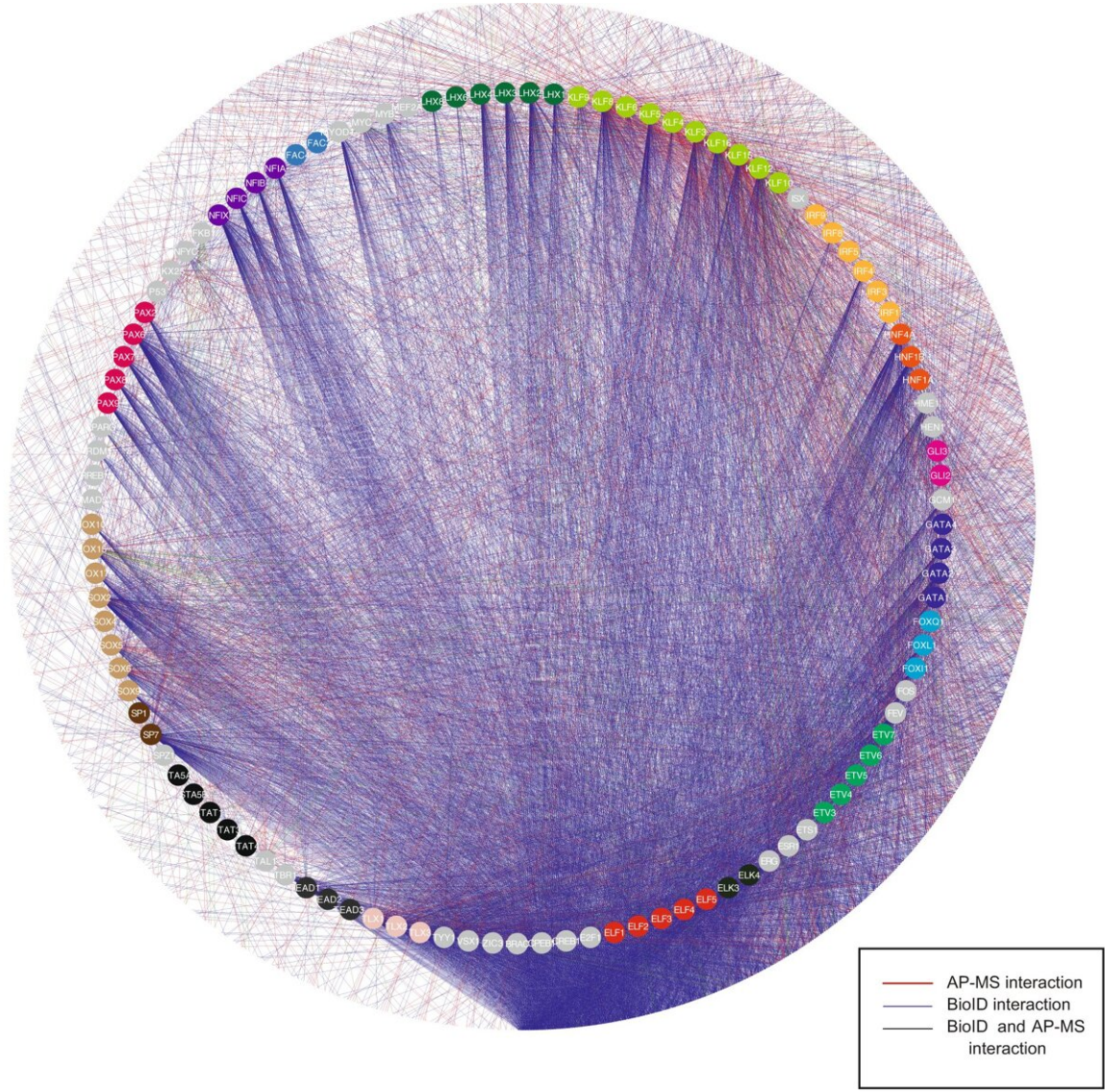


# Researchers resolved human transcription factor (TF) regulation

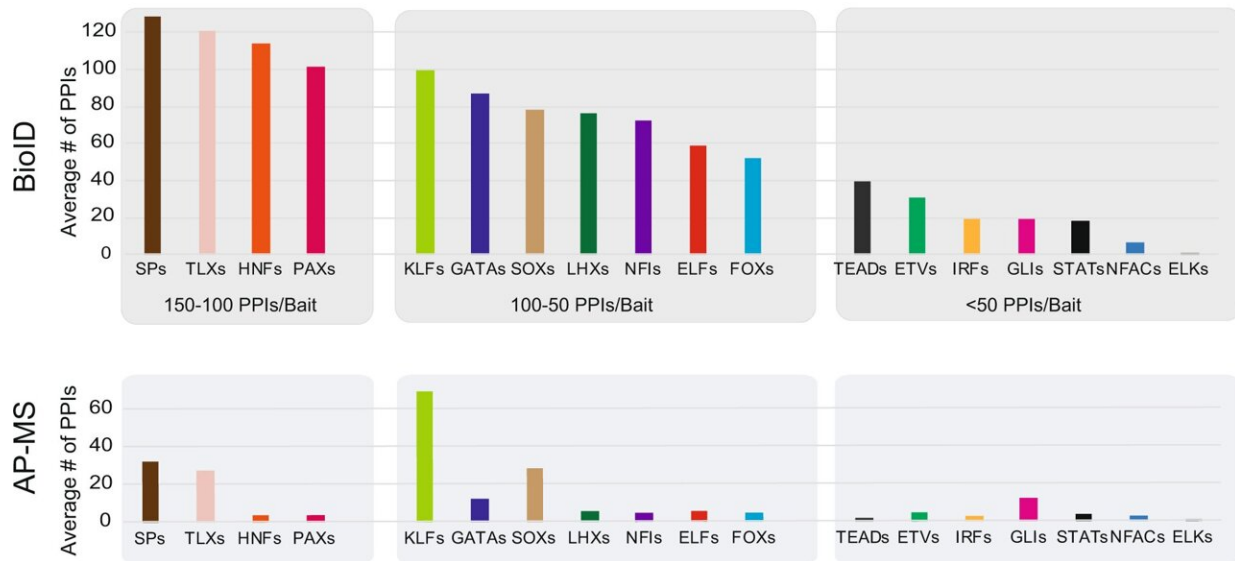
February 9 2022

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**a**



**b**



Comprehensive protein interactomes of the studied TF and TF families. a Studied TFs are organized and color-coded (node color) based on their TF families in the inner circle, and interacting proteins are shown in the outer circle with white. Blue edges indicate interactions detected with the BioID analysis, red with the AP-MS analysis and green from both. b The average number of PPIs of different TF families detected by BioID and AP-MS. Note that the color coding of the TF families is the same as in a) and the TF families are organized into three bins based on the BioID data, first having families with 150–100 high-confidence interactions/bait TF, second 100–50, and third

Citation: Researchers resolved human transcription factor (TF) regulation (2022, February 9) retrieved 3 July 2023 from <https://medicalxpress.com/news/2022-02-human-transcription-factor-tf.html>

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