



Fig. S4. Number of altered transcripts/differentially regulated genes in the epidermis of groups of patients with X-linked recessive ichthyosis (XRI) and ichthyosis vulgaris with mono- (*pink*) or biallelic (*red*) *FLG* mutations. (From the thesis of T Hoppe, Acta Universitatis Uppsaliensis, ISBN 978-91-554-8583-2, 2013).