Table W2 CDSN SNPs identified by resequencing

	SNP Position	**Nucleotide Change**
-712	Promoter	c / t
-646	Promoter	a / g
-461	Promoter	c / t
-31	Promoter	a / g
-22	Promoter	c / t
9	5′UTR (exon1)	c / t
75i	Intron 1	a / g
137	Coding(exon 2)	c / t
180	Coding(exon 2)	c / t
206	Coding(exon 2)	c / t
442	Coding(exon 2)	a / g
614	Coding(exon 2)	a / g
619	Coding(exon 2)	c / t
722	Coding(exon 2)	c / t
767	Coding(exon 2)	g / a
971	Coding(exon 2)	c / t
1118	Coding(exon 2)	a / g
1215	Coding(exon 2)	a / g
1236	Coding(exon 2)	g / t
1243	Coding(exon 2)	c / t
1331	Coding(exon 2)	c / g
1593	Coding(exon 2)	g / a
1606	3′UTR	del / aag
1675	3′UTR	g / t
1739	3′UTR	c / t
1747	3′UTR	c / t

The variants selected for genotyping are typed in bold