the number of interactions (interaction IDs) 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 100 two-hybrid and AP-MS two-hybrid, not AP-MS Mus musculus 10 -1000 -S. cerevisiae, strain S288c 10 not two-hybrid, and not AP-MS northo-hybrid, and not AP MS nortwo-hybrid and nor AP MS not two hybrid, Ap. Ms two-hybrid and Ap-Ms not two hybrid, Ap. Ms two-hybrid, not AP-MS not two hybrid, Ap. Ms two-hybrid, not Ap-MS two-hybrid and Ap-Ms two-hybrid and Ap-Ms

interaction detection method

the number of interaction IDs, interaction detection experiments