

HGNC approved gene symbol	Transcript	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x	% covered >50x
<i>CACNA1A</i> **	NM_001127221	601011	246	100	100	100	100
<i>ATP1A2</i> **	NM_000702	182340	222	100	100	100	100
<i>SCN1A</i> **	NM_001165963	182389	334	100	100	100	100

* Familial hemiplegic migraine

** Core genes