

Table S1A: Sanger sequencing efforts.

chrom	start	stop	orientation	inds
1	1121520	1122379	revcomp	Cr1Gr1, Cr75, Cr81, Crtaal
1	2189244	2189862	revcomp	Cr1Gr1, Cr75, Cr81, Crtaal
1	3187217	3187782	revcomp	Cr75, Cr81, Crtaal
1	9252546	9253264	forward	Cr1337, Cr1Gr1, Cr81, Crtaal
1	9791410	9792193	revcomp	Cr1Gr1, Cr75, Cr81, Crtaal
1	12972008	12972691	forward	Cr1Gr1, Cr81, Crtaal
1	16978816	16979446	forward	Cr1Gr1, Crtaal
2	7158433	7159032	revcomp	Cr1Gr1, Cr75, Cr81, Crtaal
2	9111928	9112538	revcomp	Cr75, Crtaal
2	11501659	11502346	revcomp	Cr1Gr1, Cr75, Cr81, Crtaal
2	13499343	13499957	forward	Cr81
3	4380447	4381090	revcomp	Cr81
3	12783435	12784119	revcomp	Cr1Gr1, Cr75, Cr81, Crtaal
7	2781533	2782194	forward	Cr1Gr1, Cr75, Cr81, Crtaal
7	3433282	3434023	revcomp	Cr1Gr1, Cr81, Crtaal
7	3458572	3459198	revcomp	Cr1Gr1, Cr75, Cr81, Crtaal
7	4820394	4821059	forward	Cr1337, Cr1Gr1, Cr81, Crtaal
7	6079504	6080090	forward	Cr1337, Cr1Gr1, Cr75, Cr81, Crtaal
7	9123830	9124515	forward	Cr1Gr1, Cr75, Cr81
7	9214127	9214749	forward	Cr1Gr1, Cr75, Cr81, Crtaal
8	2578408	2579001	revcomp	Cr1Gr1, Cr75, Cr81, Crtaal
8	6671378	6671912	revcomp	Cr1Gr1, Cr75, Cr81, Crtaal
8	7784867	7785350	forward	Cr1Gr1, Crtaal, Cr1337, Cr75
8	9670696	9671200	forward	Cr1Gr1, Cr75, Cr81, Crtaal

Table S1B: Comparison of genotype calls across technology (Sanger/RNA-Seq).

Sample	0/0	2/2	0/NA	NA/0	2/NA	0/2	1/1	NA/NA
Cr1Gr1	12808	25	6	92	0	0	0	50
Cr75	10611	19	5	53	0	0	0	28
Cr81	13494	6	37	104	0	0	1	50
Crtaal	12904	8	369	139	2	2	0	50
Cr1337	2429	0	12	0	0	0	0	13