

**Table SIV. Microarray expression of genes relevant to this study**

Agilent Probe ID	Ref Sequence	Gene symbol	Site within lesion	Fold change	p-value
A_33_P3284345	NM_013956.3	NRG1	Centre dermis	10.71	0.0002
A_33_P3284345	NM_013956.3	NRG1	Margin dermis	22.24	1.59E-05
A_33_P3284345	NM_013956.3	NRG1	Extralesional dermis	-2	0.42
A_33_P3292596	NM_004448.3	ErbB2	Margin dermis	2.56	0.0031
A_33_P3292596	NM_004448.3	ErbB2	Margin dermis	1.47	0.1633
A_33_P3292596	NM_004448.3	ErbB2	Extralesional dermis	1.03	0.927
A_23_P349416	NM_001982.3	ErbB3	Centre dermis	-10.14	3.53E-06
A_23_P349416	NM_001982.3	ErbB3	Margin dermis	-15.6	6.82E-07
A_23_P349416	NM_001982.3	ErbB3	Extralesional dermis	-1.25	0.6075
A_32_P183765	NM_005235.2	ErbB4	Centre dermis	-6.83	0.0038
A_32_P183765	NM_005235.2	ErbB4	Margin dermis	-5.42	0.0145
A_32_P183765	NM_005235.2	ErbB4	Extralesional dermis	-5.96	0.013