Erratum to: Genome-based expression profiles study for the pathogenesis of pelvic organ prolapse: an array of 33 genes model

Ling-Hong Tseng · Ilene Chen · Yi-Hao Lin ·
Ming-Yang Chen · Tsia-Shu Lo · Chyi-Long Lee

Published online: 4 March 2010
© The International Urogynecological Association 2010

Erratum to: Int Urogynecol J
DOI 10.1007/s00192-009-0990-y

The authors would like to clarify that they did not perform the microarray experiments themselves at Chang Gung Memorial Hospital. Rather, the source of data was the publicly accessible Gene Expression Omnibus (GEO), and the dataset is deposited in GEO with the accession number GSE12852. The description of RNA extraction and microarray experiments was merely intended to help readers quickly comprehend the laboratory procedures. The authors are grateful to the original researchers (Brizzolara SS et al., *Mol Hum Reprod* 2009;15:59—67) for the use of their deposited microarray datasets in this paper.

The online version of the original article can be found at <http://dx.doi.org/10.1007/s00192-009-0990-y>.

L.-H. Tseng · Y.-H. Lin · T.-S. Lo · C.-L. Lee (✉)
Department of Obstetrics and Gynecology,
Chang Gung Memorial Hospital and University
of Chang Gung School of Medicine,
5 Fu-Hsing Street, Kwei-Shan,
Tao-Yuan 333, Taiwan
e-mail: 3g7330@yahoo.com.tw

I. Chen
School of Electrical and Information Engineering,
University of Sydney,
Sydney, Australia

M.-Y. Chen
Department of Electrical Engineering, Stanford University,
Stanford, CA, USA