

<b>PRLR, wt</b>	430	AGA	TAT	TCT	TAC	CAC	AAC	ATT	GCT	GAC	GTG	<b>TGT</b>	GAG	CTG	GCC	443
		R	Y	S	Y	H	N	I	A	D	V	C	E	L	A	
<b>PRLR, p.C440*</b>	430	AGA	TAT	TCT	TAC	CAC	AAC	ATT	GCT	GAC	GTG	<b>TGA</b>	GAG	CTG	GCC	440
		R	Y	S	Y	H	N	I	A	D	V	*				
<b>PRLR, wt</b>	456	CAA	ACA	GAC	CAA	CAT	<u>GCT</u>	TTA	AAA	GCC	<b>TCA</b>	AAA	ACC	ATT	GAA	456
		Q	T	D	Q	H	<b>A</b>	L	K	A	S	K	T	I	E	
<b>PRLR, p.L462*</b>	456	CAA	ACA	GAC	CAA	CAT	GTT	TAA	AAG	CCT	CAA	AAA	CCA	TTG	AAA	456
		Q	T	D	Q	H	<b>V</b>	*								
<b>PRLR, p.S465*</b>	456	CAA	ACA	GAC	CAA	CAT	GCT	TTA	AAA	GCC	<b>TAA</b>	AAA	ACC	ATT	GAA	465
		Q	T	D	Q	H	A	L	K	A	*					

**Supplementary Figure 2.** Sequence of wt and HDR-edited alleles. Nucleotide and amino acid sequences of relevant regions are shown for the wt allele and alleles with edited p.C440\*, p.L462\* and p.S465\* mutations. Numbers refer to amino acid sequence positions in *PRLR*. Gene sequence changes are indicated in bold, the nucleotide deleted in the p.L462\* edited allele is underlined. Premature stop codons and the amino acid change (A to V in pL462\*) are in bold.