

CORRECTIONS & AMENDMENTS

CORRIGENDUM

doi:10.1038/nature25991

Corrigendum: Cloche is a bHLH-PAS transcription factor that drives haemato-vascular specification

Sven Reischauer, Oliver A. Stone, Alethia Villasenor, Neil Chi, Suk-Won Jin, Marcel Martin, Miler T. Lee, Nana Fukuda, Michele Marass, Alec Witty, Ian Fiddes, Taiyi Kuo, Won-Suk Chung, Sherveen Salek, Robert Lerrigo, Jessica Alsiö, Shujun Luo, Dominika Tworus, Sruthy M. Augustine, Sophie Muceniaks, Björn Nystedt, Antonio J. Giraldez, Gary P. Schroth, Olov Andersson & Didier Y. R. Stainier

Nature **535**, 294–298 (2016); doi:10.1038/nature18614

In the Author Information section of this Letter, LT571435 from the SRA was provided as the accession number for the PacBio whole-genome sequencing data. Immediately after publication, we noted that LT571435 is instead the GenBank entry for the assembled genomic sequence of the *cloche* locus and this information is now included in the legend of Extended Data Fig. 2. The correct accession number for the PacBio whole-genome sequencing data, available from the European Nucleotide Archive, is PRJEB13442. The original Letter has been corrected online.