

CORRIGENDUM

p16^{INK4A} Represses the paracrine tumor-promoting effects of breast stromal fibroblasts

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Correction to: *Oncogene* (2013) 32, 2356–2364; doi:10.1038/onc.2012.270; published online 2 July 2012

The authors of this study have noticed an error in Figure 3a (top right panel, the sixteenth page) of their paper. The authors inadvertently duplicated the panel from the left column. The error

was unintentionally introduced during figure preparation and is not a correct representation of the data. The authors have provided a corrected Figure 3a, and it is reproduced here in its entirety. The error in the original published manuscript does not in any way alter the results or conclusions of this study. The authors apologize that these errors were not detected earlier.

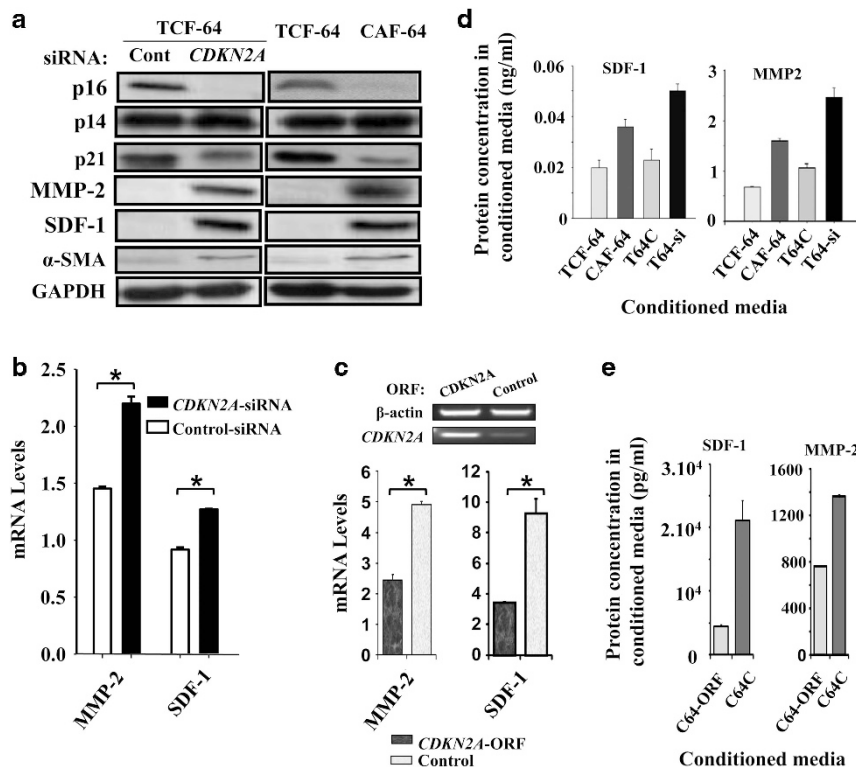


Figure 3. p16 Suppresses the expression of α -SMA, SDF1 and MMP 2. **(a)** Whole-cell lysates were prepared from TCF/CAF-64 and TCF-64 expressing either *CDKN2A*-siRNA or control-siRNA, and were used for immunoblot analysis. **(b)** Total RNA was extracted and the amount of *MMP-2* and *SDF-1* mRNAs were assessed using real-time RT-PCR. Error bars represent means \pm s.d. **P*-value < 0.05. **(c)** Total RNA was extracted from CAF-64 cells expressing either *CDKN2A* ORF or control. Upper panel: the *CDKN2A* mRNA was amplified by RT-PCR, and the obtained fragments were electrophoresed on ethidium bromide stained agarose gel. Lower panel: the amount of *MMP-2* and *SDF-1* mRNAs were assessed using real-time RT-PCR. Error bars represent means \pm s.d. **P*-value < 0.05. **(d, e)** CM from the indicated cells were collected after 24 h and the levels of the indicated proteins were determined by ELISA and were presented in the respective histograms. Error bars represent means \pm s.d.