CORRIGENDUM

The epigenetic regulator UHRF1 promotes ubiquitination-mediated degradation of the tumor-suppressor protein promyelocytic leukemia protein

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Since the publication of the above article the authors have identified an oversight error in the migration assays shown in Figure 8a (siU-1 and siU-2, 0 h), in which two images were

duplicated due to a mistake during transfer of a file from Photoshop to Canvas.

The amended figure is shown below.

The authors wish to apologise for any inconvenience caused.

Figure 8. Knockdown of UHRF1 inhibits HUVEC migration. **(a)** Representative images of wound-healing assays. HUVECs were transiently transfected with a nontargeting or two different UHRF1 siRNAs. At 60 h after transfection, wound-healing assays were performed as described in the Materials and methods. **(b)** Statistical analyses of wound-healing assays. **(c)** UHRF1 is efficiently knocked down by siRNAs. Whole-cell extracts were prepared from HUVECs 60 h after siRNA transfection. Immunoblotting was performed to determine UHRF1 knockdown efficiency. Wound-healing assays were performed as described in the Materials and methods. Unpaired two-tail *t*-tests ($*P < 0.01$) were used for statistical analyses.