



Correction for Lecona et al., “USP7 Cooperates with SCML2 To Regulate the Activity of PRC1”

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Volume 35, no. 7, p. 1157–1168, 2015, <https://doi.org/10.1128/MCB.01197-14>. The wrong data set was used in analysis for Fig. 2; therefore, several changes are needed to the figure and text as noted below. These changes do not affect any of the conclusions of the work.

Page 1159, right column, line 4 from bottom: “cells and” should read “cells with a partial knockdown of endogenous SCML2 and rescue with overexpression of SCML2A (12) and.”

Page 1159, left column, “ChIP-seq analysis,” lines 7–end: “<math><10^{10}</math>...<math><5\text{ kb}</math>” should read “<math><10^5</math>. SCML2 ERs were restricted to those that lie within 5 kb of an annotated TSS, and SCML2 ERs showing significant signal in the IgG control were further filtered from our heat map analysis.”

Page 1159, left column: The following paragraph should be added before Results. “**Accession number(s)**. The GEO number for the ChIP-seq data of USP7 is GSE61048, and that for the rest of ChIP-seq data is GSE38275.”

Page 1161, Fig. 2A to C: There was a mistake in the data sets used to generate these images, and one of the panels was inadvertently duplicated in panel A. The corrected panels appear below.

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