Genome Research 22: 947–956 (2012)

Genome-wide determination of RNA stability reveals hundreds of short-lived noncoding transcripts in mammals
Hidenori Tani, Rena Mizutani, Kazi Abdus Salam, Keiko Tano, Kenichi Ijiri, Ai Wakamatsu, Takao Isogai, Yutaka Suzuki, and Nobuyoshi Akimitsu

In the above-mentioned paper, all citations of Supplemental Tables S16 and S17 (three instances on p. 954) should have been cited as Supplemental Table S15 instead.

Genome Research 22: 1006–1014 (2012)

BRAFV600E remodels the melanocyte transcriptome and induces BANCR to regulate melanoma cell migration
Ross J. Flockhart, Dan E. Webster, Kun Qu, Nicholas Mascarenhas, Joanna Kovalski, Markus Kretz, and Paul A. Khavari

The BANCR gene noted in the abstract of this article should have been identified as BRAF-activated non-coding RNA (rather than BRAF-regulated IncRNA 1). The designation was correctly identified elsewhere in the article.