Next-generation sequencing reveals the mutational lan Usher syndrome: copy number variations, phenocopies translational read-through, and<i>PEX26</i>

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Citation Report

#	ARTICLE Next-generation sequencing reveals the mutational landscape of clinically diagnosed Usher	IF	Citations
1	syndrome: copy number variations, phenocopies, a predominant target for translational read-through, and <i>PEX26 </i> in mutated in Heimler syndrome. Molecular Genetics & Denomic Medicine, 2017, 5, 531-552.	1.2	55
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