

Panel-based NGS Reveals Novel Pathogenic Mutations in Autosomal Recessive Retinitis Pigmentosa

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Supplementary Table 1. Regions not covered in our RD panel.

Chr	Start position	End position	Gene	Exon
2	96967395	96967407	<i>SNRNP200</i>	4
2	182468774	182468826	<i>CERKL</i>	2
6	64497950	64498012	<i>EYS</i>	39
12	88471664	88471677	<i>CEP290</i>	40
12	88510789	88510829	<i>CEP290</i>	18
17	58232683	58232718	<i>CA4</i>	2
X	46713082	46713137	<i>RP2</i>	2