

Supplemental Table S1Summary of the quantification analyses of *Skn-1a*-, *Trpm5*-, and *Mash1*-expressing cells in the MOE.

Mouse ID		S384	S656	S660	Average	SD
Number of cells	Skn1a+ cells	787	199	214	400	335
	Trpm5+ cells	1395	731	378	835	516
	Skn1a+Trpm5+ cells	603	144	180	309	255
Population (%)	Skn1a+Trpm5+ / Skn1a+	76.6%	72.4%	84.1%	77.7%	5.9%
	Skn1a+ Trpm5+ / Trpm5+	43.2%	19.7%	47.6%	36.8%	15.0%

Mouse ID		S384	S656	S660	Average	SD
Number of cells	Skn1a+ cells	566	290	272	376	164.8
	Mash1+ cells	2963	2493	1269	2242	874.5
	Skn1a+ Mash1+ cells	29	30	26	28	2.1
Population (%)	Skn1a+ Mash1+ / Skn1a+	5.1%	10.3%	9.6%	8.3%	2.8%
	Skn1a+ Mash1+ / Mash1+	1.0%	1.2%	2.0%	1.4%	0.56%