the number of interacting partners associated with the protein detection method and the scale of the study all UniprotAC were coerced to generic UniprotAC, grouped by interaction detection method (colour), large and small scale experiments | small\_scale large\_scale | large\_scale | small\_scale 1000 Escherichia coli, strain K12 10 -1000 Homo sapiens two\_hybrid\_vs\_AP\_MS 10 not two-hybrid, and not AP-MS not two-hybrid, AP-MS two-hybrid and AP-MS 1000 two-hybrid, not AP-MS Mus musculus 10-1000 -S. cerevisiae, strain S288c 10 northo-hybrid, and nor AP-MS not two-hybrid, and not AP-MS nortwo-hybrid and nor AP-MS not two hybrid, Ap. Ms two-hybrid and Ap-Ms two-hybrid, not AP-MS not two hybrid, Ap. Ms two-hybrid and Ap-Ms two-hybrid, not Ap-Ms not mo-hybrid, Ap. Ms two-hybrid and Ap-Ms two-hybrid, not Ap-Ms

interaction detection method

the number of interacting partners