

Figure S1. Sequence alignment between mouse and human HS-E1

Mouse (top line) and human (bottom line) genomic DNA sequences from the region to which HS-E1 maps in both species were aligned using Pipmaker (<http://pipmaker.bx.psu.edu/pipmaker/>).²² Coordinates for the mouse sequence refer to the transcription start site for the mouse $\epsilon\gamma$ -globin gene, in which the first A in the sequence “GTACGTA~~CTT~~GCTTCTG” is defined as +1. Coordinates for the human sequence refer to the transcription start site for the human ϵ -globin gene, in which the first A in the sequence “GCACATATCTGCTTCCG” is defined as +1.

Figure S2. Blood cytopins comparing wild type and mutant peripheral blood from E12.5 litter mates

Representative Wright stain microscope fields from two wild type (WT) and two mutant (Δ HS-E1) littermates show no gross differences in red cell maturation and morphology. Cells were cytopun (Shandon cytopsin II, Thermo-Scientific, Waltham, MA) on plain glass slides, Wright stained (Sigma Aldrich WSHT), and mounted with Permount. Images were acquired on a Nikon DS-Fi1 camera on a Nikon Eclipse 80i microscope (Nikon, Chiyoda-ku, Tokyo, Japan) using Nikon Plan Fluor objective lens (Nikon, 20 \times magnification, 0.5 NA). Images were processed by Nikon Elements software for white balance and brightness. The scale bars in the lower right corners of each panel indicate 50 μ m.

Figure S1

Mouse 7569 TAAAGAATCAGCAGTCACAGGATTTCTGACTCAAGAAATGGTATTACTAAGAAAAATATTCTT AAGGATCCAATTT
Human 2427 TAAACGATAAGCA TTAGGATGTTAAGTGACTCAGGAAATAAGATTGTTGGAAAAAATAATCTGCTTATGTGCACAAAATGGATTCAGTTTGCAGAT

Mouse 7645 AACTTAAATGCTTAGGAATCAATTACAG AGATATAAAAAGTTTAAAATGAAGAGGTGAAGCAAGGC ATGCAACCCGAGGTAATGTTCAAAGGTGGG
Human 2524 AAAATAAAATA TGATGATGATGATTCAAAGGACAGATACAATGGTTCAAACCAAGAGGAGCAGTGAGTCTGTGAAATTTGAAAGGATGACAAAGGTGGG

Mouse 7739 GTGGGGGAGCAGCGT CAGATTAAC TCTAATATATGAGAAGAAACAAGA CAGTAGAAGA TGAAATCAAAAATAC TGAGTGGACATAACTACAT
Human 2624 GTGAGAAA GACATAGTATTCGACCTGACTGTGGGAGATGAGAAGGAAGAAGGAGGTGATTAATGACTGAAAGCTCCGACAGTG GTGAAGA TAACAGGAG

Mouse 7833 AAAG TGTACATTGACTTGGA AAACTCATGTGGGTAAAA CACACTTGAGAGAATTTAAAGGTGCTCAAAGAAATTC
Human 2723 GAAACCATGCAC TTGACCCTGGTGACTCTCATGTGTGAAGGGTAGAGGGATATTAACAGATT TAC TTTTAGGAAAGTCTAGATTGGTCAAGGAGTTT

Mouse 7908 AAGCC CTAGTCAATCTCTTGTCAAGGCTGGTGATCTTTAAGTGGTAGAGGTAAGAGAACCACATTTTAACTAATA AACTATATTCCCTCATCC
Human 2822 GACCTTCAGGCTTGTGTCTTCATATCAAGG AACCTTTCATTTTCCAAGTTAGAGTGCCATATTTTGGCAAATAA ACTTTATTAG

Mouse 8001 TGATTCATAG ATTGGTTAGAGTATGACTGGTGGATTGTATTTAAGTTGCTTTGTCCACTTTAAATAAG AAAGAACAATGATCT
Human 2910 TAA TTTTATAGTGCTCTCACATTGATCAGACTTTTCTGTGAATTACTTTTGAATTTGGCTGTATATATCCAGAAATGAGGAGAGACAAATAATAT

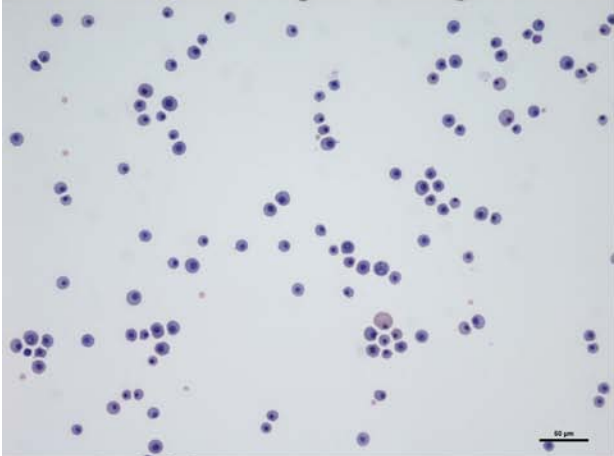
Mouse 8088 TGTAGTTT CAGGCATCAACAACATTGGCCTTTTGGTCTACAGCATTTCATATG GCAA TAAACAGGAAAACAACC AAACCAAACAACAATAAC
Human 3010 TGTAGTTG CAGGCATCAACAATCTGGTCTCTCTGAGCCTTATAA CCTTTCAATA TGCCCA TAAACAGAGTAA

Mouse 8185 AACAAAAATG CAGGTTGTATATGACACTCAAATAGTTTCTCTGGACCTGTC AAGAGAT GAGACAATGCCAATTTTT ATAAA TTTCTTTTCC
Human 3085 ACAGGGA TTTATTCA TGCCACTAAA TATTTTCACTAGTCACTACAAAATGAGGCAATGTGCAATTTTGTATACATATTTTATA

Mouse 8283 CTTATG
Human 3174 TTTATG

Figure S2

WT



Δ HS-E1

