

Correction

Correction: *BCoR-L1* variation and breast cancer

Felicity Lose^{1,2}, Jeremy Arnold¹, David B Young¹, Carolyn J Brown³, Graham J Mann⁴,
Gulietta M Pupo⁴, The Kathleen Cuninghame Foundation Consortium for Research into Familial
Breast Cancer, Kum Kum Khanna¹, Georgia Chenevix-Trench¹ and Amanda B Spurdle¹

¹Cancer and Cell Biology Division, Queensland Institute of Medical Research, 300 Herston Road, Brisbane, Queensland, Australia, 4006

²School of Medicine, Central Clinical Division, University of Queensland, Royal Brisbane Hospital, Corner Butterfield Street and Bowen Bridge Road, Brisbane, Queensland, Australia, 4029

³Department of Medical Genetics, Molecular Epigenetics Group, University of British Columbia, 2329 West Mall, Vancouver, BC, Canada, V6T 1Z4

⁴Westmead Institute for Cancer Research, University of Sydney at Westmead Millennium Institute, Westmead Hospital, Darcy Road, Westmead, New South Wales, Australia, 2145

Corresponding author: Amanda B Spurdle, Amanda.Spurdle@qimr.edu.au

Published: 21 October 2008

This article is online at <http://breast-cancer-research.com/content/10/5/406>

© 2008 BioMed Central Ltd

Breast Cancer Research 2008, **10**:406 (doi:10.1186/bcr2153)

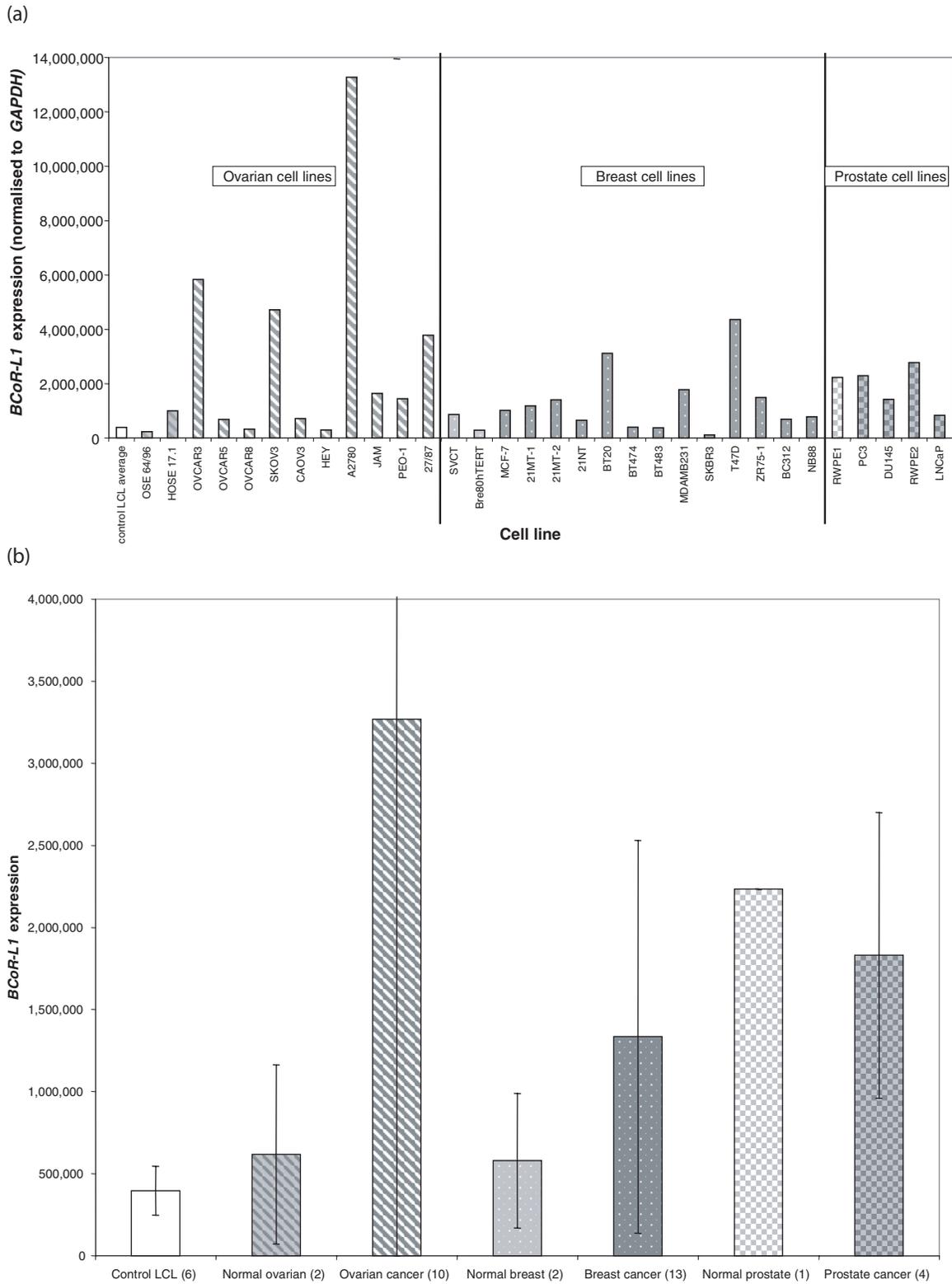
Following the publication of our article [1] we noticed that, due to a production error, the figure legends and images were incorrectly matched. All legends were correctly placed, and cited in the text, but were associated with the wrong image.

The figures should therefore appear in the order shown in this correction.

Reference

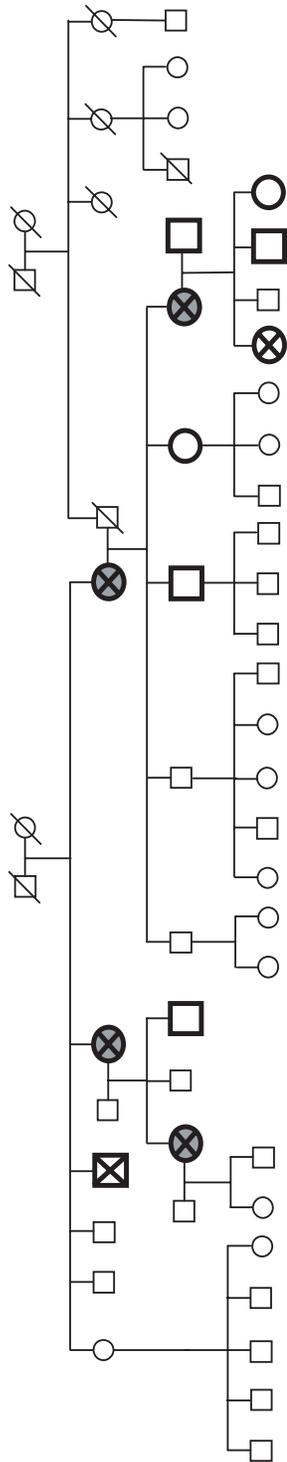
1. Lose F, Arnold J, Young DB, Brown CJ, Mann GJ, Pupo GM, The Kathleen Cuninghame Foundation Consortium for Research into Familial Breast Cancer, Khanna KK, Chenevix-Trench G, Spurdle AB: **BcoR-L1 variation and breast cancer.** *Breast Cancer Res* 2007, **9**:R54.

Figure 1



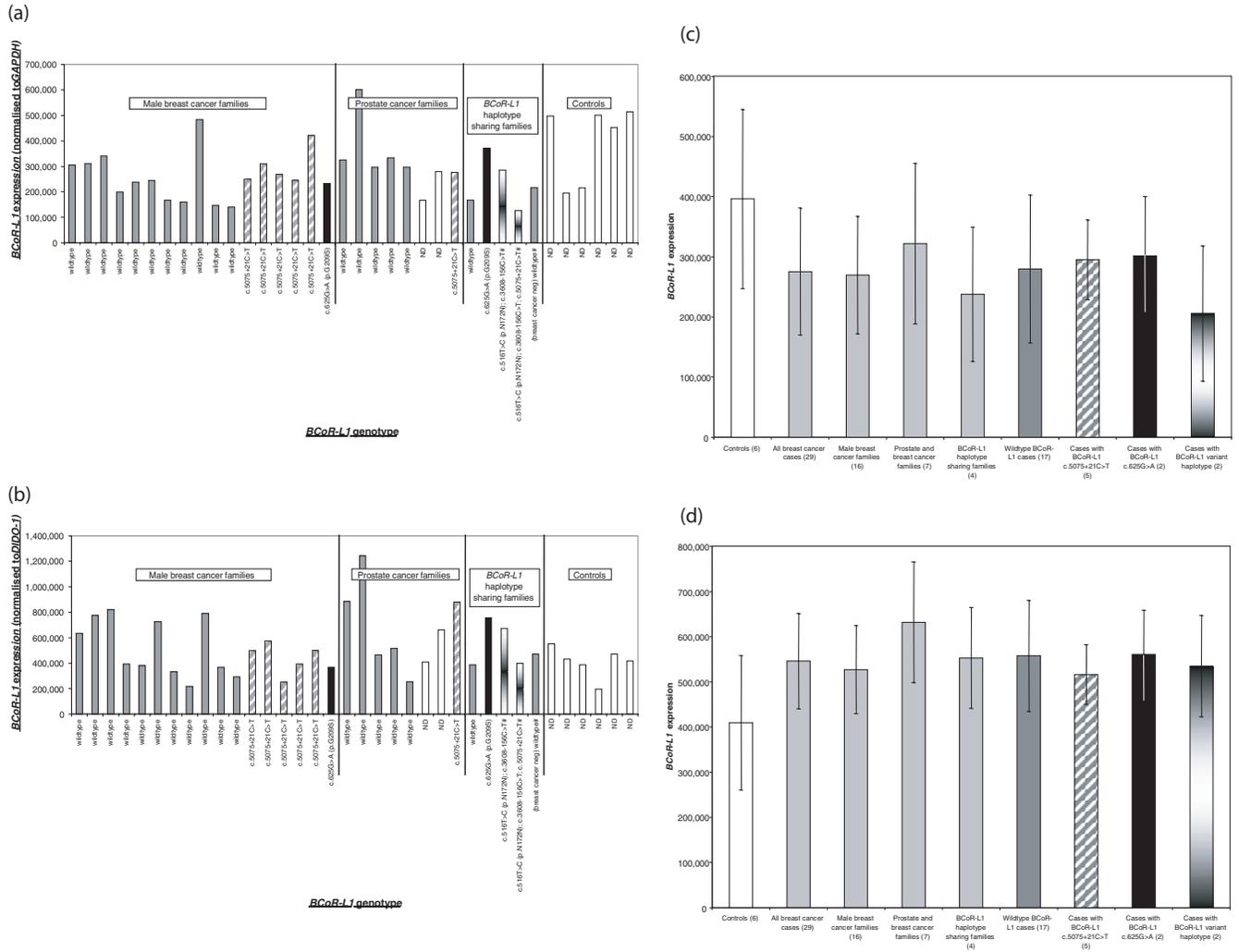
BCoR-L1 expression in cancer and normal cell lines. (a) BCoR-L1 expression in cancer and normal cell lines. **(b)** Mean and standard deviation of BCoR-L1 expression in cancer and normal cell lines. Normal cell lines: ovarian – OSE 64/96, HOSE 17.1; breast – SVCT, Bre80hTERT; prostate – RWPE1. BCoR-L1, BCL6 corepressor-like 1; GAPDH, glyceraldehyde-3-phosphate dehydrogenase.

Figure 2



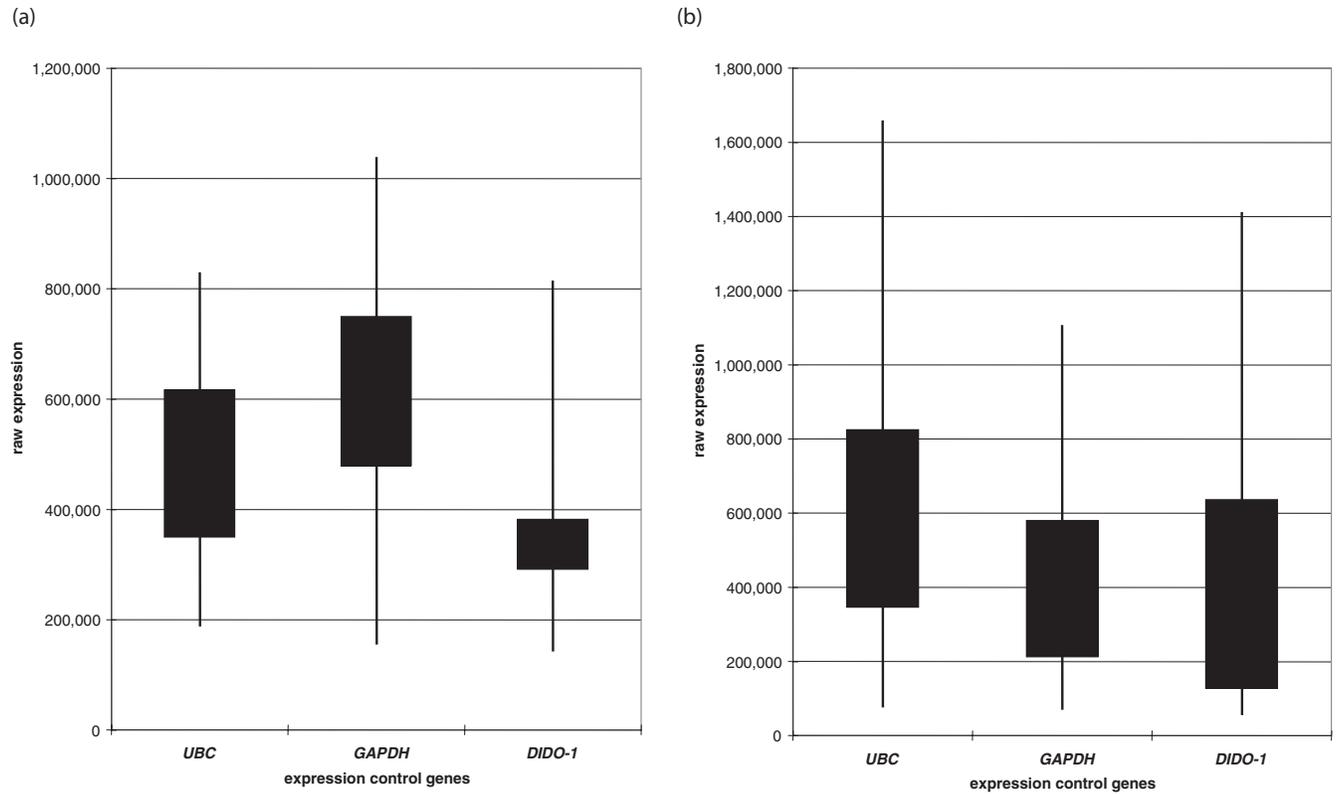
***BCoR-L1* haplotype sharing family pedigree detailing carriers of the c.516T>C and c.3608-156C>T variants.** ● = breast cancer-positive; c.516T>C and c.3608-156C>T-positive. ⊗ = breast cancer-negative; c.516T>C and c.3608-156C>T-positive. ■ = breast cancer-negative; c.516T>C and c.3608-156C>T-negative. ○ = female, □ = male; subjects marked by small shapes were not available for genotyping. *BCoR-L1*, *BCL6* corepressor-like 1.

Figure 3



BcoR-L1 expression in lymphoblastoid cell lines (LCLs) from breast cancer families. (a) *BCoR-L1* expression in LCLs from breast cancer families (normalised to *GAPDH*). **(b)** *BCoR-L1* expression in LCLs from breast cancer families (normalised to *DIDO-1*). **(c)** Mean and standard deviation of *BCoR-L1* expression in samples, grouped according to type of family cancer or *BCoR-L1* genotype (normalised to *GAPDH*). **(d)** Mean and standard deviation of *BCoR-L1* expression in samples, grouped according to type of family cancer or *BCoR-L1* genotype (normalised to *DIDO-1*). *Subject also carries a *BRCA2* mutation. #Subjects from the same *BCoR-L1* haplotype sharing family. *BCoR-L1*, *BCL6* corepressor-like 1; *DIDO-1*, death inducer-obliterator 1; *GAPDH*, glyceraldehyde-3-phosphate dehydrogenase.

Figure 4



Variation in control gene expression. (a) Variation in control gene expression in lymphoblastoid cell lines. **(b)** Variation in control gene expression in cell lines. *DIDO-1*, death inducer-obliterator 1; *GAPDH*, glyceraldehyde-3-phosphate dehydrogenase; *UBC*, ubiquitin C.