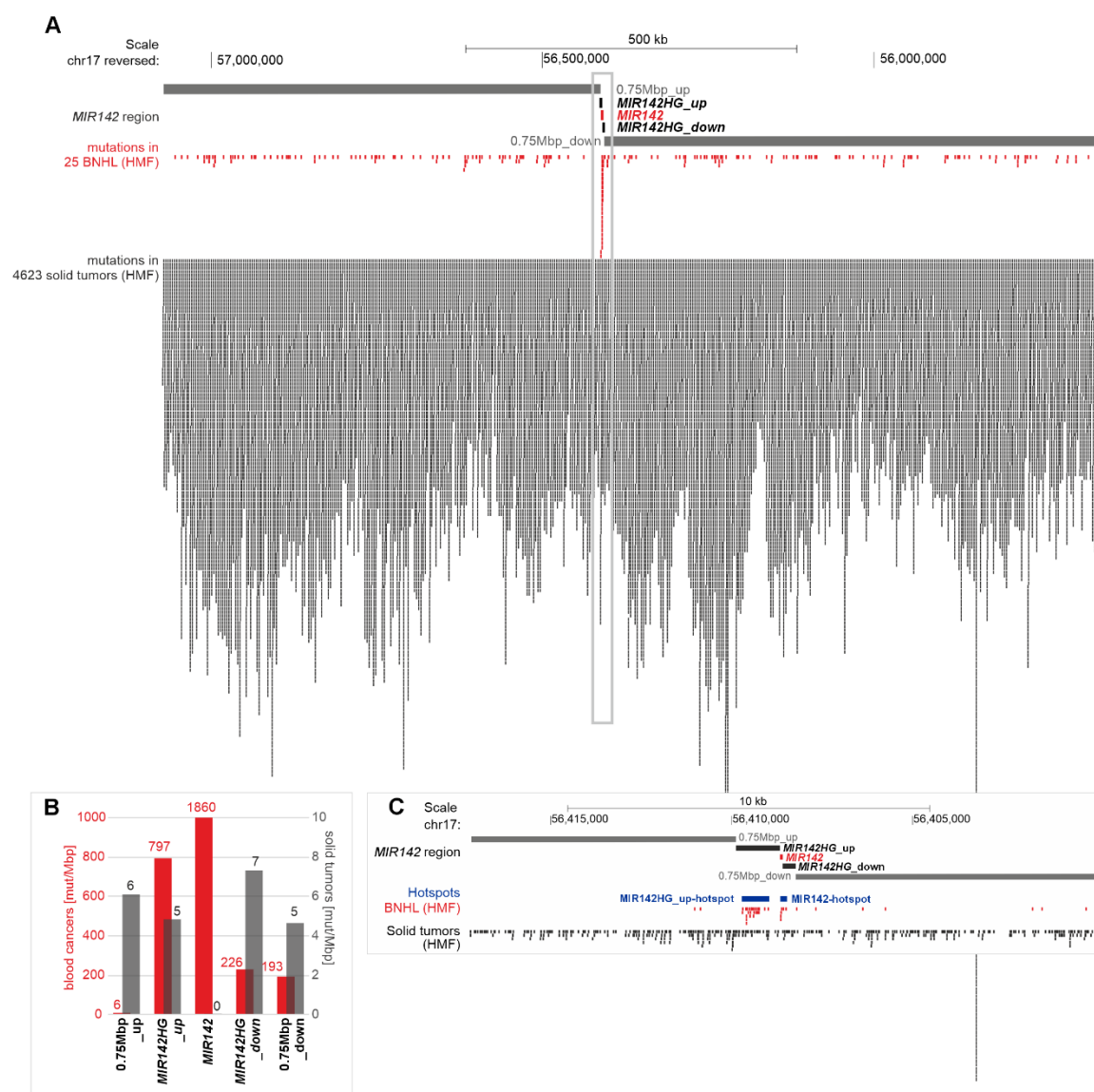




Supplementary Figure S1. **A)** Map of somatic mutations identified in HMF-deposited blood cancer (BNHLs, N=25) and solid tumor (N=4,623) samples (visualized with the UCSC Genome Browser). **B)** Graph showing mutation density. **C)** Zoomed-in view of the MIR142HG region. The overall figure schemes are as in Figures 1C, 1D, and 1E, respectively.