

	geneID	sampleID	pValue	padjust	zScore	l2fc	rawcounts	normcounts	meanCorrected	aberrant	AberrantBySample	padj_rank	FC	gene_type	OMIM	PINH	PMIM
1	ESYT1	patient 3	4.60E-12	6.34E-07	-7.42	-1.01	5863	6530.32	13130.49	TRUE	3	1	0.5	protein_coding	616670		
2	ZNF160	patient 3	2.24E-08	1.55E-03	-6.02	-0.96	745	499.99	974.34	TRUE	3	2	0.51	protein_coding	600398		
3	DPM1	patient 3	7.08E-08	3.26E-03	-5.59	-0.56	1451	1338.1	1978.9	TRUE	3	3	0.68	protein_coding	603503	Autosomal recessive	608799
2439	CLDN11	patient 3	0.1475797	1.00E+00	-1.49	-0.83	12706	10093.92	17991.61	FALSE	3	6838	0.562529242				

Supplementary table 1

Significantly differentially expressed genes (adjusted $p < 0.05$). *CLDN11* is not significantly differentially expressed compared to 156 control fibroblasts.