Title: Effect of BRCA2 sequence variants predicted to disrupt exonic splice enhancers on BRCA2 transcripts

Authors:

Phillip J Whiley (philip.whiley@qimr.edu.au)
Christopher A Pettigrew (christopher.pettigrew@ucc.ie)
Brooke L Brewster (b.brewster@uq.edu.au)
Logan C Walker (logan.walker@qimr.edu.au)
kConFab Investigators (heather.thorne@petermac.org)
Amanda B Spurdle (amanda.spurdle@qimr.edu.au)
Melissa A Brown (melissa.brown@uq.edu.au)

Version: 3 Date: 3 May 2010

Author's response to reviews: see over
30th April, 2010.

Scott Edmunds PhD
The BioMed Central Editorial Team
Tel: +44 (0) 20 3192 2013
e-mail: editorial@biomedcentral.com
Web: http://www.biomedcentral.com/

Dear Dr Edmunds,

Re: MS: 7672096462965717
Functional analysis of BRCA2 sequence variants predicted to disrupt exonic splice enhancers.
Phillip J Whiley, Christopher A Pettigrew, Brooke L Brewster, Logan C Walker, kConFab Investigators, Amanda B Spurdle and Melissa A Brown

Thank you for your assessment of the revised version of the above manuscript. We appreciate the helpful comments of the reviewers and are delighted to submit a re-revised manuscript, addressing the remaining minor comment, requesting that we indicate we have used HGVS nomenclature and cite the den Dunnen et al, 2001 paper.

My sincerest apologies for the delay in resubmitting this manuscript. I had some unanticipated surgery in March, which delayed progress.

We hope that the revised manuscript will be to your satisfaction and that the manuscript will now be suitable for publication in BMC Medical Genetics.

Yours sincerely,
Melissa Brown, PhD.