- hypermethylation in colorectal cancer. *Cancer Epidemiol Biomarkers Prev* 2009;**18**:3086–96.
- ⁴¹ Bolton-Smith C, Woodward M, Tunstall-Pedoe H et al. Accuracy of the estimated prevalence of obesity from self reported height and weight in an adult Scottish population. J Epidemiol Community Health 2000;54:143–48.
- ⁴² Nyholm M, Gullberg B, Merlo J et al. The validity of obesity based on self-reported weight and height: implications for population studies. *Obesity*; 15:197–208.
- ⁴³ Spencer EA, Appleby PN, Davey GK et al. Validity of self-reported height and weight in 4808 EPIC-Oxford participants. Public Health Nutr 2002;5:561–65.
- ⁴⁴ Wada K, Tamakoshi K, Tsunekawa T et al. Validity of self-reported height and weight in a Japanese workplace population. Int J Obes 2005;29:1093–99.
- Gordon I, Boffetta P, Demers PA. A case study comparing a meta-analysis and a pooled analysis of studies of sinonasal cancer among wood workers. *Epidemiology* 1998;**9:**518–24.
- ⁴⁶ Slattery ML, Curtin K, Anderson K *et al*. Associations between cigarette smoking, lifestyle factors, and microsatellite instability in colon tumors. *J Natl Cancer Inst* 2000;**92**: 1831–36.
- ⁴⁷ Smits KM, Cleven AH, Weijenberg MP *et al.* Pharmacoepigenomics in colorectal cancer: a step forward in predicting prognosis and treatment response. *Pharmacogenomics* 2008;**9**:1903–16.
- ⁴⁸ Imai K, Yamamoto H. Carcinogenesis and microsatellite instability: the interrelationship between genetics and epigenetics. *Carcinogenesis* 2008;**29**:673–80.
- ⁴⁹ Hughes LA, Khalid-de Bakker CA, Smits KM *et al*. The CpG island methylator phenotype in colorectal cancer: progress and problems. *Biochim Biophys Acta* 2012;**1825**: 77–85.

- ⁵⁰ Hinoue T, Weisenberger DJ, Lange CP *et al*. Genome-scale analysis of aberrant DNA methylation in colorectal cancer. *Genome Res* 2012;**22**:271–82.
- ⁵¹ Ogino S, Kawasaki T, Kirkner GJ et al. CpG island methylator phenotype-low (CIMP-low) in colorectal cancer: possible associations with male sex and KRAS mutations. J Mol Diagn 2006;8:582–88.
- ⁵² Yagi K, Akagi K, Hayashi H et al. Three DNA methylation epigenotypes in human colorectal cancer. Clin Cancer Res 2010:16:21–33.
- ⁵³ Dirx MJ, van den Brandt PA, Goldbohm RA et al. Energy restriction early in life and colon carcinoma risk: results of The Netherlands Cohort Study after 7.3 years of follow-up. Cancer 2003;97:46–55.
- Frankel S, Gunnell DJ, Peters TJ et al. Childhood energy intake and adult mortality from cancer: the Boyd Orr Cohort Study. BMJ 1998;316:499–504.
- 55 Hughes LA, van den Brandt PA, Goldbohm RA *et al.* Childhood and adolescent energy restriction and subsequent colorectal cancer risk: results from The Netherlands Cohort Study. *Int J Epidemiol* 2010;**39**:1333–44.
- Svensson E, Grotmol T, Hoff G et al. Trends in colorectal cancer incidence in Norway by gender and anatomic site: an age-period-cohort analysis. Eur J Cancer Prev 2002;11: 489–95.
- ⁵⁷ Svensson E, Moller B, Tretli S *et al*. Early life events and later risk of colorectal cancer: age-period-cohort modelling in the Nordic countries and Estonia. *Cancer Causes Control* 2005;**16**:215–23.
- ⁵⁸ Hughes LA, van den Brandt PA, de Bruine AP et al. Early life exposure to famine and colorectal cancer risk: a role for epigenetic mechanisms. PloS One 2009;4:e7951.

Published by Oxford University Press on behalf of the International Epidemiological Association © The Author 2012; all rights reserved. Advance Access publication 16 May 2012

International Journal of Epidemiology 2012;**41**:1072–1075 doi:10.1093/ije/dys076

Commentary: Lifestyle factors and colorectal cancer microsatellite instability—molecular pathological epidemiology science, based on unique tumour principle

Shuji Ogino^{1,2,3,4}* and Edward Giovannucci^{1,4,5,6}

¹Cancer Epidemiology Program, Dana-Farber/Harvard Cancer Center, ²Department of Pathology, Brigham and Women's Hospital, and Harvard Medical School, ³Department of Medical Oncology, Dana-Farber Cancer Institute, ⁴Department of Epidemiology, Harvard School of Public Health, ⁵Department of Nutrition, Harvard School of Public Health and ⁶Channing Laboratory, Department of Medicine, Brigham and Women's Hospital, Boston, MA, USA

*Corresponding author. Center for Molecular Oncologic Pathology, Dana-Farber Cancer Institute, Brigham and Women's Hospital, 450 Brookline Ave, Room JF-215C, Boston, MA 02215, USA. E-mail: shuji ogino@dfci.harvard.edu

Colorectal cancer encompasses fundamentally heterogeneous multifactorial diseases, 1-3 as do breast, lung and other common cancers. Each tumour is unique in terms of the tumour microenvironment, interactome and intra-tumour heterogeneity, as well as host genomic variation and lifestyle and environmental exposures. There is likely a subtle difference in the local microenvironment even in a single organ system^{4,5} or within a single tumour. This unique tumour concept is supported by technologies that have enabled reading whole genome, epigenome, transcriptome, etc. in human tumour specimens. Essentially, each tumour follows its own unique pathway of tumour evolution and progression, and we classify tumours according to similarities in molecular signatures accumulated during the carcinogenesis process.⁶ Accumulating evidence suggests that aetiological factors influence the carcinogenesis process differentially according to tumour pathway (hence, tumour classification).^{2,3} Therefore, just as different tumours respond variably to therapy, causation appears to differ by tumour subtype. However, traditional cancer epidemiology approaches (including many genome-wide association studies) have not generally taken tumour heterogeneity into consideration or analysis.

Recently, molecular pathological epidemiology (MPE) has been established as a transdisciplinary field, ^{1–3} which examines a relationship between exposures and molecular signatures in tumour, as well as interactive influences of the exposures and molecular features on cancer progression. ^{7,8} MPE is philosophically based on the concept of the uniqueness and heterogeneity of neoplastic diseases. Through MPE research, we can refine risk estimates for specific subtypes, and gain pathogenic insights on how potential aetiological factors influence different carcinogenesis pathways. ^{1–3} MPE may uncover causal associations in tumour subtypes, which had been masked when all tumours in an anatomical site were pooled together. ^{2,3}

In this issue of IJE, utilizing the MPE approach, Hughes et al.⁹ prospectively examined the relationship between anthropometric measurements and risk of colorectal cancer according to status of BRAF mutation and microsatellite instability (MSI). Notably, this study represents the first MPE study to conduct a pooled prospective analysis using geographically and operationally distinct cohort studies. One substantial challenge in MPE research is limited statistical power, because MPE research is essentially a subset analysis using tumour classification.³ The strategy of pooling multiple cohorts may potentially alleviate this issue. Hughes et al.9 successfully demonstrated that an increase in body mass index (BMI) was associated with increased risk of microsatellite stable (MSS) cancer but not of MSI-high cancer, although the risk difference was not statistically significant. These data are generally in agreement with the previous case-control studies. 10–12

Energetics and inflammation have been implicated in colorectal carcinogenesis. Obesity was associated with CpG island methylator phenotype (CIMP)-low/ negative colon cancer in a case-control study, 13 although in a case-cohort study, body size and physical activity were associated with colorectal cancer risk but not differentially by CIMP status.¹⁴ Of note, in colorectal cancer, MSI-high is associated with CIMPhigh, which is associated with BRAF mutation. 15-18 Thus, these molecular correlations can confound the apparent relationship between an exposure and a tumour variable. This 'tumour molecular confounding' is not typical confounding in an epidemiological sense because the nature of the relationships among molecular markers is not always understood. Recently, a prospective study of women showed that obesity was associated with colorectal cancer risk differentially by fatty acid synthase (FASN) expression.¹⁹ FASN has been known to be physiologically regulated by energy metabolic status. FASN is implicated in carcinogenesis, and its expression is associated with MSI-high colorectal cancer, independent of CIMP status.²⁰ The apparent relationship between obesity and MSS cancer might be due to the link between obesity and FASN-negative tumour. 19 Therefore, the interrelationship between energetics and tumour molecular features seems complex and more investigations are needed in this area.

Hughes *et al.*⁹ also showed that body height was associated with increased MSI-high (or *BRAF*-mutated) cancer risk to a significantly greater degree than MSS (or *BRAF*-wild-type) cancer risk. Body height may reflect exposure to energy metabolic status and hormonal milieu in the growth period. Interestingly, in the Netherlands Cohort Study, calorie restriction in early life might be associated with a lower risk of CIMP-high colorectal cancer.²¹ Although confirmation by independent data sets is necessary, these data suggest that energy metabolic status in early life to adolescence may influence carcinogenesis pathway that involves epigenetic instability, whereas later in life, energy metabolism is more relevant to MSS or FASN-negative tumour development.

Although MPE appears to be a promising science, ^{2,3} as a largely observational endeavour it encompasses all limitations of observational epidemiology. In addition, there are specific caveats, which have been discussed in detail elsewhere.³ We believe that MPE research should be conducted in a rigorous manner, so that findings can be generalized and appropriate public health measures can be taken based on new knowledge. To that end, we need to develop international guidelines for MPE, an extension of the STROBE (STrengthening the Reporting of OBservational Studies in Epidemiology) guidelines, which can be termed 'STROBE-MPE'.²²

For centuries, organ-based cancer classification has been useful. However, we are now geared to enter an era towards more personalized treatment in medicine. Epidemiologists should also regard each tumour as unique, and use molecular classification to better design epidemiology studies. Eventually, population cancer registries worldwide should record classification based on molecular pathogenesis and disease heterogeneity, which will accelerate the advancement of population health science. To advance the integrated interdisciplinary field of MPE, there is great need to educate epidemiologists in molecular pathology, as well as great need to educate pathologists in epidemiology and biostatistics.²² We believe that MPE can serve as a successful platform for such an interdisciplinary integration of diverse fields.

In summary, the study by Hughes *et al.*⁹ underscores the importance of the MPE approach in our quest for cancer aetiologies as well as the potential of a strategy of pooling multiple studies to overcome challenges and gain generalizable knowledge. In addition, to increase statistical power of individual population-based MPE studies, cooperation of all hospitals and pathology laboratories in provision of medical information and biospecimens is crucial. We genuinely call for collaboration and cooperation for the advancement of population science and public health.

Funding

US National Institute of Health [R01 CA151993, to S.O.; P01 CA87969, to S.E. Hankinson; P01 CA55075, to W.C. Willett].

Acknowledgements

The content is solely the responsibility of the authors and does not necessarily represent the official views of US National Institute of Health.

Conflict of interest: None declared.

References

- Ogino S, Galon J, Fuchs CS et al. Cancer immunology-analysis of host and tumor factors for personalized medicine. Nat Rev Clin Oncol 2011;8:711–19.
- ² Ogino S, Stampfer M. Lifestyle factors and microsatellite instability in colorectal cancer: the evolving field of molecular pathological epidemiology. *J Natl Cancer Inst* 2010; 102:365–67.
- ³ Ogino S, Chan AT, Fuchs CS *et al*. Molecular pathological epidemiology of colorectal neoplasia: an emerging transdisciplinary and interdisciplinary field. *Gut* 2011;**60**: 397–411.
- ⁴ Yamauchi M, Morikawa T, Kuchiba A *et al*. Assessment of colorectal cancer molecular features along bowel subsites challenges the conception of distinct dichotomy of proximal versus distal colorectum. *Gut* 2012;**61**:847–54.

- ⁵ Yamauchi M, Lochhead P, Morikawa T *et al.* Colorectal cancer: a tale of two sides or a continuum? *Gut* 2012;**61:** 794–97
- ⁶ Ogino S, Goel A. Molecular classification and correlates in colorectal cancer. *J Mol Diagn* 2008;**10**:13–27.
- Ogino S, Nosho K, Meyerhardt JA et al. Cohort study of fatty acid synthase expression and patient survival in colon cancer. J Clin Oncol 2008;26:5713–20.
- ⁸ Morikawa T, Kuchiba A, Yamauchi M *et al*. Association of CTNNB1 (beta-catenin) alterations, body mass index, and physical activity with survival in patients with colorectal cancer. *JAMA* 2011;**305**:1685–94.
- ⁹ Hughes LA, Williamson EJ, van Engeland M *et al.* Body size and risk for colorectal cancers showing BRAF mutation or microsatellite instability: a pooled analysis. *Int J Epidemiol* 2012;**41**:1060–72.
- Slattery ML, Curtin K, Anderson K et al. Associations between cigarette smoking, lifestyle factors, and microsatellite instability in colon tumors. J Natl Cancer Inst 2000;92: 1831–36.
- Satia JA, Keku T, Galanko JA et al. Diet, lifestyle, and genomic instability in the north Carolina colon cancer study. Cancer Epidemiol Biomarkers Prev 2005;14:429–36.
- ¹² Campbell PT, Jacobs ET, Ulrich CM et al. Case-control study of overweight, obesity, and colorectal cancer risk, overall and by tumor microsatellite instability status. J Natl Cancer Inst 2010;102:391–400.
- ¹³ Slattery ML, Curtin K, Sweeney C et al. Diet and lifestyle factor associations with CpG island methylator phenotype and BRAF mutations in colon cancer. Int J Cancer 2007; 120:656–63.
- ¹⁴ Hughes LA, Simons CC, van den Brandt PA *et al*. Body size, physical activity and risk of colorectal cancer with or without the CpG island methylator phenotype (CIMP). *PLoS One* 2011;**6**:e18571.
- Nosho K, Irahara N, Shima K et al. Comprehensive biostatistical analysis of CpG island methylator phenotype in colorectal cancer using a large population-based sample. PLoS One 2008;3:e3698.
- ¹⁶ Curtin K, Slattery ML, Samowitz WS. CpG island methylation in colorectal cancer: past, present and future. *Pathol Res Int* 2011;**2011**:902674.
- Hughes LA, Khalid-de Bakker CA, Smits KM et al. The CpG island methylator phenotype in colorectal cancer: progress and problems. Biochim Biophys Acta 2012;1825: 77–85.
- ¹⁸ Lao VV, Grady WM. Epigenetics and colorectal cancer. Nat Rev Gastroenterol Hepatol 2011;8:686–700.
- ¹⁹ Kuchiba A, Morikawa T, Yamauchi M et al. Body mass index and risk of colorectal cancer according to fatty acid synthase expression in the nurses' health study. J Natl Cancer Inst 2012;104:415–20.
- Ogino S, Kawasaki T, Ogawa A et al. Fatty acid synthase overexpression in colorectal cancer is associated with microsatellite instability, independent of CpG island methylator phenotype. Hum Pathol 2007;38:842–49.
- ²¹ Hughes LA, van den Brandt PA, de Bruine AP *et al.* Early life exposure to famine and colorectal cancer risk: a role for epigenetic mechanisms. PLoS One. 2009;**4:** e7951.
- ²² Ogino S, King EE, Beck AH *et al*. Interdisciplinary education to integrate pathology and epidemiology: towards molecular and population-level health science. *Am J Epidemiol* 2012; In press.