

**Supplemental Table 3. qPCR data on *LCE* gene expression in epidermis of normal skin before and after tape-stripping**

HUGO symbol	mRNA <i>in vivo</i>		Ratio Relative Q after TS/before TS	P value <sup>4</sup>
	dCt <sup>1</sup> before TS <sup>2</sup>	dCt after TS <sup>3</sup>		
<b><i>LCE1 group</i></b>				
LCE1A	4.7	8.1	0.14	<b>0.002</b>
LCE1B	4.8	8.5	0.11	<b>0.002</b>
LCE1C	4.4	7.3	0.17	<b>0.002</b>
LCE1D&E	6.8	9.7	0.06	0.093
LCE1F	8.2	11.4	0.12	<b>0.001</b>
<b><i>LCE2 group</i></b>				
LCE2A	4.7	7.4	0.19	<b>0.006</b>
LCE2B	5.0	8.6	0.09	<b>0.000</b>
LCE2C	5.0	7.8	0.18	<b>0.002</b>
LCE2D	5.2	8.4	0.14	<b>0.006</b>
<b><i>LCE3 group</i></b>				
LCE3A	11.5	6.5	12	<b>0.015</b>
LCE3C	12.4	8.2	20	0.078
LCE3D	5.5	2.5	5.8	<b>0.044</b>
LCE3E	7.2	4.6	5.3	<b>0.046</b>
<b><i>LCE5 group</i></b>				
LCE5A	5.4	11.3	0.03	<b>0.001</b>
<b><i>LCE6 group</i></b>				
LCE6A	7.6	10.9	0.13	<b>0.001</b>

<sup>1</sup>dCt: Delta PCR cycle time. <sup>2</sup>mRNA expression data from epidermal cells of normal skin before tape-stripping (TS) (N=5), <sup>3</sup> after tape-stripping (N=5), <sup>4</sup>P-values of paired t-test. Bold: p-value < 0.05.