

Hypermethylation of the 5' CpG island of the gene encoding the serine protease Testisin promotes its loss in testicular tumorigenesis

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Correction to: *British Journal of Cancer* (2005) **92**, 760–769; doi:10.1038/sj.bjc.6602373

The corresponding author of the above paper, Dr TM Antalis, has received suggestions that Figure 2A in this paper may have been manipulated. Since the samples from the experiment are no longer available, Figure 2A cannot be repeated, so should be discounted.

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Experimental data and analysis have been found providing support for other figures in this paper. Furthermore, the result of the paper, that the absence of Testisin mRNA in testicular tumour tissue is associated with aberrant methylation of the Testisin gene, was confirmed in a publication in 2006 by an unrelated laboratory (*J Cancer Res Clin Oncol*, **13**: 765–770).

Exogenous administration of protease-resistant, non-matrix-binding IGFBP-2 inhibits tumour growth in a murine model of breast cancer

C-L Soh, K McNeil, C M Owczarek, M P Hardy, L J Fabri, M Pearce, C A Delaine and B E Forbes

Correction to: *British Journal of Cancer* (2014) **110**, 2855–2864; doi:10.1038/bjc.2014.232

Upon publication of the above paper in *British Journal of Cancer*, the authors noted an error in Table 1. The relative K_D values were not correct. The corrected Table 1 is reproduced below.

Table 1. Binding affinities of WT IGFBP-2 and mutants

	IGF-I			IGF-II		
	K_D (nM)	s.e.m.	Relative K_D	K_D (nM)	s.e.m.	Relative K_D
WT	1.38	0.13	1.00	1.37	0.18	1.00
PR	3.74	0.50	2.71 ^a	1.15	0.12	0.84 ^a
PR/NMB	3.16	0.38	2.29 ^a	2.17	0.07	1.58 ^a

K_D = dissociation constant (k_d/k_a , dissociation rate/association rate).
^aNot statistically significant.
 Relative K_D = K_D mutant/ K_D WT.

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Ethnicity, deprivation and screening: survival from breast cancer among screening-eligible women in the West Midlands diagnosed from 1989 to 2011

M Morris, L M Woods, N Rogers, E O'Sullivan, O Kearins and B Rachet

Correction to: *British Journal of Cancer* (2015) **113**, 548–555; doi:10.1038/bjc.2015.204

It has been brought to our attention that there are errors in the paragraph 'Assigning deprivation score' on page 549 in the Materials and Methods section of the above paper published in volume 113, issue 3 (28 July 2015). The errors have now been corrected and the amended version of the paragraph in question reads as follows:

'Two different ecological measures of deprivation were used according to the study period: a score based on Townsend's system (Townsend *et al*,

1988) for women diagnosed with a breast cancer between 1989 and 2003 (40% of cases); the 2004, 2007 and 2010 scores of the Income Domain of the Index of Multiple Deprivation (IMD) (21.4%, 26.5% and 12.2%, respectively) for women diagnosed from 2004 onwards. The women were assigned a score according to their geographical area of residence at the time of their diagnosis (Communities and Local Government, 2008; Communities and Local Government, 2011; Neighbourhood Renewal Unit, 2004). Both the Townsend and IMD scores used Lower Super Output Areas (LSOAs, approximately 1500 people). The scores were split into five categories based on the quintiles of the national distribution of the areas.'

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HLA class I is most tightly linked to levels of tapasin compared with other antigen-processing proteins in glioblastoma

Camilla Thuring, Elna Follin, Linda Geironson, Eva Freyhult, Victoria Junghans, Mikkel Harndahl, Søren Buus and Kajsa M Paulsson

Correction to: *British Journal of Cancer* (2015) **113**, 952–962; doi:10.1038/bjc.2015.297

It has been brought to our attention that there is an error in the title of the above paper published in advance online. The original title was

'The environment HLA class I is most tightly linked to levels of tapasin compared with other antigen-processing proteins in glioblastoma'. This has been corrected above and in volume 113, issue 6 in which it appears.

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