

CORRECTION

## Correction: Consensus Analysis of Whole Transcriptome Profiles from Two Breast Cancer Patient Cohorts Reveals Long Non-Coding RNAs Associated with Intrinsic Subtype and the Tumour Microenvironment

James R. Bradford, Angela Cox, Philip Bernard, Nicola J. Camp

There is information missing from the Competing interests statement. The Competing interests statement should read: Philip Bernard is an inventor of the PAM50 signature and a stakeholder in BioClassifier LLC, a company that licensed the PAM50 know-how to Nanostring Inc. for commercialization of Prosigna®. PB also has been a Medical Director at ARUP Laboratories Inc since 2003. This does not alter our adherence to PLOS ONE policies on sharing data and materials.

## Reference

 Bradford JR, Cox A, Bernard P, Camp NJ (2016) Consensus Analysis of Whole Transcriptome Profiles from Two Breast Cancer Patient Cohorts Reveals Long Non-Coding RNAs Associated with Intrinsic Subtype and the Tumour Microenvironment. PLoS ONE 11(9): e0163238. https://doi.org/10.1371/ journal.pone.0163238 PMID: 27685983



## GOPEN ACCESS

Citation: Bradford JR, Cox A, Bernard P, Camp NJ (2018) Correction: Consensus Analysis of Whole Transcriptome Profiles from Two Breast Cancer Patient Cohorts Reveals Long Non-Coding RNAs Associated with Intrinsic Subtype and the Tumour Microenvironment. PLoS ONE 13(2): e0192589. https://doi.org/10.1371/journal.pone.0192589

Published: February 6, 2018

Copyright: © 2018 Bradford et al. This is an open access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.