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Table 1

Patient no.	Allele 1	Allele 2
<i>SLC4A1</i>		
1	c.577_578del p.(Ser193 fs)	Normal
2	c.1030C>T p.(Arg344*)	Normal
3	c.1421C>A p.(Ala474Asp)	Normal
4	c.2057+1G>A (splicing)	Normal
5	c.2057+1G>A (splicing)	Normal
6#	c.2057+1G>A (splicing)	Normal
7	c.2348T>A p.(Ile783Asn)	Normal
<i>ANK1</i>		
8#	c.341C>T p.(Pro114Leu)	Normal
9	c.341C>T p.(Pro114Leu)	Normal
10#	c.344T>C p.(Leu115Pro)	Normal
11	c.344T>C p.(Leu115Pro)	Normal
12	c.498C>G p.(Tyr166*)	Normal

Patient no.	Allele 1	Allele 2
13	c.2559-2A>G (?)	Normal
14	c.5201_5202insTCAG p.(Thr1734 fs)	Normal
15	c.5201_5202insTCAG p.(Thr1734 fs)	Normal
	<i>SPTA1</i>	
16	c.678G>A p.(Glu227 fs) + α^{LELY}	α^{LELY}
17	c.2755G>T (p.Glu919*)	α^{LELY}
18	c.4339-99C>T p.(?)	c.4339-99C>T p.(?)
19	c.4339-99C>T p.(?)	Normal
20#	c.4339-99C>T p.(?)	c.7201C>T p.(Arg2401*)
21	c.4738-1G>A p.(?)	Normal
	<i>SPTB</i>	
22#	c.2136_2137delinsTT	Normal
23#	c.3449G>A (p.Trp1150*)	Normal

- Splenectomized patients are noted with # and mutations not previously reported are depicted in boldface. α^{LELY} represents the c.[5572C>G;6531-12C>T] p.[(Leu1858Val);(?)] mutations in *SPTA1*.

Figure 1

Figure 2