HGF upregulation contributes to angiogenesis in mice with keratinocyte-specific *Smad2* deletion

Kristina E. Hoot, …, Qinghong Zhang, Xiao-Jing Wang


HGF levels were determined using Power SYBR Green Master Mix (Applied Biosystems) and the following custom primers: 5ʹ-AGGAACAGGGGCTTTACGTT-3ʹ (forward) and 5ʹ-GTCAAATTCATGGCCAAACC -3ʹ (reverse). Human HGF quantitative PCR assays were performed using the TaqMan probe set Hs00300159_m1 The authors regret the error.

Find the latest version:

https://jci.me/69077/pdf
Corrigendum

Hypoxia-inducible factor–dependent breast cancer–mesenchymal stem cell bidirectional signaling promotes metastasis

Pallavi Chaturvedi, Daniele M. Gilkes, Carmen Chak Lui Wong, Kshitiz, Weibo Luo, Huafeng Zhang, Hong Wei, Naoharu Takano, Luana Schito, Andre Levchenko, and Gregg L. Semenza


Citation for this corrigendum: J Clin Invest. 2013;123(3):1402. doi:10.1172/JCI69244.

Figure 5C was incorrect. The correct figure is below.

The authors regret the error.

Corrigendum

HGF upregulation contributes to angiogenesis in mice with keratinocyte-specific Smad2 deletion

Kristina E. Hoot, Masako Oka, Gangwen Han, Erwin Bottinger, Qinghong Zhang, and Xiao-Jing Wang


Citation for this corrigendum: J Clin Invest. 2013;123(3):1402. doi:10.1172/JCI69077.

In the Methods section, the authors inadvertently provided incorrect quantitative PCR primer sequences for amplification of mouse and human HGF. The correct passage appears below.

HGF levels were determined using Power SYBR Green Master Mix (Applied Biosystems) and the following custom primers: 5’-AGGAACAGGGCTTTACGTT-3’ (forward) and 5’-GTCAAATTCATGGCCAAACC-3’ (reverse). Human HGF quantitative PCR assays were performed using the TaqMan probe set Hs00300159_m1.

The authors regret the error.