**Supplemental Table 1.** Hematological indices of the proband heterozygous for PIEZO1 V598M (II:1) and an unaffected family member (I:1), from 2016. Some of these indices are more severe than usually encountered in HX patients with other causative PIEZO1 mutations (Andolfo et al., Am J Hematol 2018; 93:107-121). RR, reference range.

**Supplemental Figure 1.** Proband's pedigree. The sporadic heterozygous PIEZO1 mutation V598M present in proband is absent from other family members. The common variant  $\Delta$ K1878 previously proposed as pathogenic for HX does not cosegregate with symptomatic disease in this family, and was not reported in the French HX patient heterozygous for PIEZO1 V598M [5].

**Supplemental Figure 2**. **A**. Whole blood total RNA prepared from the proband and unaffected family members was reverse-transcribed to cDNA, PCR-amplified and Sanger-sequenced. The heterozygous PIEZO1 variant c.1792G>A encoding missense mutant 598V>M was detected in cDNA from the proband (II:1) only, a finding confirmed in genomic DNA (not shown). **B.** cDNA sequencing trace of the same stretch of PIEZO1 sequence from one unaffected family member (II:2) representative of that sequence in four additional unaffected family members (not shown). The sequence was confirmed in genomic DNA (not shown). **C.** Amino acid sequence alignment showing evolutionary conservation of human PIEZO1 residue V598 in PIEZO1 from mammals, birds, amphibia and fish, and in human PIEZO2.

**Supplemental Figure 3.** Peripheral blood smears (Wright Giemsa) of (**A**) proband (II:1) and her (**B**) Mother (I:2), from 2008.

**Supplemental Figure 4.** Ektacytometry scans from 2008 of red cells from proband (II.1, red), mother (I:1, purple), father (I:2, green) and a healthy control (blue), on a Lorca ektacytometer lacking capability of continuous osmotic gradient generation

(RR Mechatronics, Netherlands). **A.** Deformability at 400 mOsm and at discrete shear stress values. **B**. Deformability at 16.86 Pa and at discrete osmolarity values.

## Supplemental Table 1.

## Hematological indices of PIEZO1 V598M and WT erythrocytes.

Mature Erythrocytes	WT	V598M	RR		
Red Blood Cells (X 10 <sup>6</sup> cells/uL)	4.30	2.86	3.75-5.07		
Hematocrit (%)	40.8	32.8	31.7-43.4		
Hemoglobin (g/dL)	12.4	9.9	11.4-15.1		
Mean Cellular Volume (fL)	94.9	114.8	82.7-90.2		
MCHC (g/dL)	30.4	30.1	33.6-35.3		
CHCM (g/dL)	29.3	29.5	32.2-37.7		
RDW (%)	13.3	18.1	12-15*		
HDW (g/dL)	1.96	3.44	1.9-3.0*		
% Hyperchromic cells	0.0	0.1	-		
Reticulocytes					
Reticulocytes (X 10 <sup>6</sup> cells/uL)	0.038	0.424	0.0406-0.140		
Retics (%)	0.87	14.82	0.8-2		
Mean Cellular Volume (fL)	102.8	118.7	93-121*		
CHCM (g/dL)	25.1	27.5	27-34*		
RDW (%)	9	11.7	-		
HDW (g/dL)	2.33	3.28	-		
% Hyperchromic reticulocytes	0.0	0.2	-		

Supplemental Figure 1







Figure 2

C		*	
Human PIEZO2	685	KYWIYVCGGMFFFVSFEGKI <b>V</b> MYKIIYMVLFLFCVALYQVHY	726
		KYWIYVC GMF VSF G++V+YKI+YM LFL C+ L+QV+Y	
Human PIEZO1	578	KYWIYVCAGMFIVVSFAGRL <b>V</b> VYKIVYMFLFLLCLTLFQVYY	619
M. musculus	583	KYWIYVCAGMFIVVSFAGRL <b>V</b> VYKIVYMFLFLLCLTLFQVYY	624
C. porcellus	586	KYWIYVCAGMFIVVSFAGRL <b>V</b> VYKIVYMFLFLLCLTLFQVYY	627
C. lupus	590	KYWIYVCAGMFIVVSFAGRL <b>V</b> VYKIVYMLLFLLCLTLFQVYY	631
B. taurus	383	KYWIYVCAGMFIVVSFAGRL <b>V</b> VYKIVYMLLFLLCLILFQVYY	424
G. gallus	566	KYWICVCAGMFIVVSFAGRL <b>V</b> VYKIVYMFLFLLCLTLFQVYY	607
X. laevis	584	KYWIYVCGGMFIVVSFAGRL <b>V</b> VYKIVYMLLFLLCLVSFQVYY	625
D. rerio	597	KYWIYVCGGMFIMVSFAGKL <b>V</b> AYKIVYMLLFLLCMCLYQVYY	638
0. niloticus	602	KYWIYVCGGMFIMVSFAGKL <b>V</b> AYKIIYMLLFLLCMCLNQVYY	643

Supplemental Figure 3



## Supplemental Figure 4



Control subject Proband (II:1) Unaffected Father I:2 Unaffected Mother 1:1