

HGNC approved gene symbol	Transcript	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x	% covered >50x
<i>AGPAT2</i> **	NM_006412**	603100	326	100	100	100	100
<i>AKT2</i>	NM_001626	164731	288	100	100	100	100
<i>BSCL2</i> **	NM_032667**	606158	347	100	100	100	100
<i>CAV1</i>	NM_001753	601047	538	100	100	100	100
<i>LMNA</i> **	NM_170707**	150330	295	100	100	100	100
<i>PLIN1</i>	NM_002666	170290	203	100	100	100	100
<i>PPARG</i> **	NM_015869**	601487	374	100	100	100	100
<i>CAVIN1 (alias PTRF)</i> **	NM_012232**	603198	322	100	100	100	100

\* Lipodystrophy

\*\* Core genes