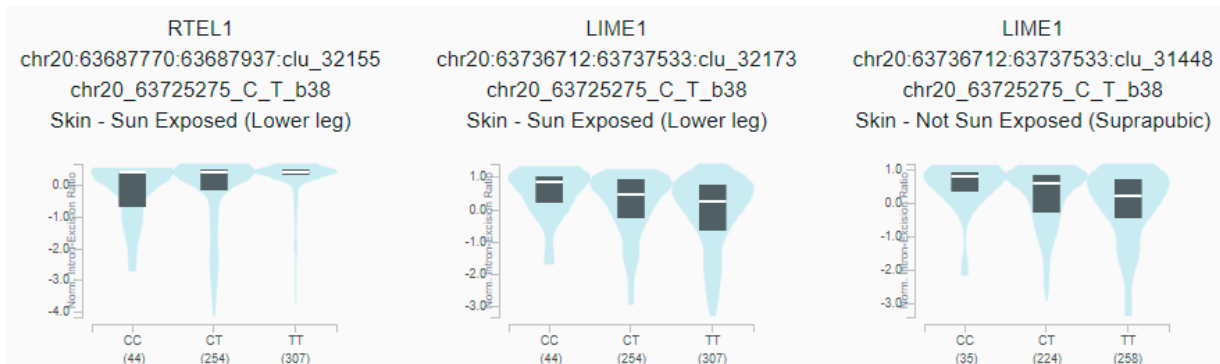
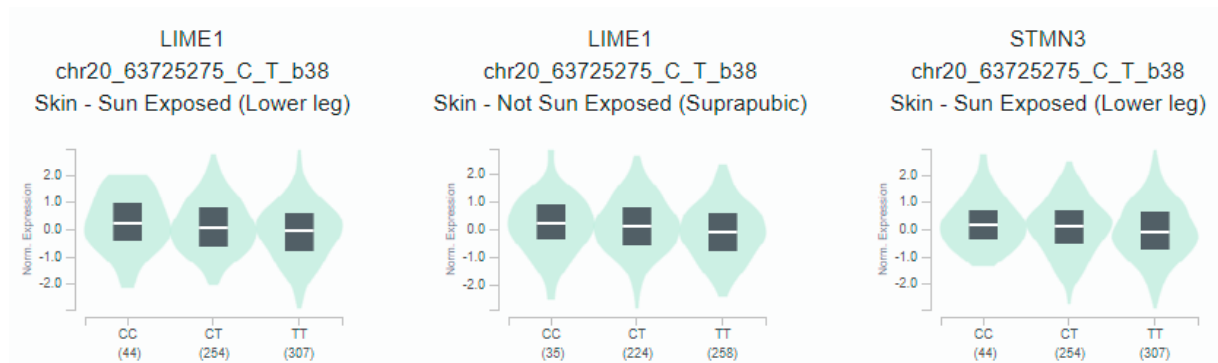


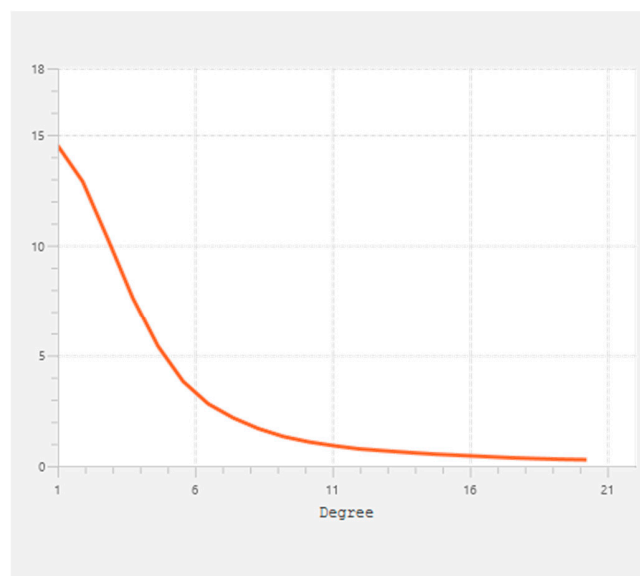
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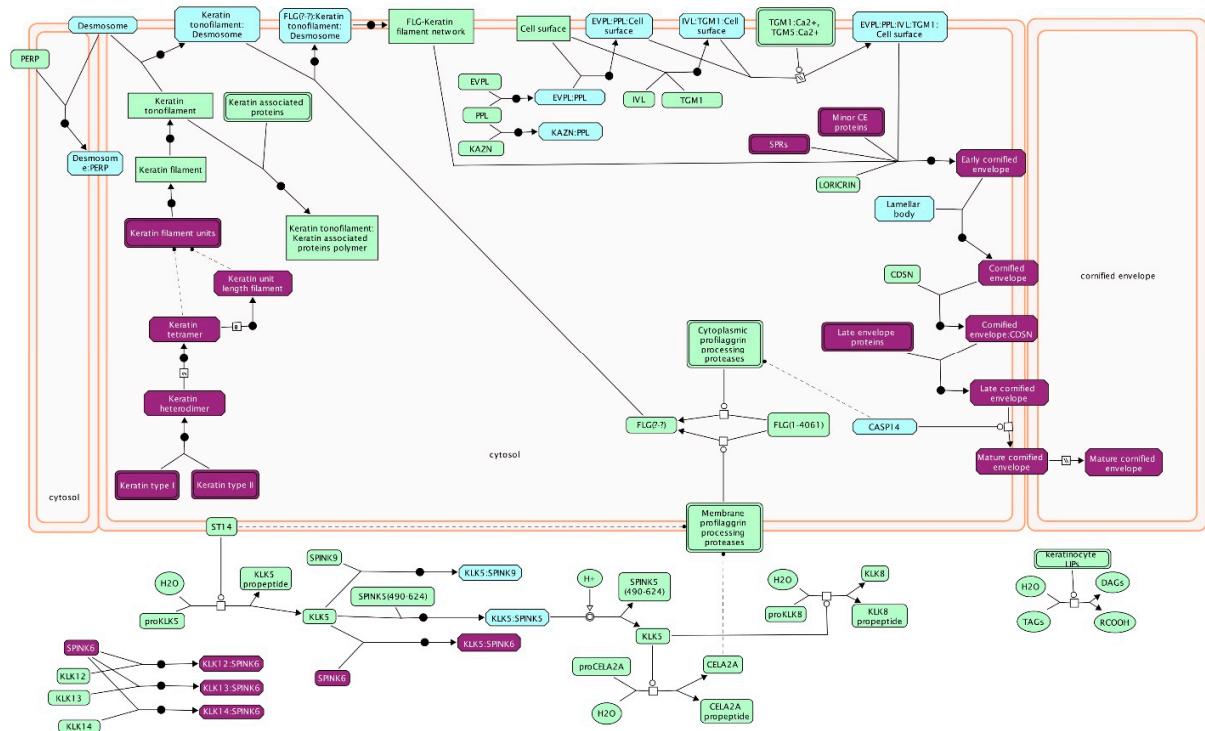
B



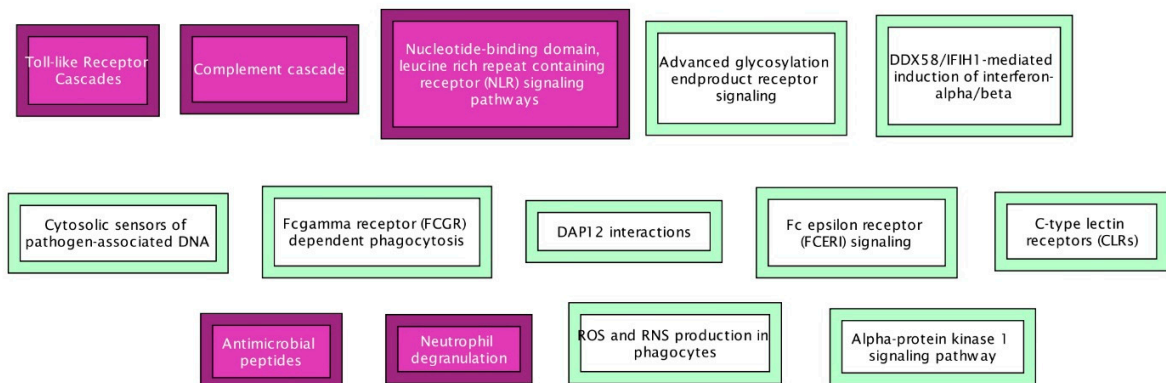
Supplementary Figure S1. Splicing-Quantitative Trait Locus (QTL) A) and expression-QTL (B) genes associated with hand eczema identified in skin tissue based on the GTEx database.



Supplementary Figure S2. The distribution of node degrees for the constructed (vesicular) hand eczema network. Y-axis shows the number of nodes with a certain calculated degree of centrality.



Supplementary Figure S3. Reactome diagram of the keratinization pathway with nodes involving our (vesicular) hand eczema genes highlighted in purple.



Supplementary Figure S4. Immune pathways involving our (vesicular) hand eczema genes highlighted in purple.

