

EDITORIAL

The press, press releases, and raising unwarranted expectations

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“£30 cancer tests on the NHS within five years” was the front page headline to an article in *The Times* on 28 March 2013, raising expectations that are, unfortunately, unlikely to be fulfilled. The article was based on a study identifying DNA variations (SNPs, single nucleotide polymorphisms) in 100,000 people with cancer of the prostate, breast, or ovary and in 100,000 people without cancer.^{1–5} The article suggested that the tests would be useful in screening, stating that: “When all the [DNA] markers linked to prostate cancer were present, they increased a man’s lifetime risk of contracting the disease five-fold from about 10 per cent to 50 per cent”. The press release from Cancer Research UK⁶ that publicized the research also suggested that the findings would be useful in screening, but included a critical limitation. The press release stated that “Each (DNA) alteration raised the risk of cancer by a small amount, but the one per cent of people who have lots of these alterations could see their risk of developing prostate cancer increase to nearly 50 per cent and breast cancer to around 30 per cent”. The “one per cent” is the critical limitation; it means that the proposed SNP analysis would be a poor screening test. Although it would identify an increased likelihood of developing either cancer by about five-fold, it would do so in only a small minority (1%) of people, and most cases of the disease would arise in screen-negative individuals. This can be seen from a flow diagram (see figure), based on people who develop a hypothetical

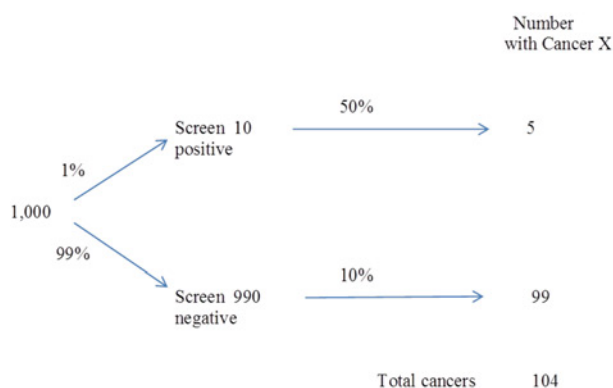
cancer X with an incidence of about 10% over a given time span, in which one percent of individuals have a five-fold increased risk relative to the remaining 99% (50% v 10%). The figure shows how 99 of the 104 cancers that arise would be missed by the SNP test although the risk of having the cancer given a positive result is high (50%).

There is no evidence that such SNP testing would be a worthwhile general screening test for any of the three cancers that were the subject of the study. Relative risks of 10 or less are unlikely to be useful in screening, though this may appear counter-intuitive. This can be seen from the risk screening convertor on <http://www.wolfson.qmul.ac.uk/rsc/>.^{7,8} The reason that etiologically important relative risks usually make poor screening tests lies in the fact that relative risk estimates focus on the risk of people in the tail of the risk distribution. It is also unlikely that the poor screening performance of a risk factor can be overcome by using the tests in sequence, or in combination with other screening markers⁹ unless the association between these markers and disease is very strong.

It has been suggested that modest improvements in programme efficacy might be achieved by using polygenic risk (so-called “risk stratification”) in addition to age to determine the target population, or to vary the frequency of screening.^{10,11} However, the potential gains are likely to be too small to justify the added complexity. A key element of a successful population screening programme is a simple system of eligibility and delivery.

The value of the study lies in showing the inherited etiology of some cases of these common cancers and that risk is influenced by many genetic variations, each with a small effect, with possible insights into the mechanisms that lead to cancer developing. The value does not lie in medical screening.

Organizations that support research need to avoid overstating the expectation of screening benefits arising from the research they support, so the press will have less opportunity to exaggerate the implications of the findings. The same concern, of course, applies to inferring harm from results showing epidemiological associations such as coffee drinking and, say, pancreatic cancer, when usually (including in this example) these associations are not causal. The limitations affecting the interpretation of research findings are often stated in the published reports, but these tend to be underestimated or ignored in the drive for publicity. They should be mentioned in press releases, together with the reference to the scientific paper concerned, so that readers can access the papers for



Percent of cancers X detected $5/104 = 5\%$
Percent of cancers X missed $99/104 = 95\%$

Odds of having cancer X given a positive result = $5:(10-5)$, or 1:1

Figure: Illustration of screening for a hypothetical cancer X with an incidence of about 10% (104/1000) over a given time span; 1% of individuals have a five-fold increased risk relative to the remaining 99% (50% v 10%).

themselves. This would help to improve the dissemination of scientific discovery to the public without raising unwarranted expectations.

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REFERENCES

- 1 Michailidou K, *et al.* Large scale genotyping identifies 41 new loci associated with breast cancer risk. *Nat Genet* 2013;**45**:353–361
- 2 Pharoah PD, *et al.* GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. *Nat Genet* 2013;**45**:362–370
- 3 Bojesen SE, *et al.* Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. *Nat Genet* 2013;**45**:371–384
- 4 Eeles RA, *et al.* Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. *Nat Genet* 2013;**45**:385–391
- 5 Garcia-Closas M, *et al.* Genome-wide association studies identify four ER negative-specific breast cancer risk loci. *Nat Genet* 2013;**45**:392–398
- 6 Press release: Study reveals the genetic variations that raise the risk of breast, prostate or ovarian cancer. 27 March 2013. <http://www.cancerresearchuk.org/cancer-info/news/archive/pressrelease/2013-03-27-genetic-variation-cancer-risk>
- 7 Wald NJ, Morris JK. Assessing risk factors as potential screening tests. A simple assessment tool. *Arch Intern Med* 2010
- 8 Wald NJ, Hackshaw AK, Frost CD. When can a risk factor be used as a worthwhile screening test? *BMJ* 1999;**319**:1562–65
- 9 Wald NJ, Morris JK, Rish S. The efficacy of combining several risk factors as a screening test. *J Med Screen* 2005;**12**:197–201
- 10 Pashayan N, Duffy SW, Chowdhury S, Dent T, Burton H, Neal DE, Easton DP, Eeles R, Pharoah P. Polygenic susceptibility to prostate and breast cancer: implications for personalised screening. *Br J Cancer* 2011;**104**:1656–1663
- 11 Burton H, Chowdhury S, Dent T, Hall A, Pashayan N, Pharoah P. Public health implications from COGS and potential for risk stratification and screening. *Nat Genet* 2013;**45**:349–351