation	University of Texas Southwestern Medical Center at Dallas, Department of Internal Medicine Dallas United States

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