



Fig. S1. Putative RUNX1 and FLI1 binding motifs in 5'UTR of ANKRD26 are not conserved in mouse and zebrafish. **A**, Schematic representation of 5'UTR of human *ANKRD26* gene indicates RUNX1 (red) and FLI1 (green) binding sequences. **B**, Sequence alignment of 5'UTR of *Ankrd26* in human (NM_014915), mouse (NM_001081112) and zebrafish (XM_021470586). RUNX1 and Fli1 binding sites in human *ANKRD26* are indicated by red and green box, respectively. Black box indicates the 2 nucleotides deletion mutation created in the zebrafish *ankrd26* gene. Dots represent same nucleotides and dashes represent nucleotide gaps among species.

Table S1. Proteomic analysis of *ankrd26*^{ku6} and wt young thrombocytes.

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Table S2. Proteomic analysis of *ankrd26*^{ku6} and wt mature thrombocytes.

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