Proteome coverag	e by interaction databases, overlap between IMEx and BioGRID
	IMEx whole_proteome_Uniprot IMEx whole_proteome_Uniprot
Homo sapiens	121 3959 13839 1525138214 2659 1421 13839 1525 3413 BioGRID_from_Mentha BioGRID_from_Mentha
	IMEx whole_proteome_Uniprot IMEx whole_proteome_Uniprot
Mus musculus	21
	IMEx whole_proteome_Uniprot IMEx whole_proteome_Uniprot
strain ATCC 204508	52
	IMEx whole_proteome_Uniprot IMEx whole_proteome_Uniprot
Drosophila melanogaster	13 8173 1982 102 32262 7431 755 1982 102 621 BioGRID_from_Mentha
	IMEx whole_proteome_Uniprot IMEx whole_proteome_Uniprot
Caenorhabditis elegans	3 4214 829 55 22412 3393 824 829 55 2173 BioGRID_from_Mentha BioGRID_from_Mentha
	IMEx whole_proteome_Uniprot IMEx whole_proteome_Uniprot
Arabidopsis thaliana	24 1580 3097 3662 81138 1264 340 3097 3662 8170 BioGRID_from_Mentha
	UniprotKB, isoforms excluded SwissProt, isoforms excluded