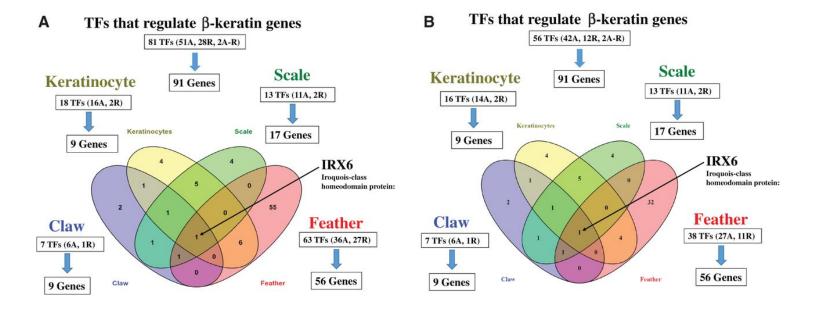
Fig. 2



Summary of annotated TF-target gene relationships. (*A*) Overall predictions and (*B*) predictions supported by differential expression of pTF-target gene. The figure summarizes the number of annotated TFs and their corresponding numbers of target  $\beta$ -keratin genes. The annotated TFs are further classified according to the sub-class (scale, claw, keratinocytes and feather) of the  $\beta$ -keratin genes they regulate. The entries "A" and "R" denotes "activator" and "repressor", respectively. The Venn-diagram shows the numbers of common and unique regulators among the four sub-classes of  $\beta$ -keratins. Iroquois-class homeodomain (IRX6) was found the sole TF protein which commonly regulates all the four sub-classes of  $\beta$ -keratin genes.