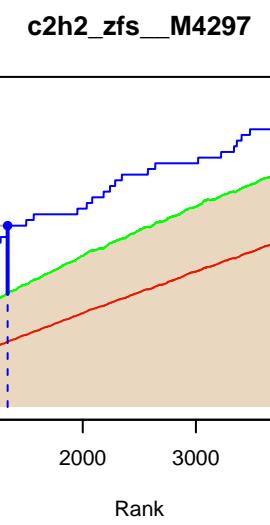
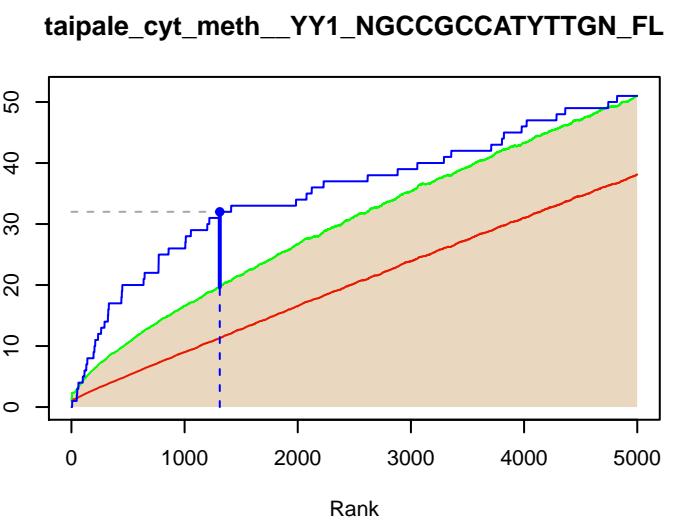
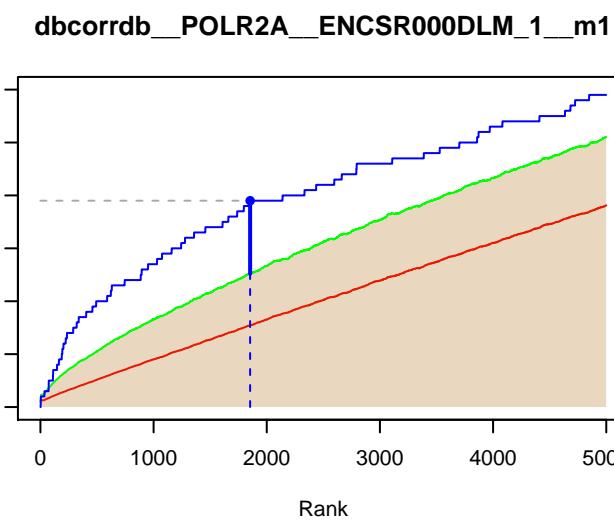
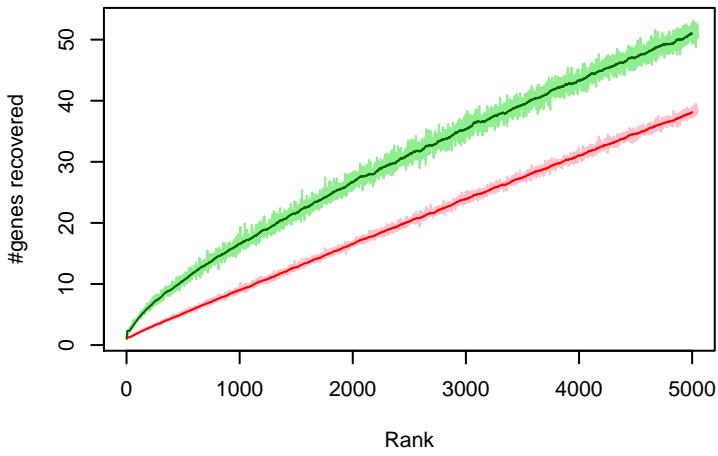
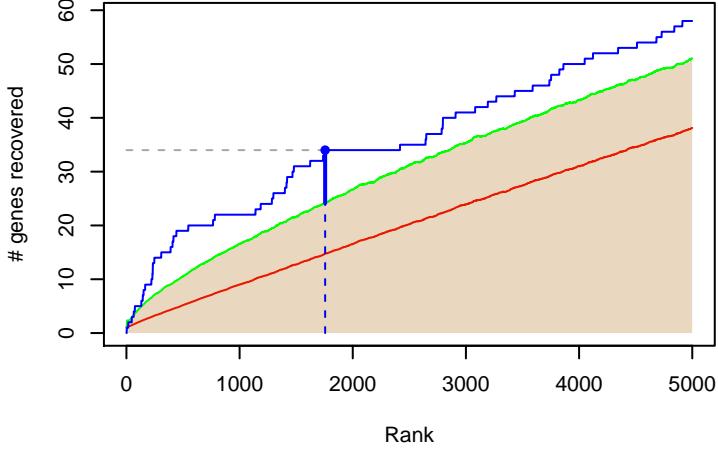
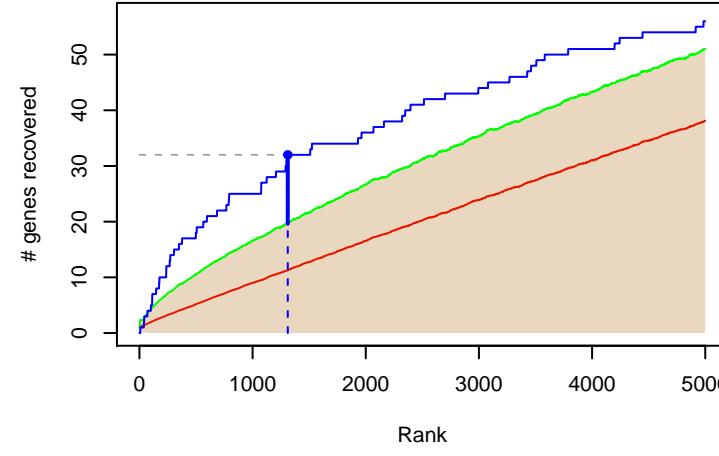
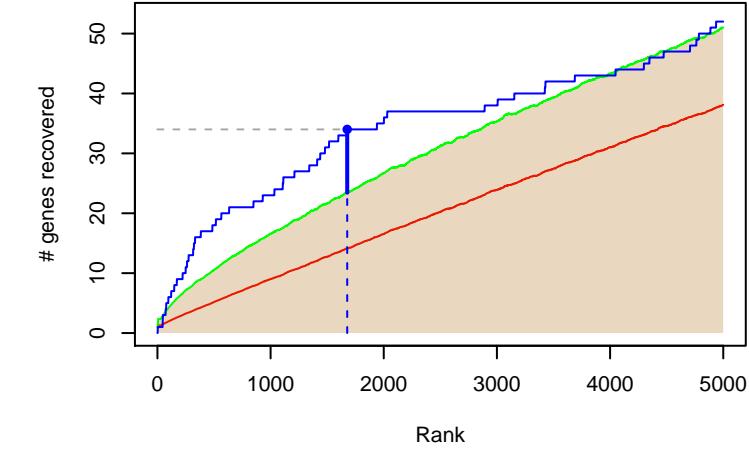
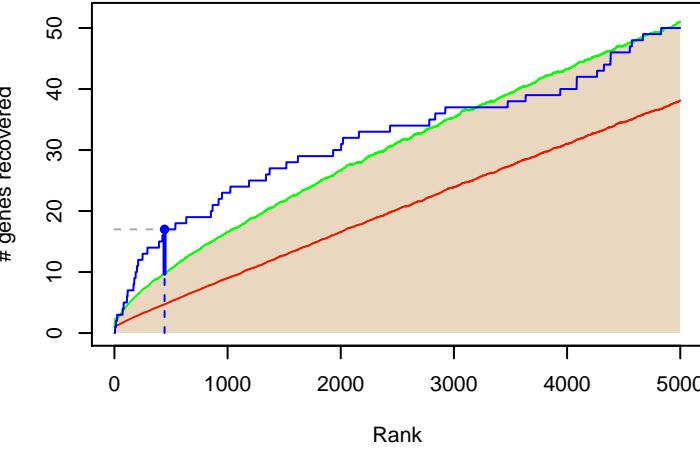
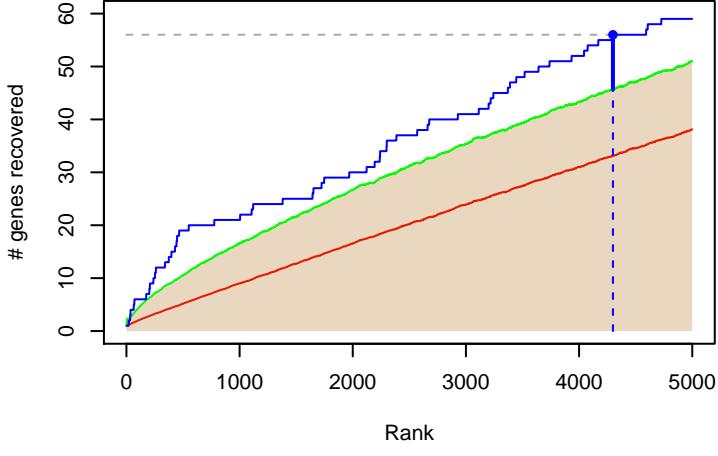
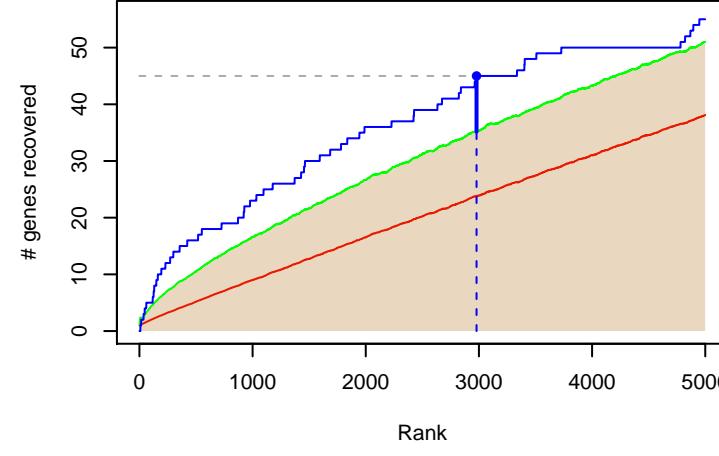
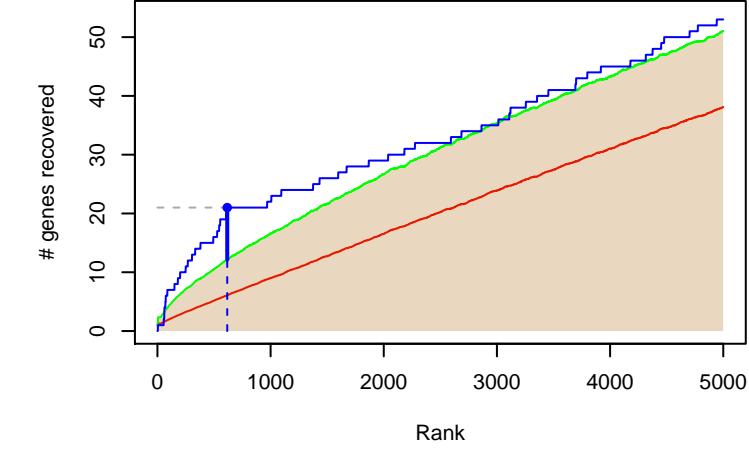
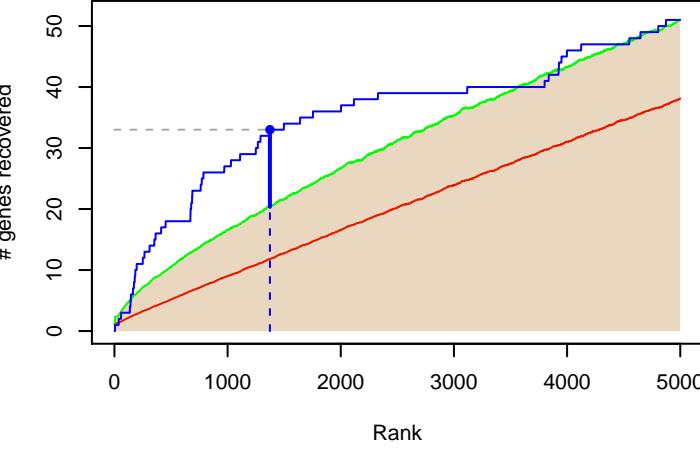
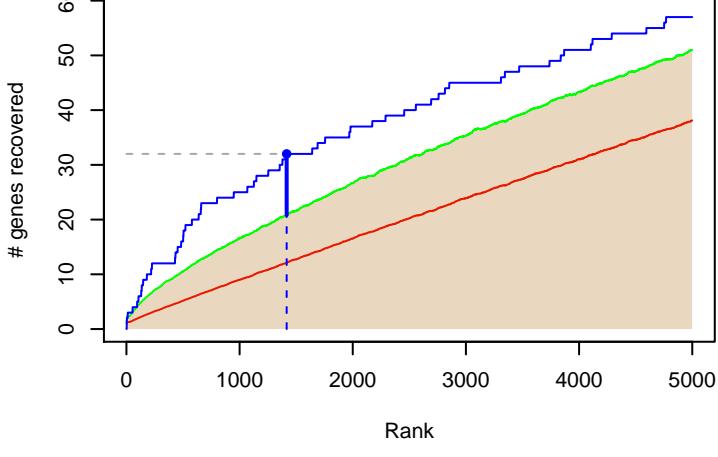
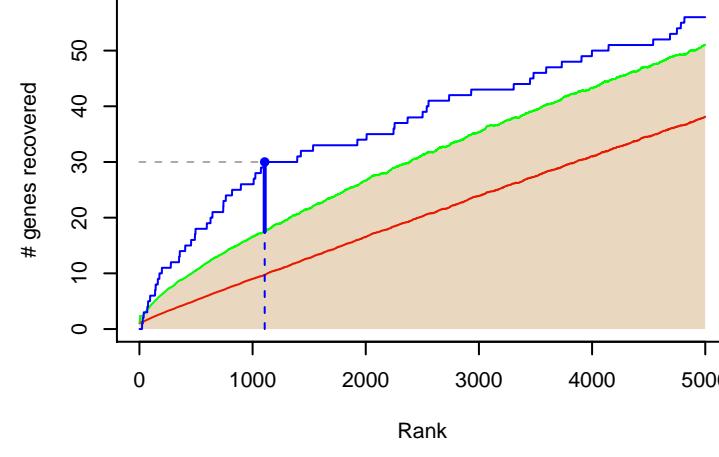
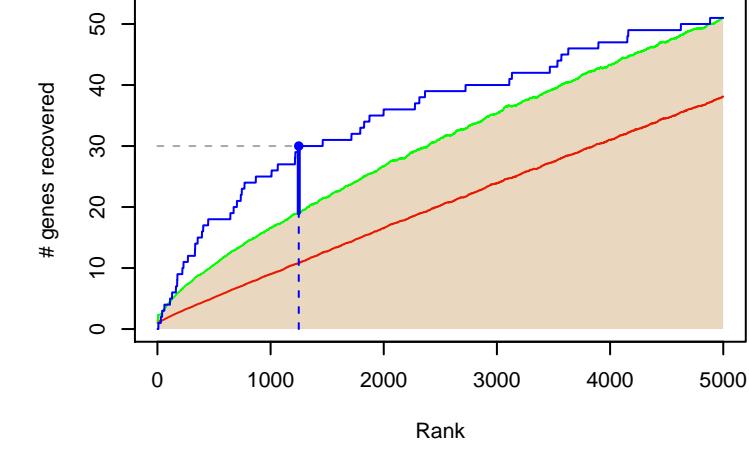
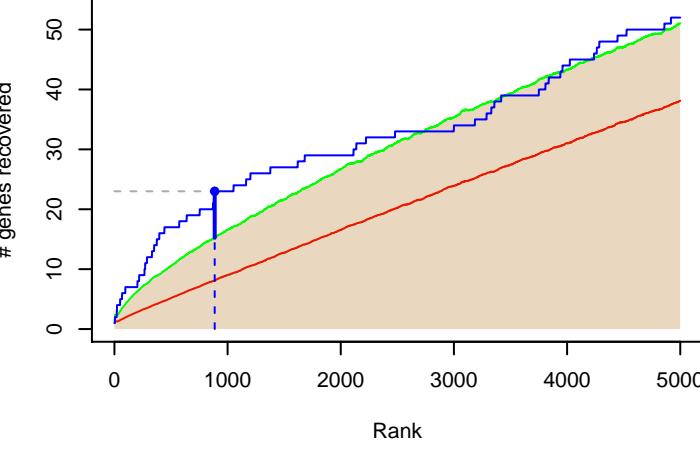
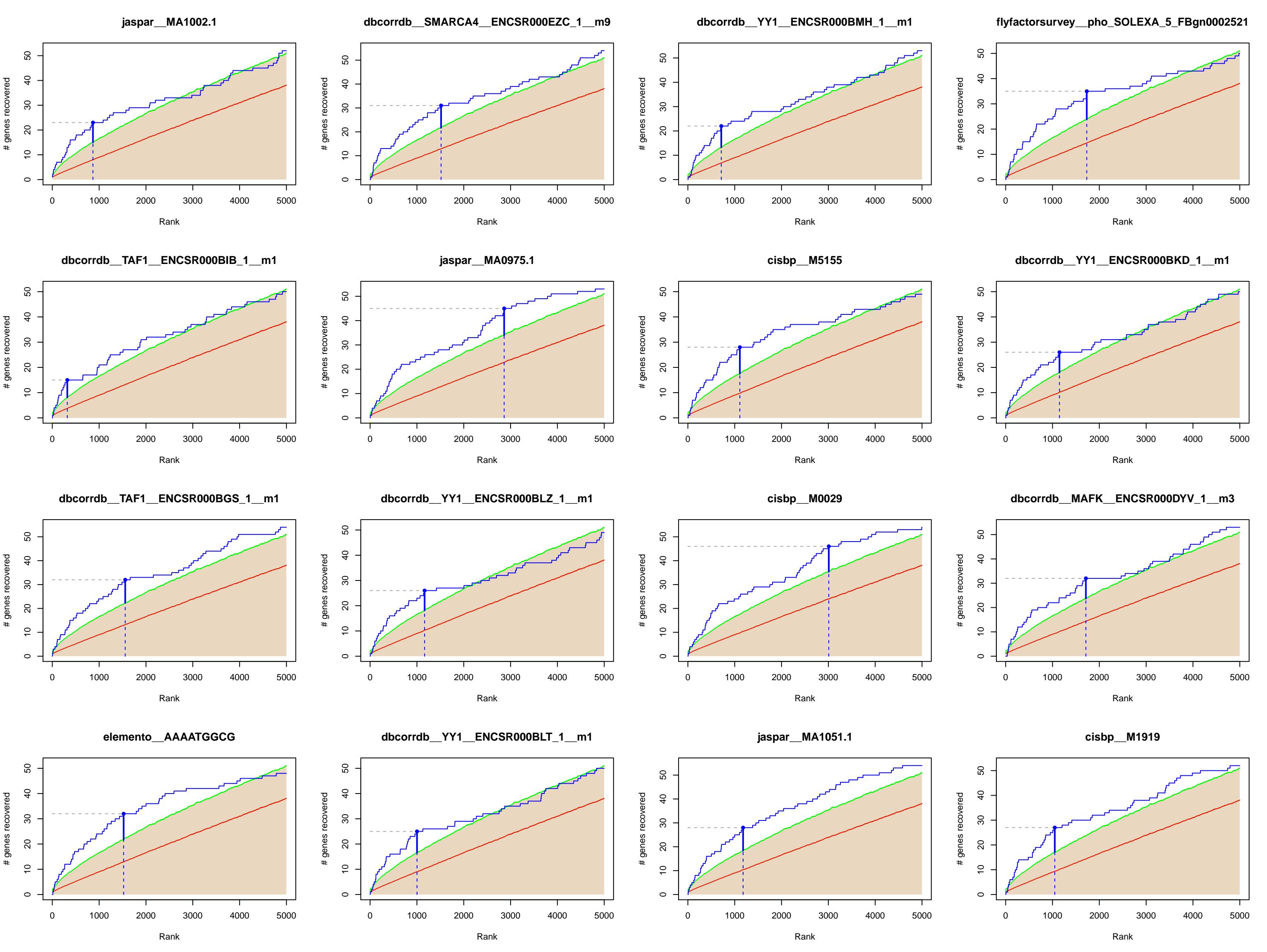
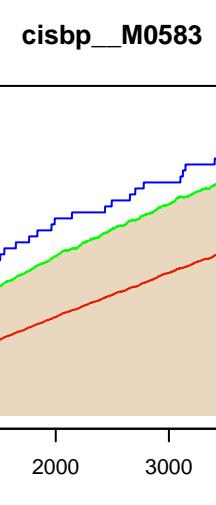
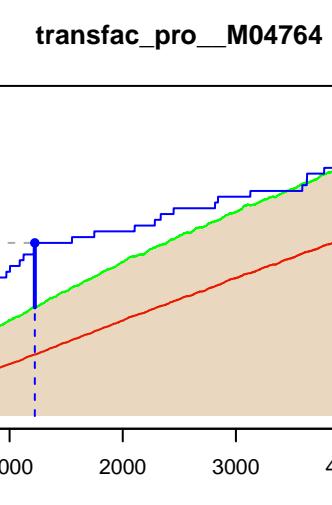
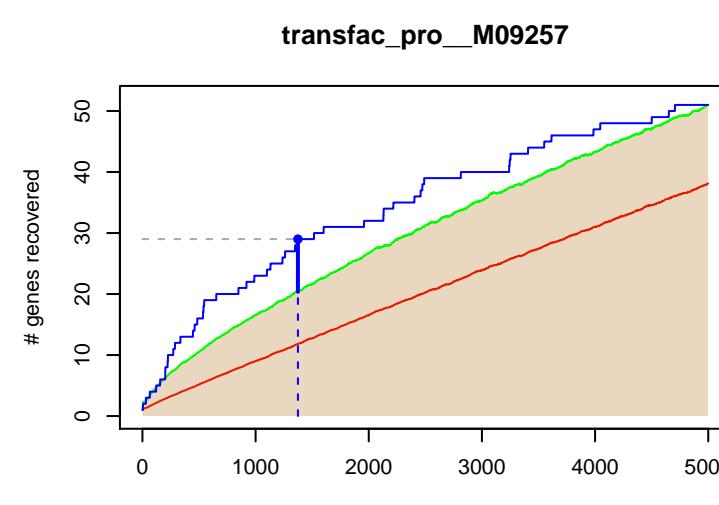
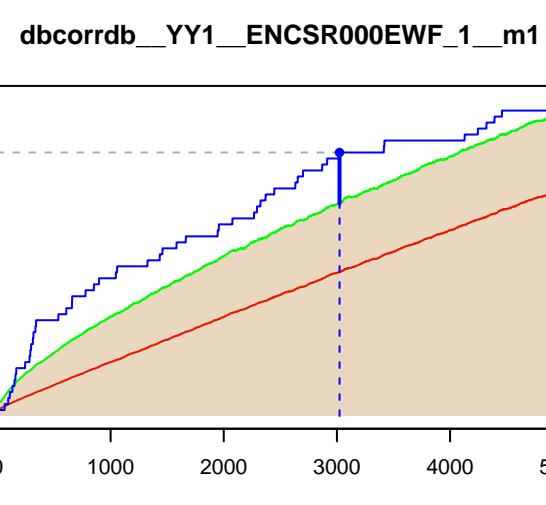
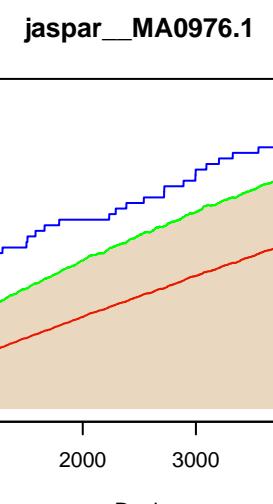
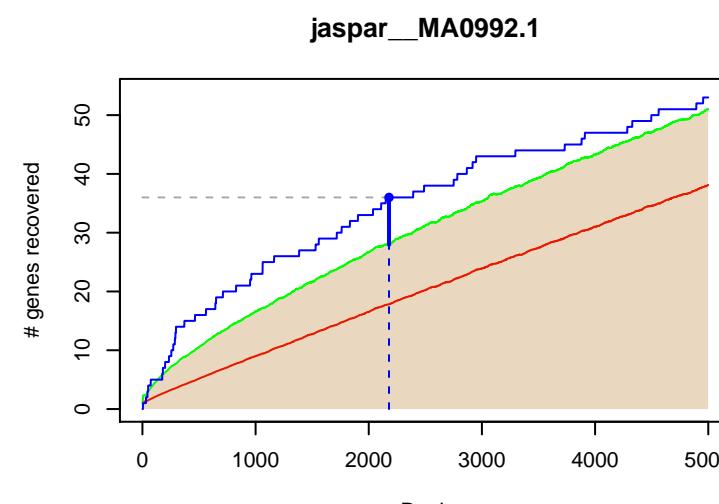
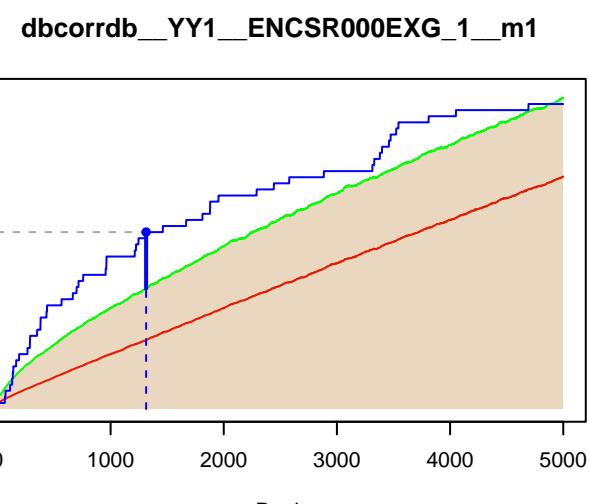
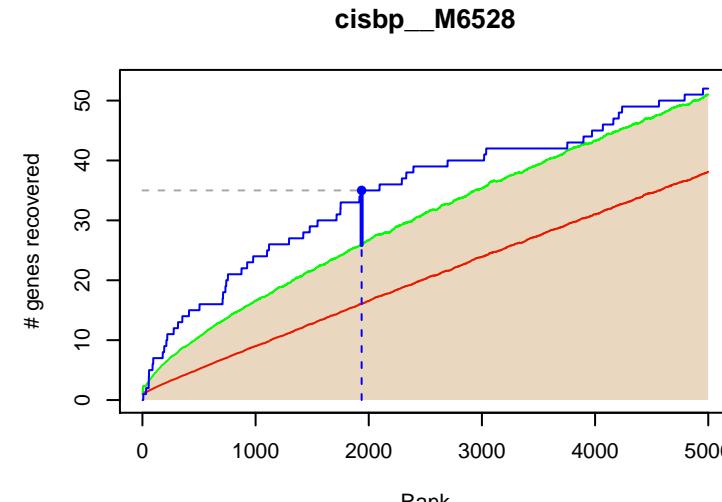
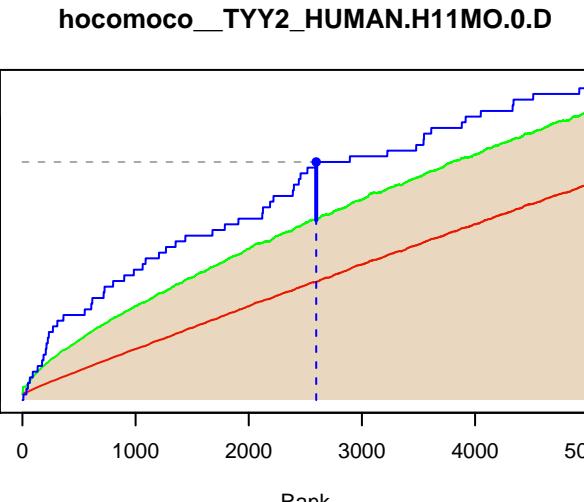
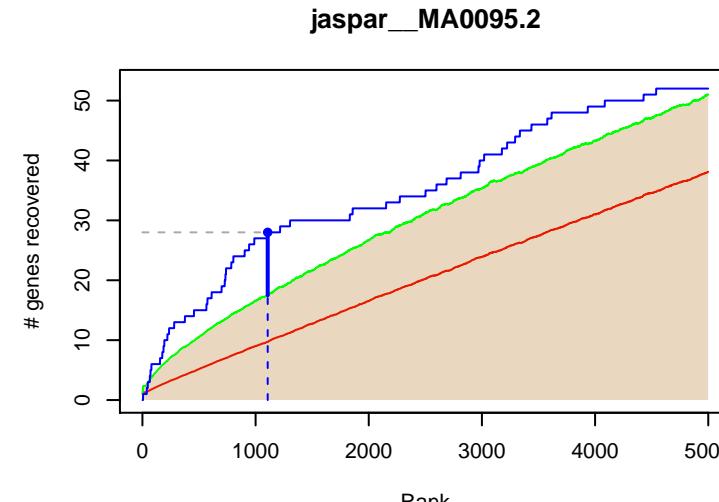
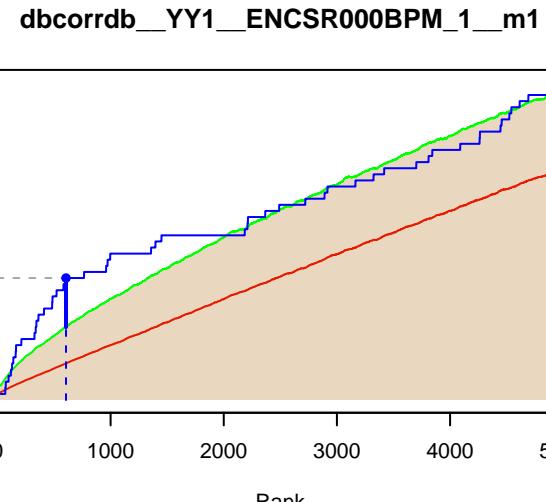
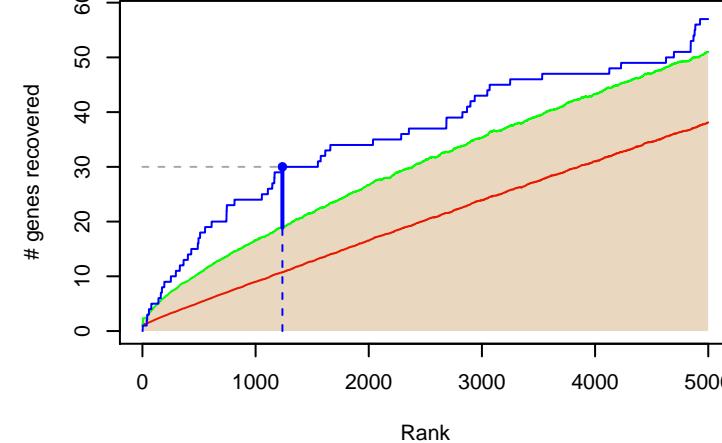
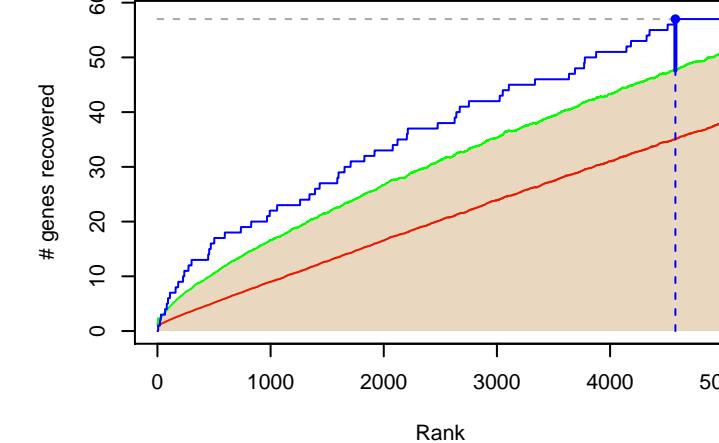
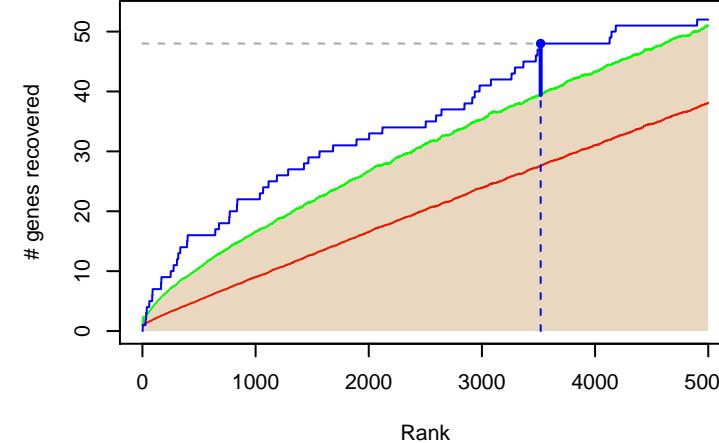
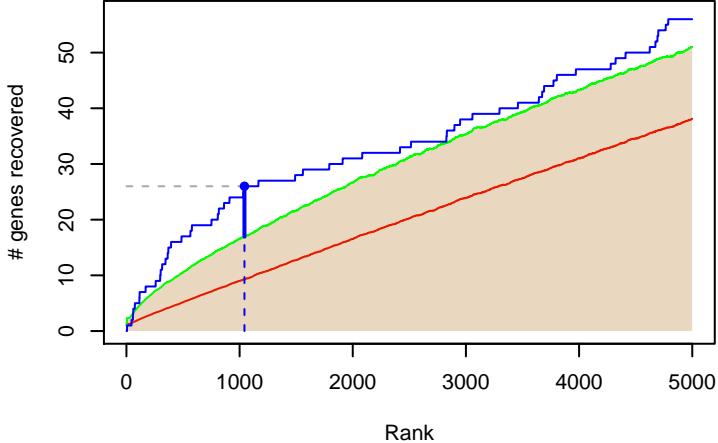
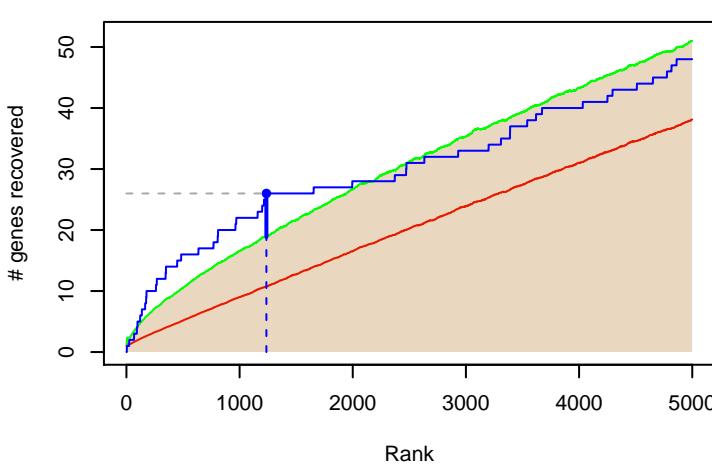


Global mean and SD estimation**dbccorrdb_POLR2A_ENCSR000BGD_1_m1****flyfactorsurvey_phol_SOLEXA_5_FBgn0035997****taipale_cyt_meth_YY1_NGCSGCCATYTTGN_FL_meth****transfac_pro_M08982****dbccorrdb_POLR2A_ENCSR000BHZ_1_m1****dbccorrdb_YY1_ENCSR000EUM_1_m1****dbccorrdb_YY1_ENCSR000BKJ_1_m1****taipale_cyt_meth_ZNF597_NGCCGCCATYTTGN_FL****dbccorrdb_POLR2A_ENCSR000EFK_1_m1****dbccorrdb_ZNF384_ENCSR000DYP_1_m1****cisbp_M4639****cisbp_M0018**

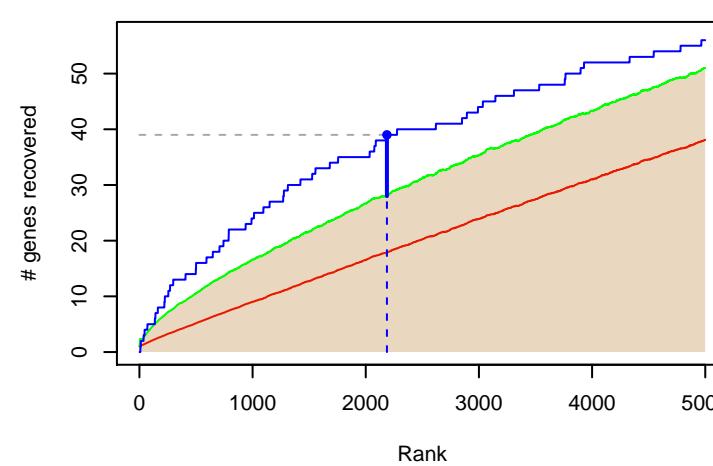




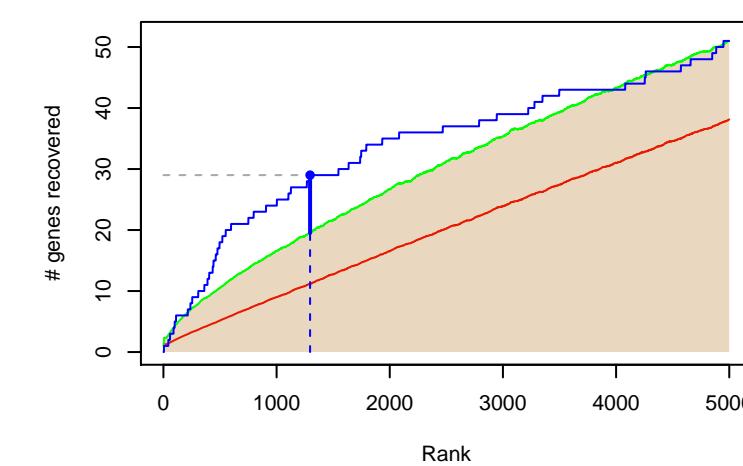
dbcorrdb__YY1__ENCSR000BNX_1_m1



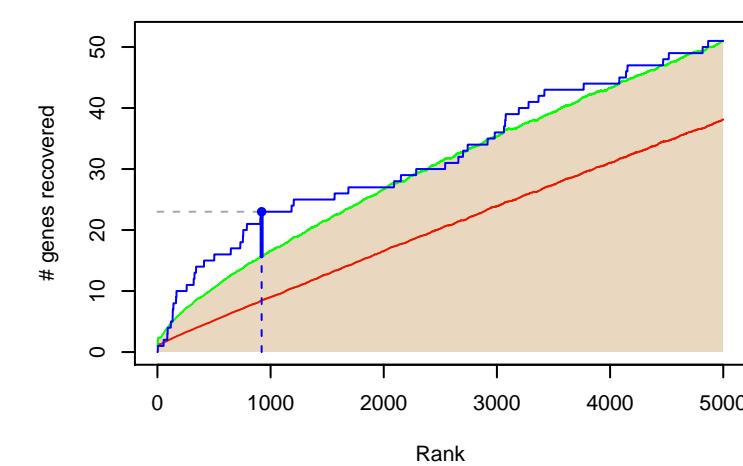
dbcorrdb__ZNF384__ENCSR000EFP_1_m1



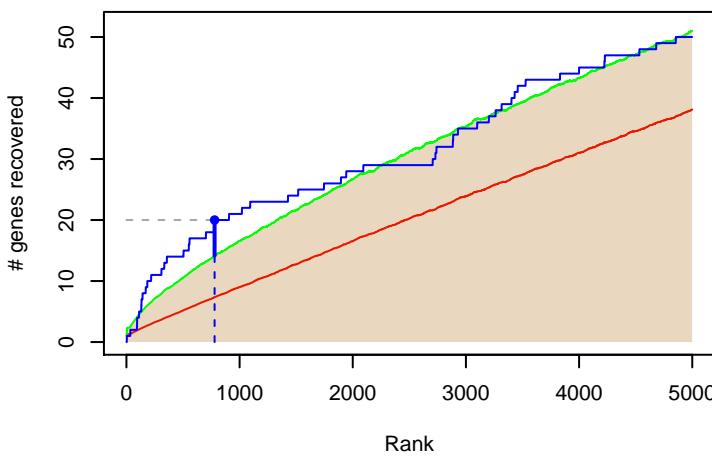
hocomoco__TYY1_HUMAN.H11MO.0.A



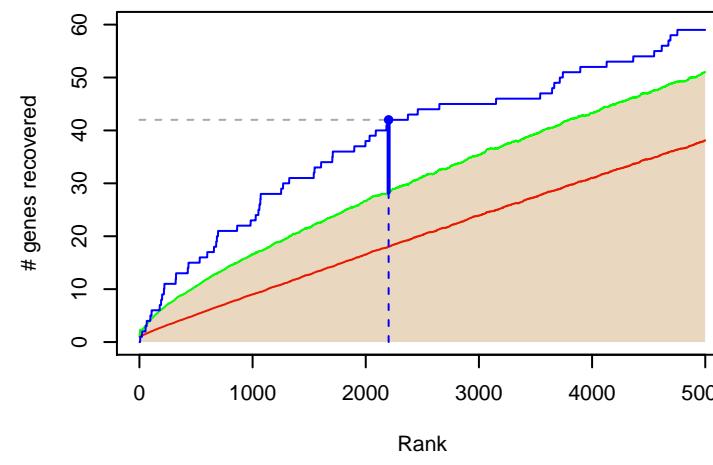
dbcorrdb__YY1__ENCSR000BKU_1_m1



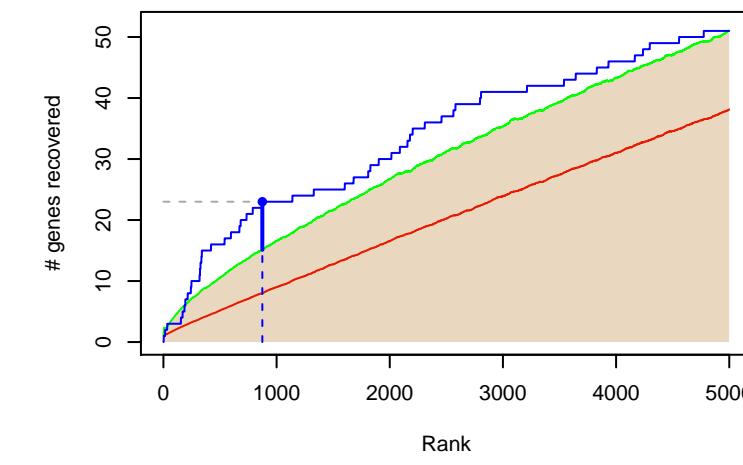
dbcorrdb__YY1__ENCSR000BNT_1_m1



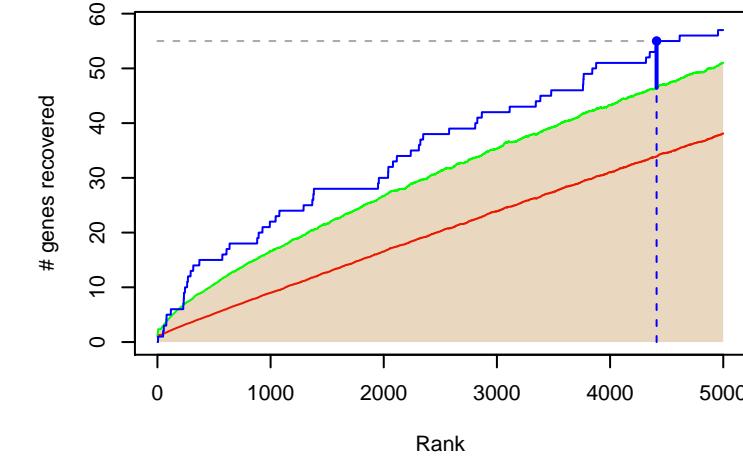
dbcorrdb__POLR2A__ENCSR000BKI_1_m1



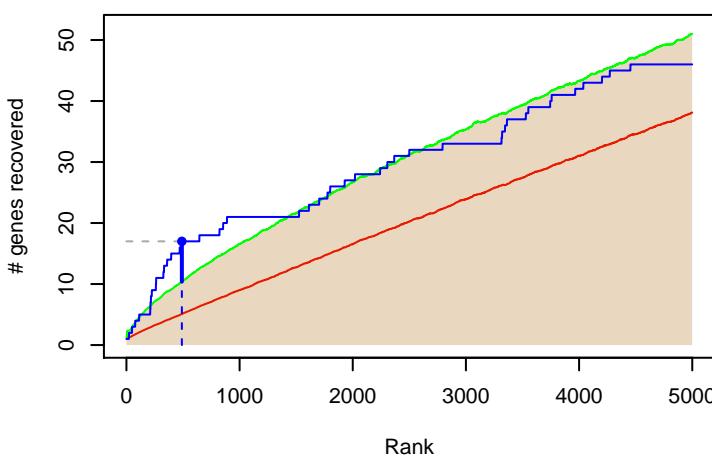
transfac_pro__M04696



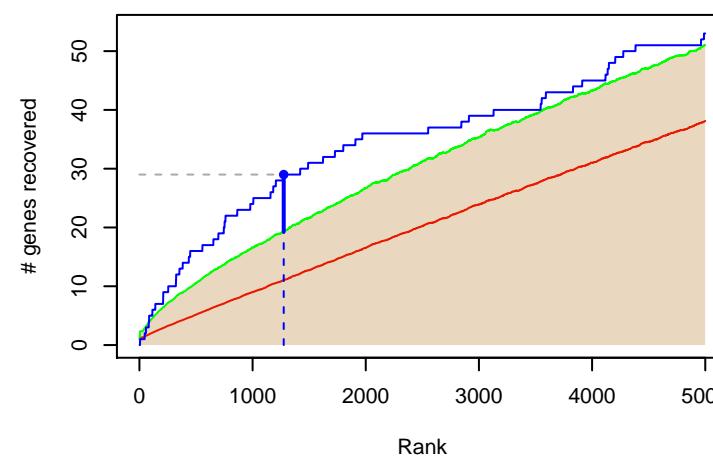
flyfactorsurvey__pho_SANGER_10_FBgn002521



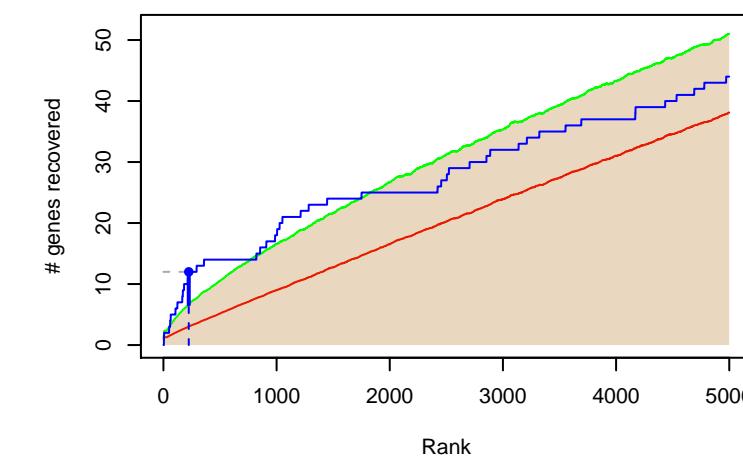
transfac_pro__M07514



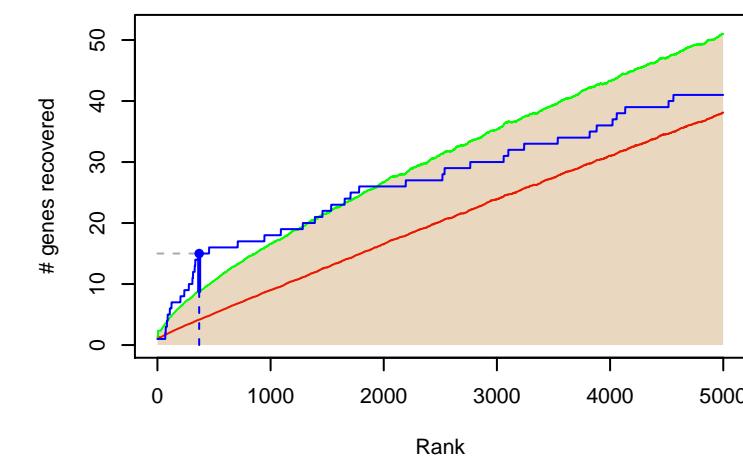
cisbp__M4652



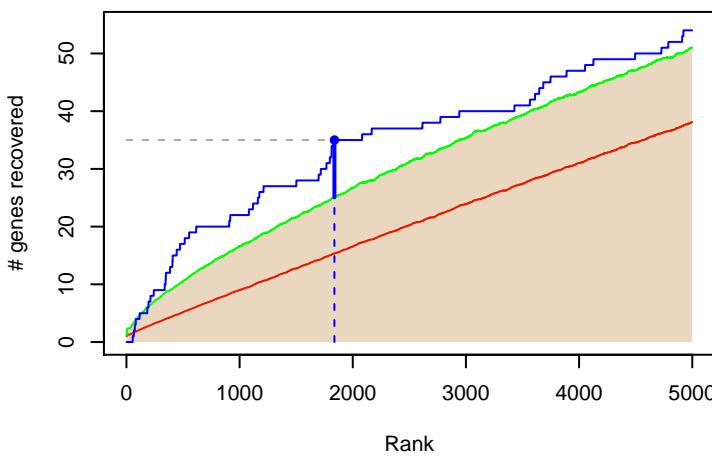
hocomoco__ZFP42_HUMAN.H11MO.0.A



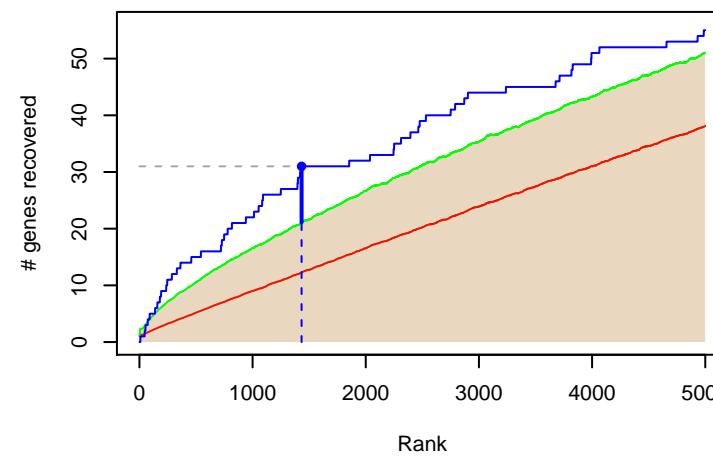
dbcorrdb__POLR2A__ENCSR000BHN_1_m1



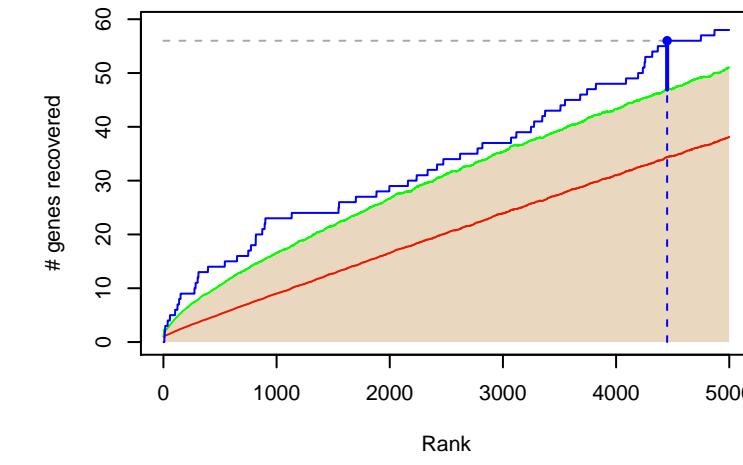
predrem__nrMotif233



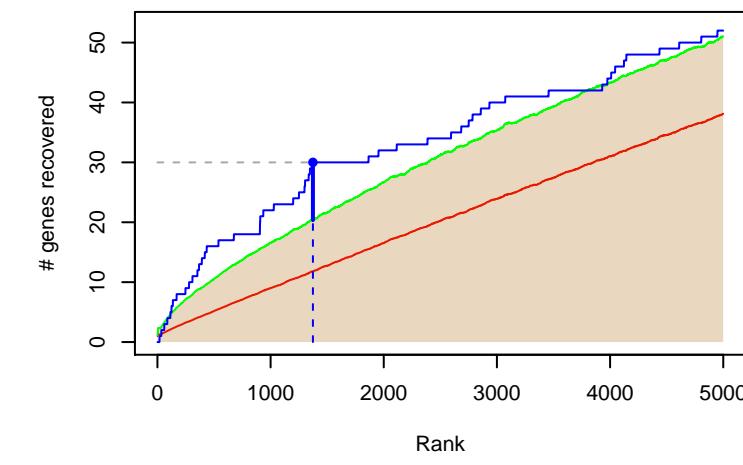
transfac_pro__M02044

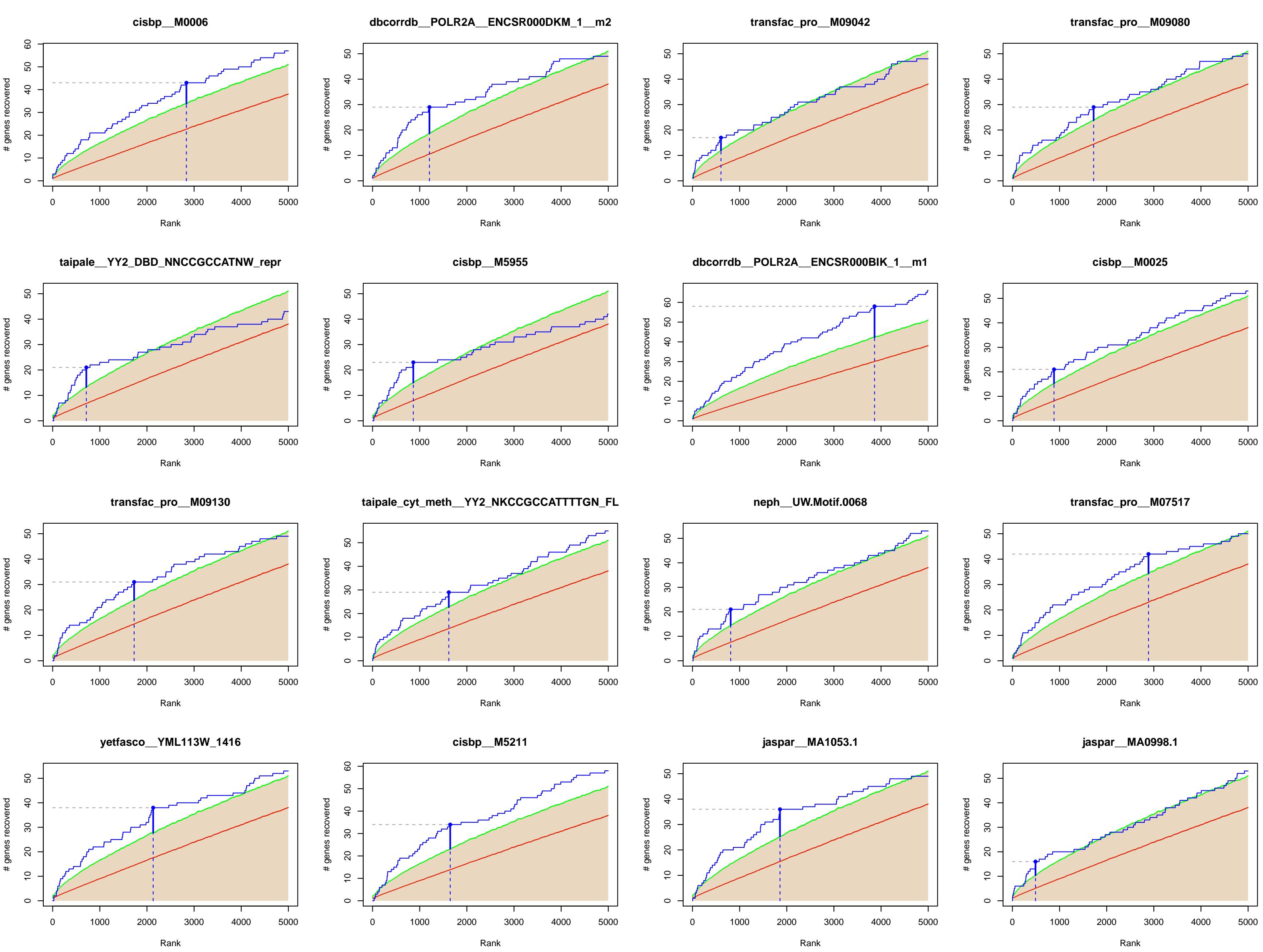


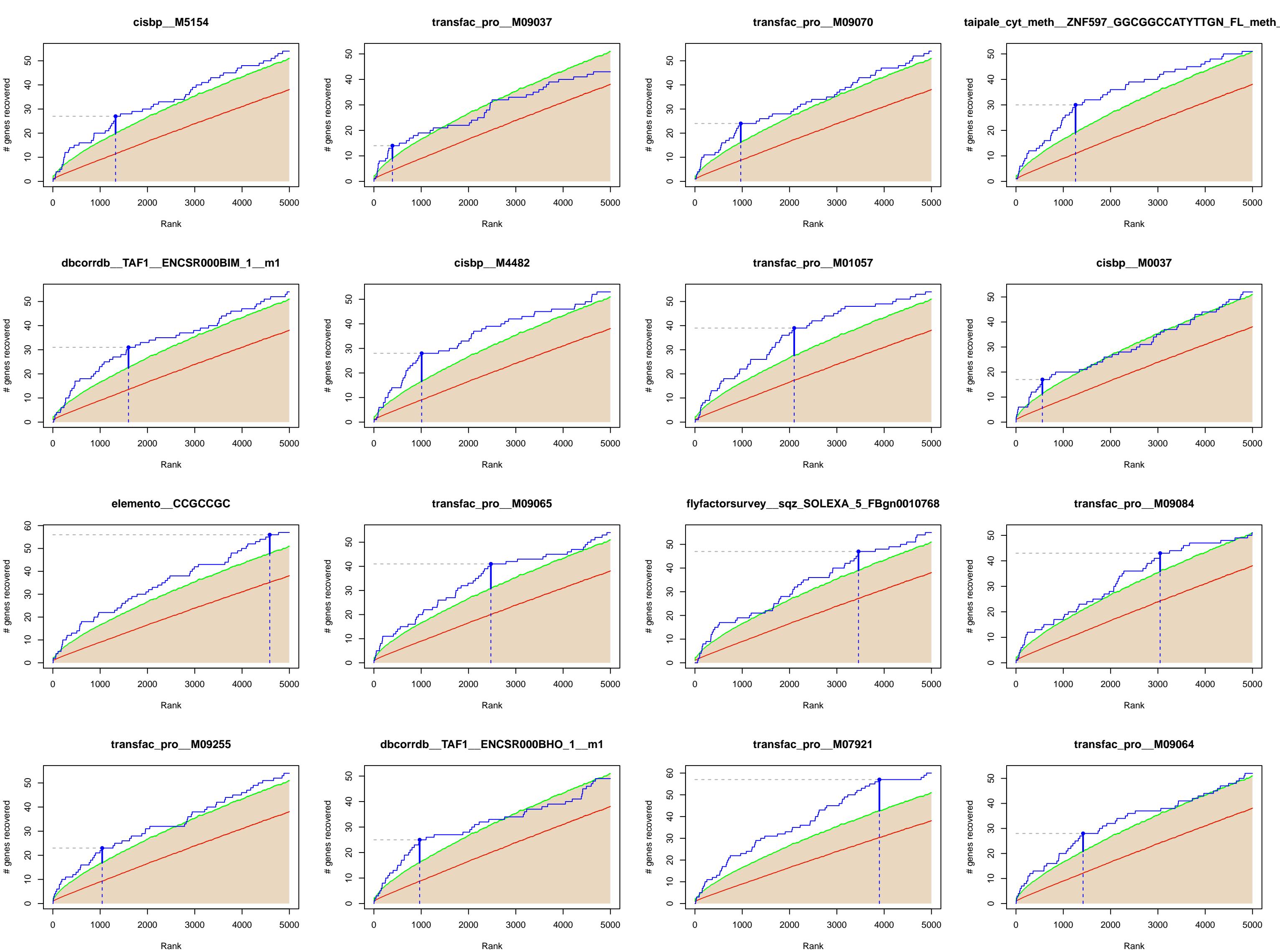
dbcorrdb__YY1__ENCSR000EXG_1_m2

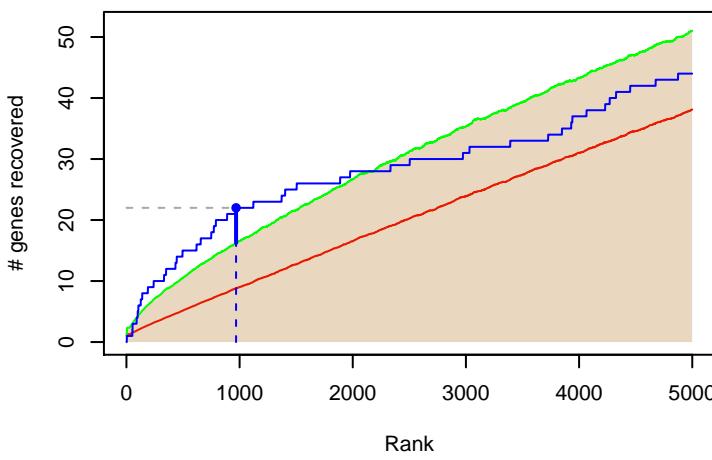
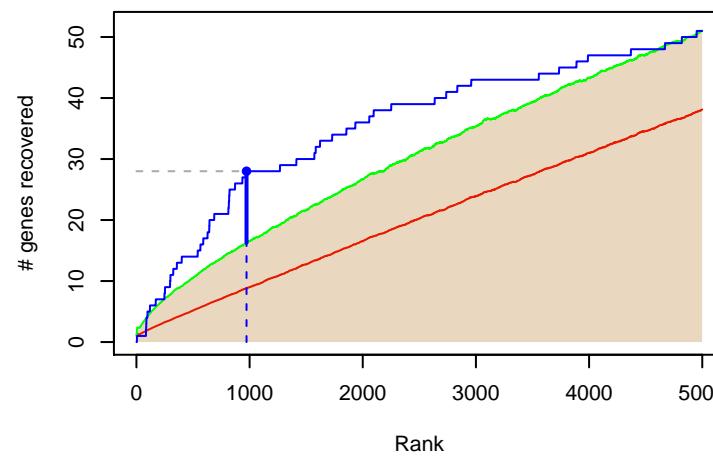
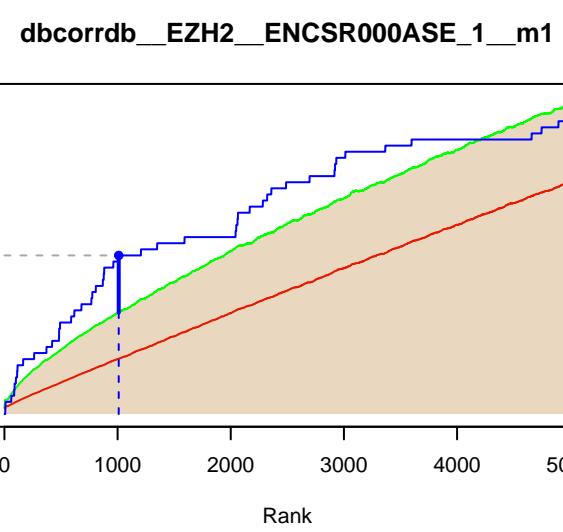
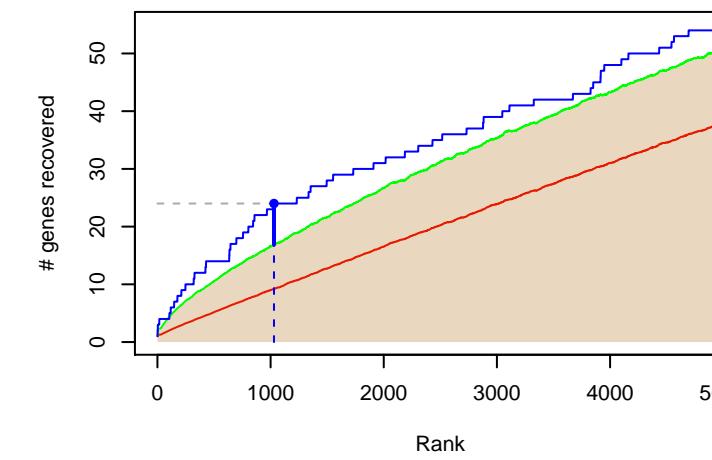
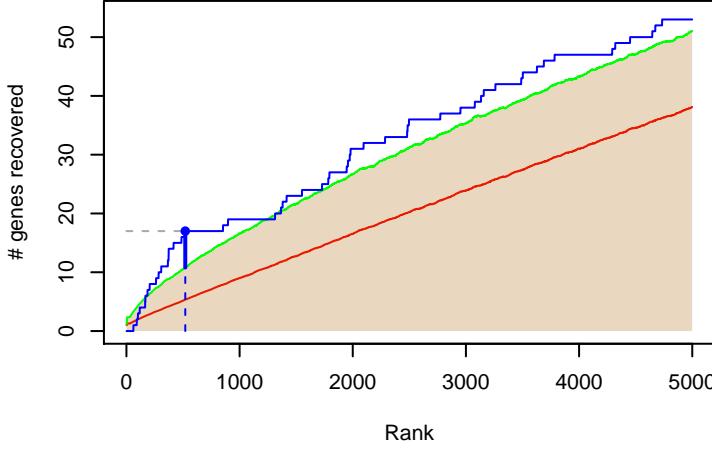
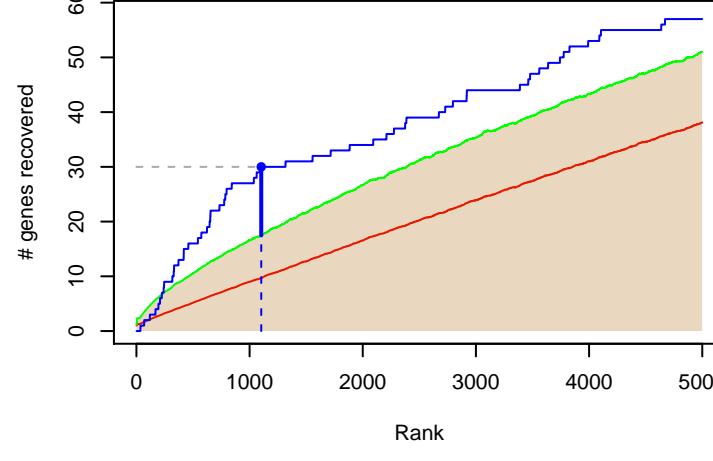
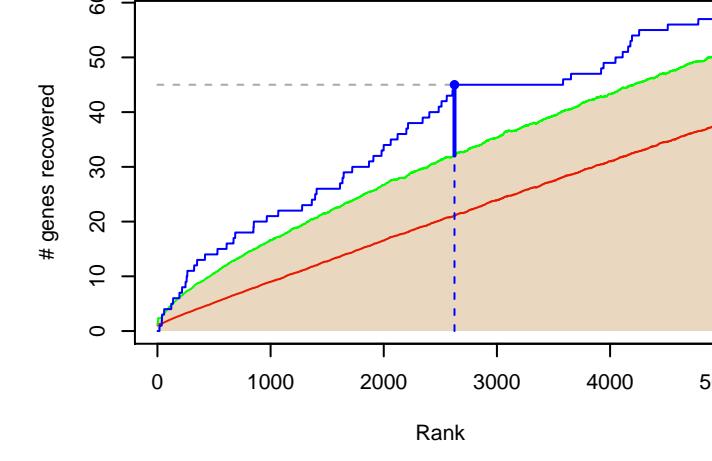
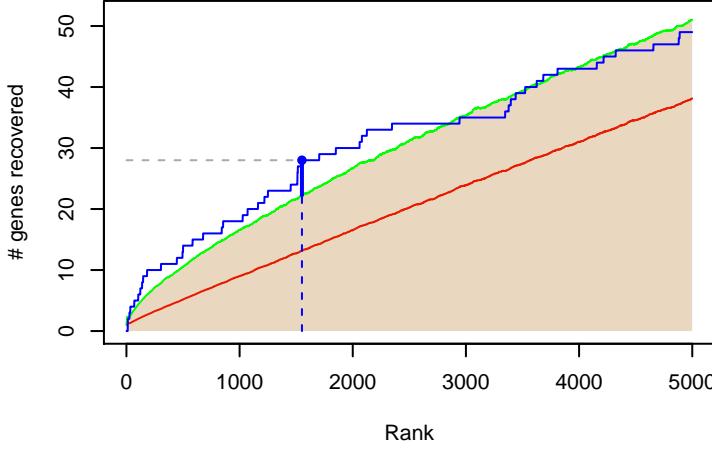
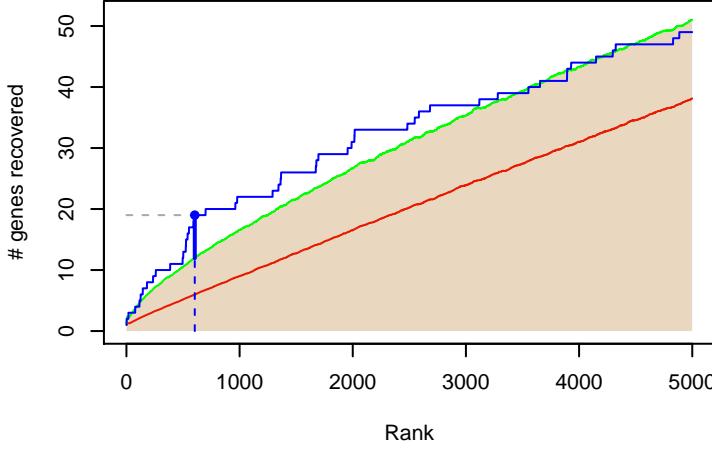
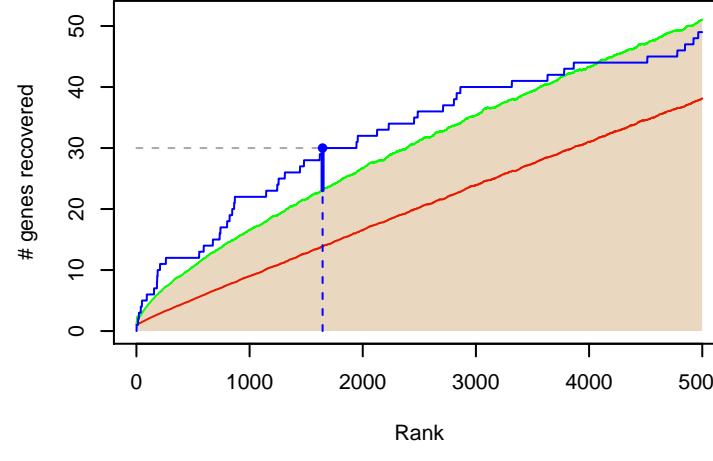
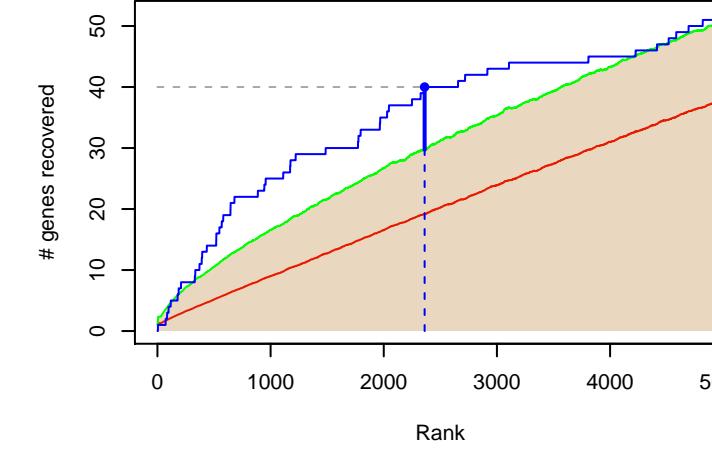
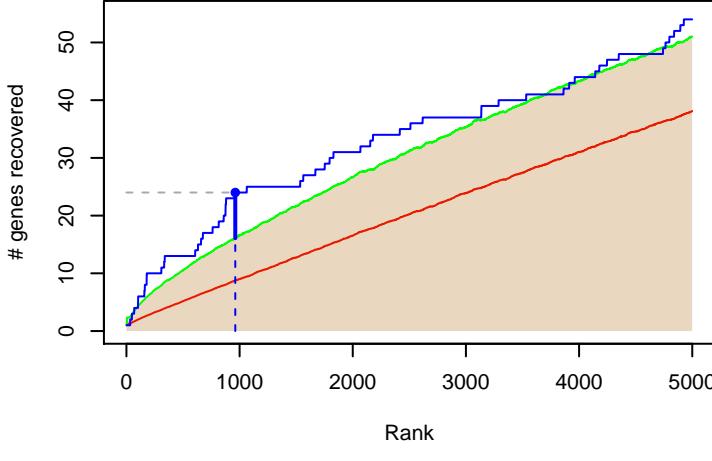
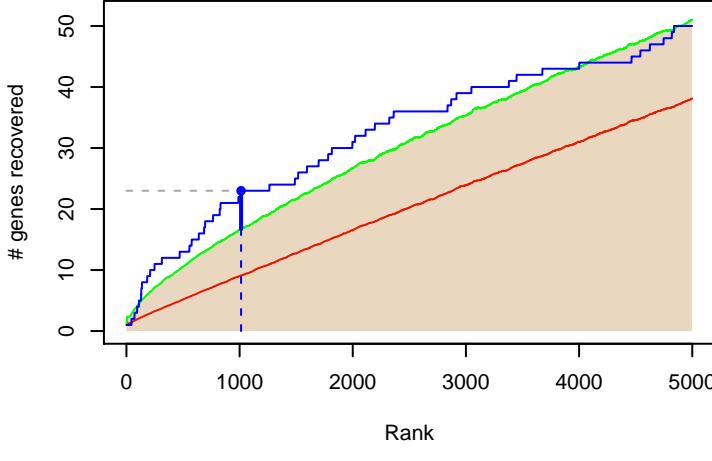
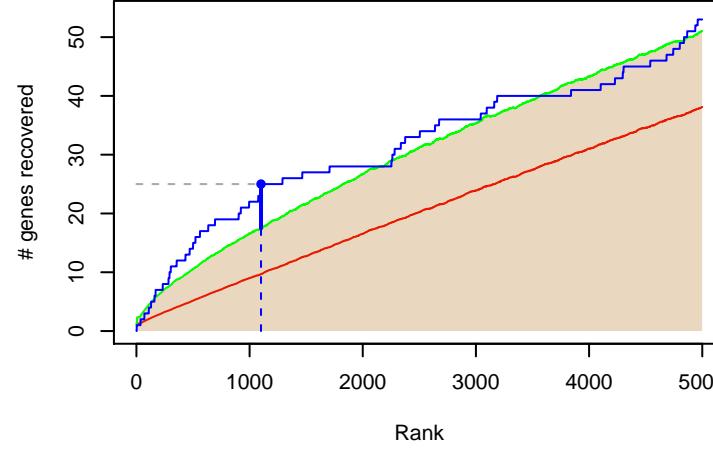
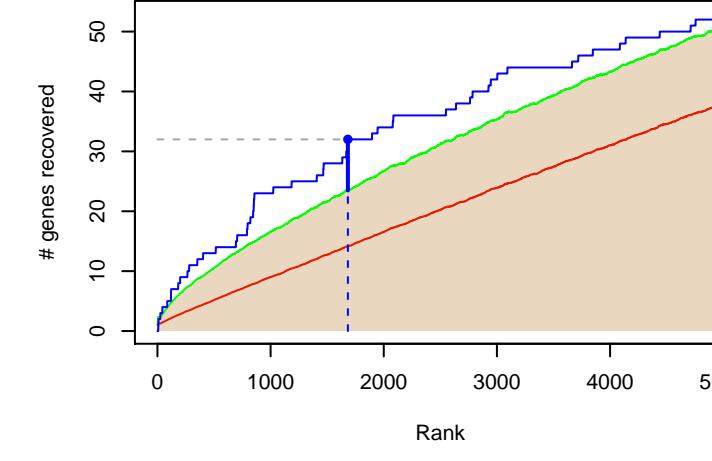
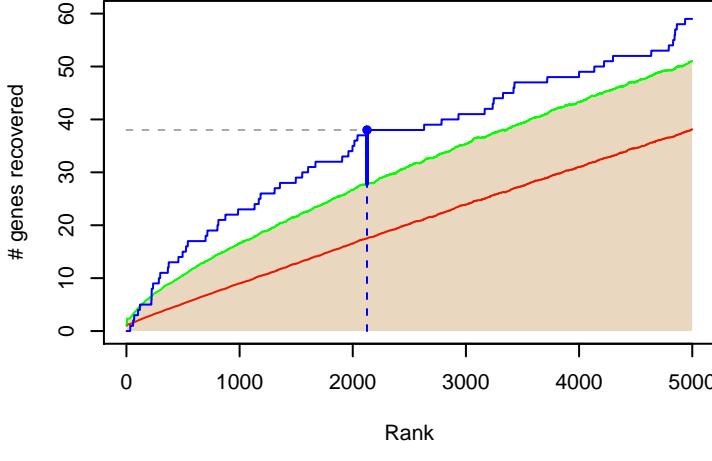


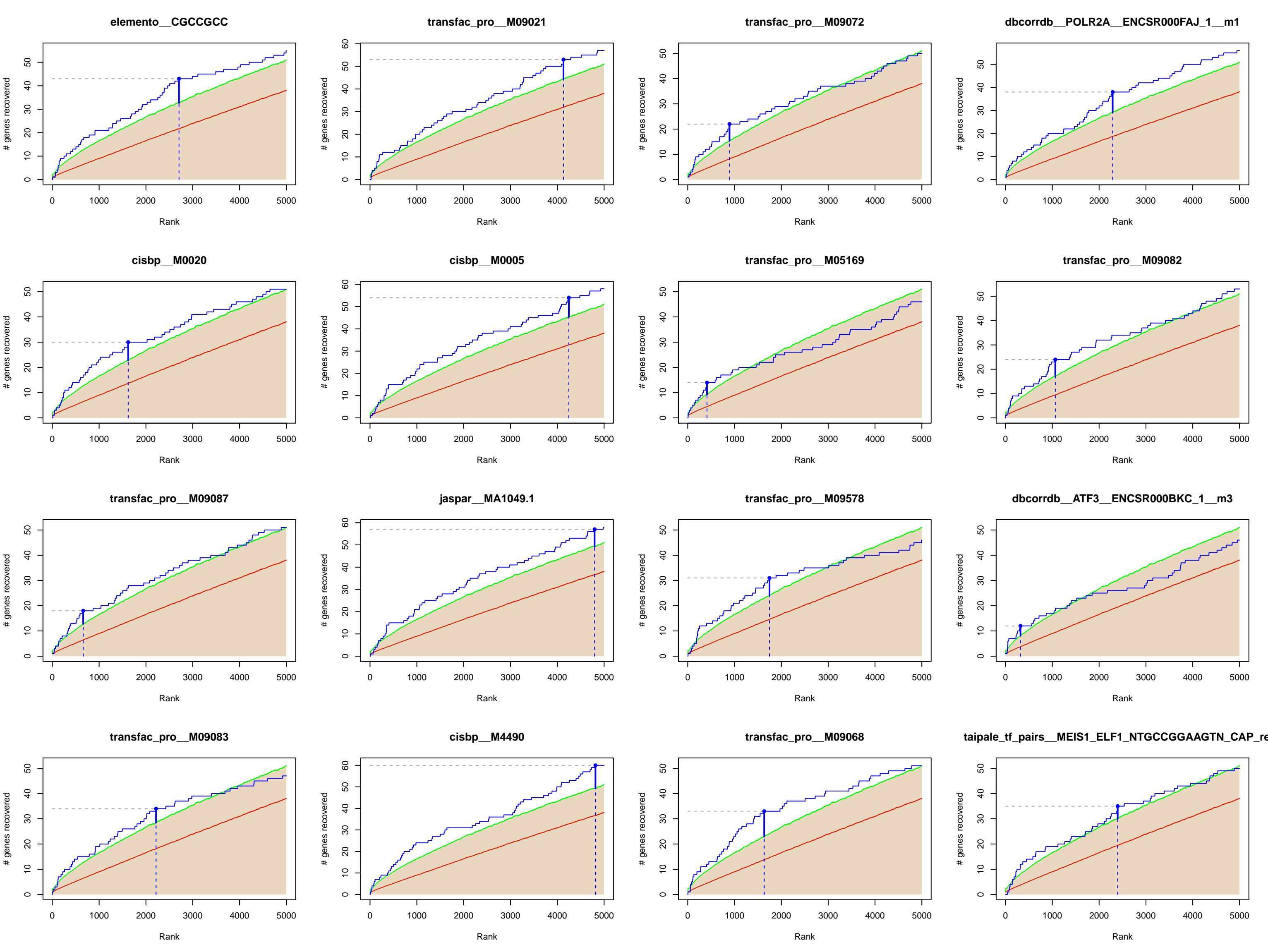
flyfactorsurvey__phol_SANGER_5_FBgn0035997

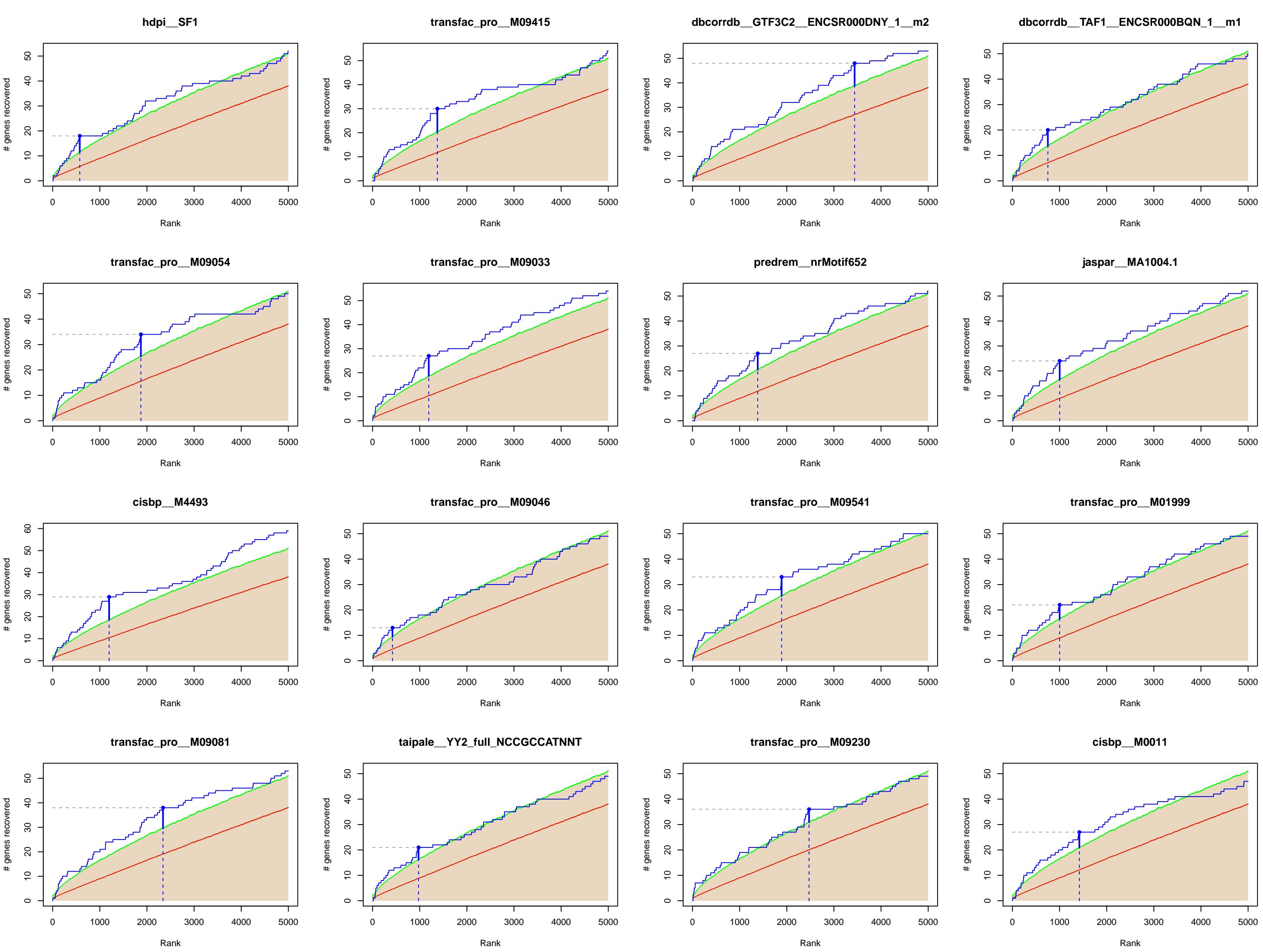


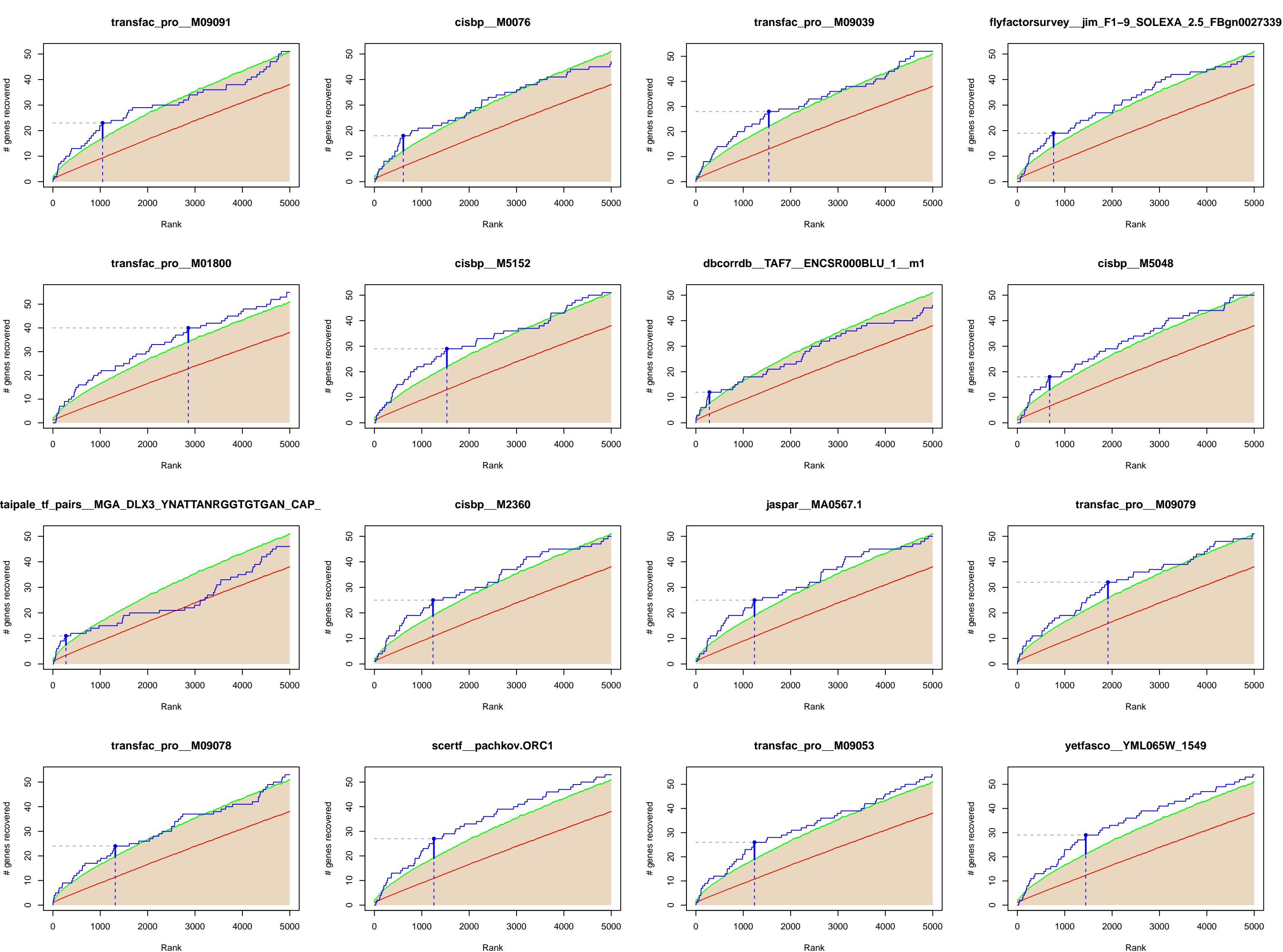


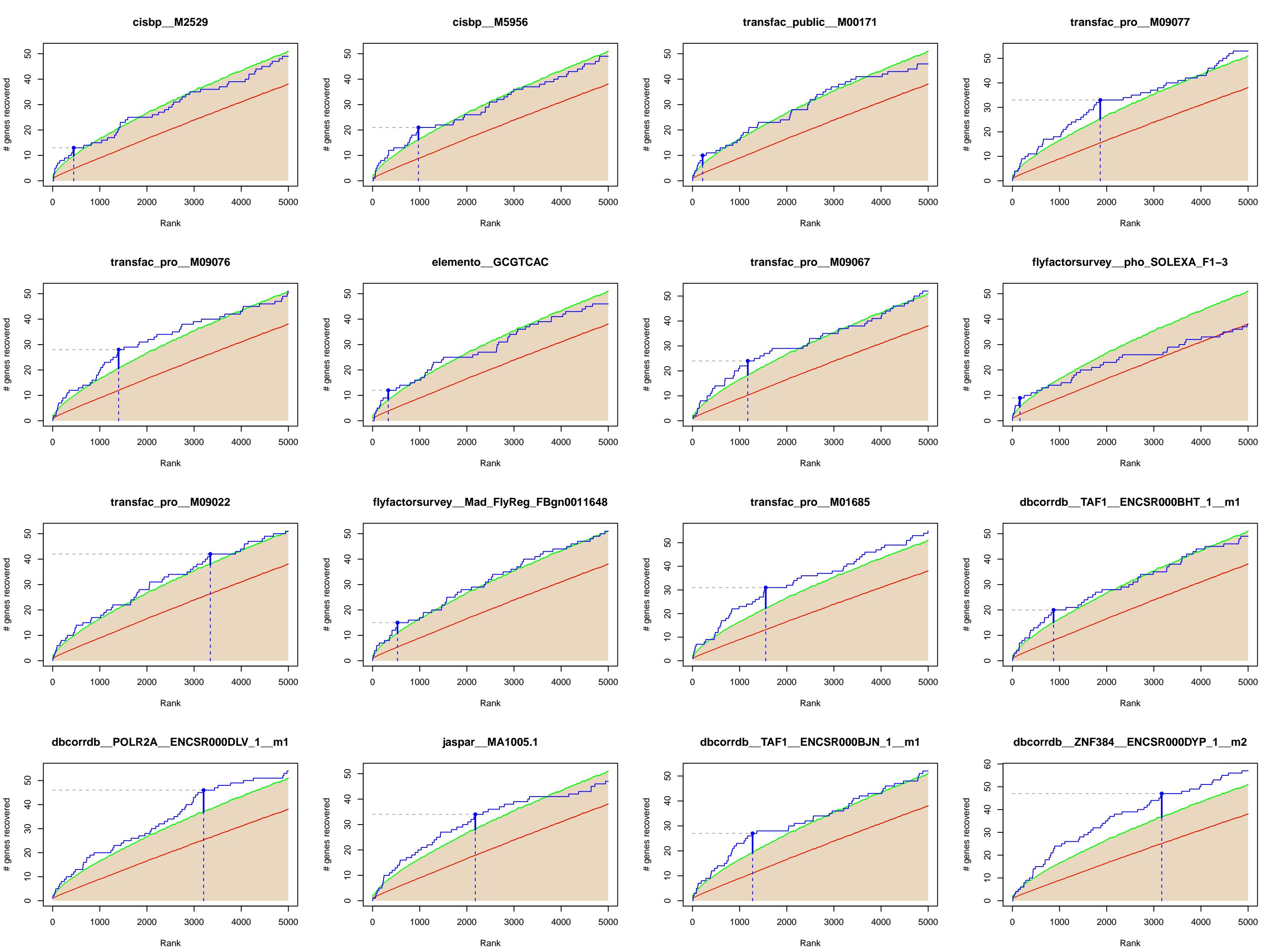


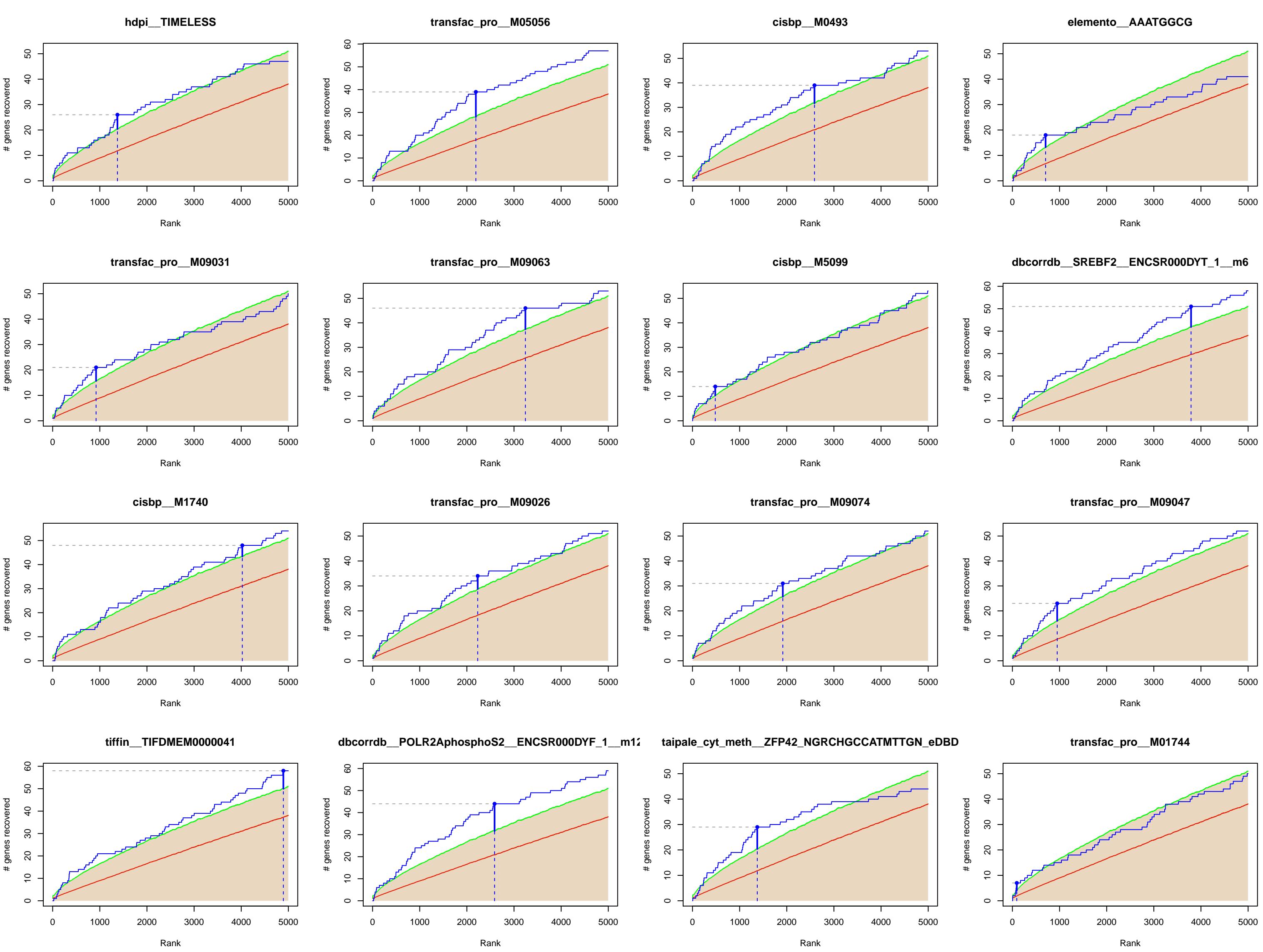
dbcrrdb YY1 ENCSR000BNP_1_m1**cisbp_M4516****transfac_pro_M09562****cisbp_M5212****dbcrrdb_POLR2AphosphoS5_ENCSR000BIL_1_m1****dbcrrdb_POLR2A_ENCSR000DPC_1_m1****transfac_pro_M04770****transfac_pro_M09586****transfac_pro_M09071****cisbp_M4603****transfac_pro_M09066****cisbp_M0004****transfac_pro_M07520****transfac_pro_M09069****flyfactorsurvey_sqz_SANGER_5_FBgn0010768**



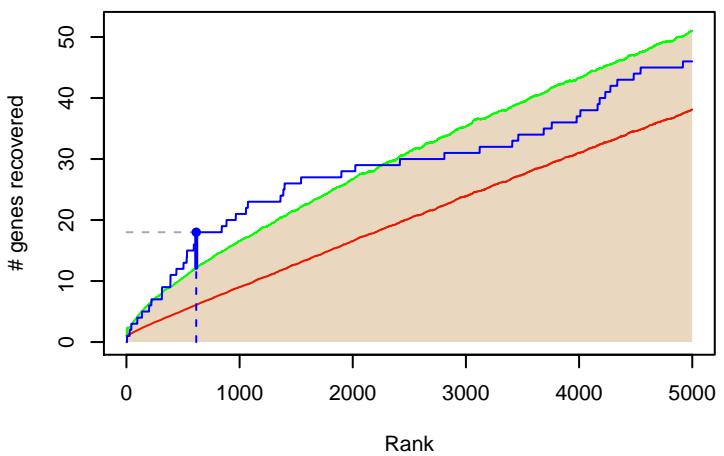




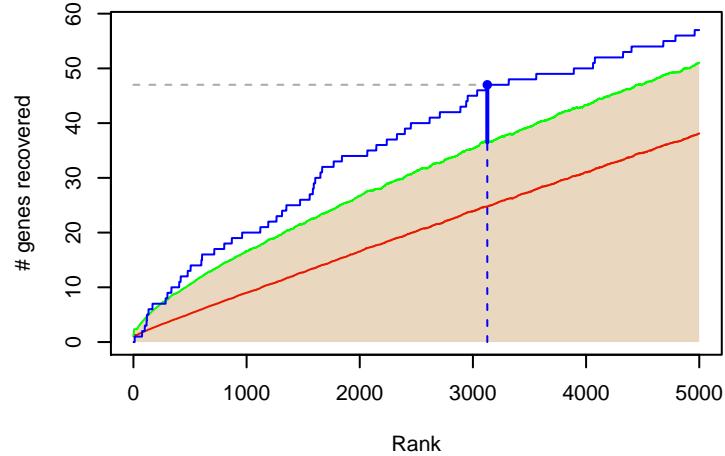




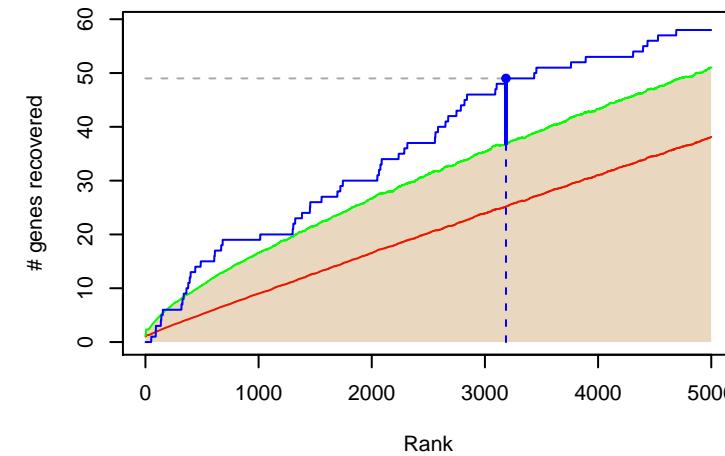
transfac_pro_M09045



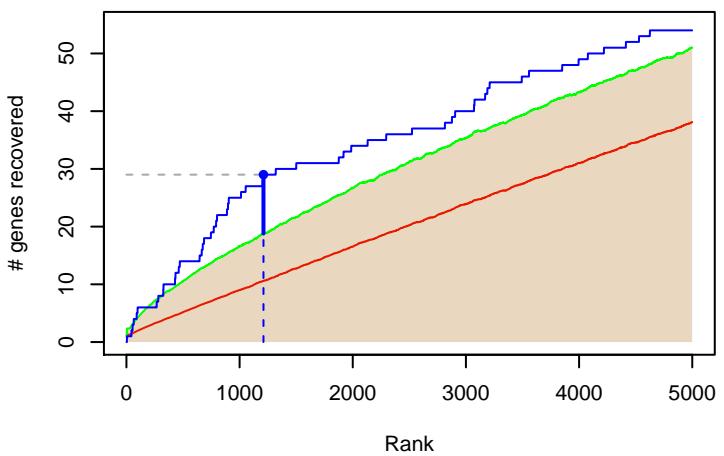
dbcrrdb_POLR2A_ENCSR000FAY_1_m1



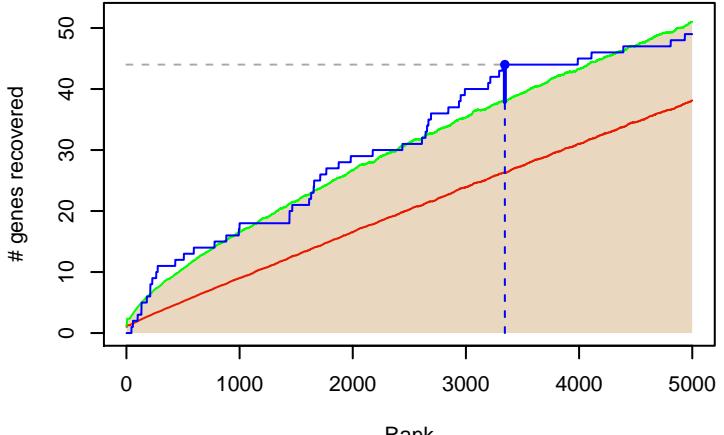
dbcrrdb_POLR2AphosphoS2_ENCSR000ECT_1_m1



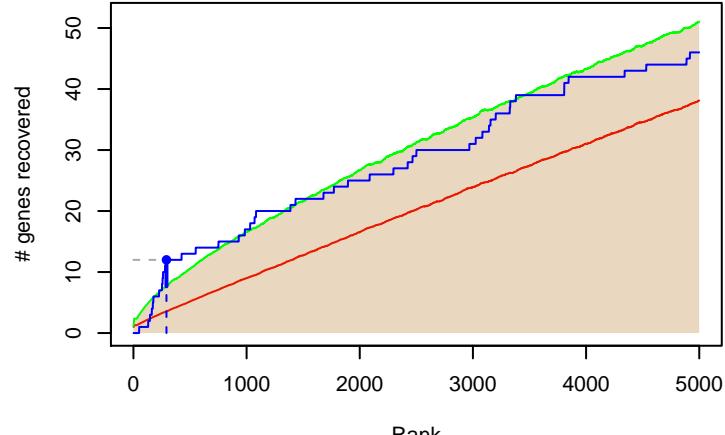
factorbook YY1



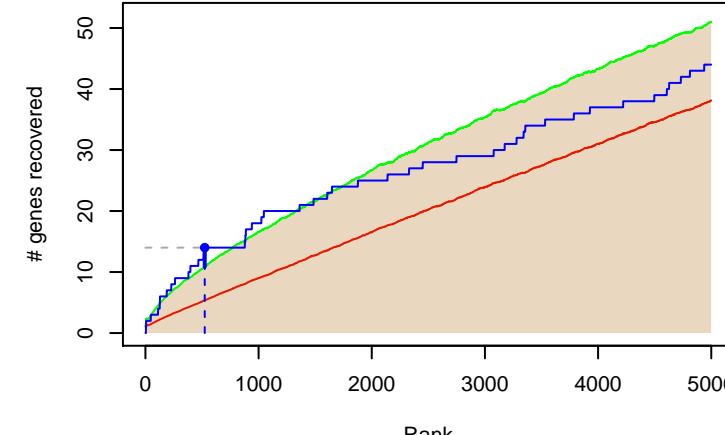
transfac_pro_M06351



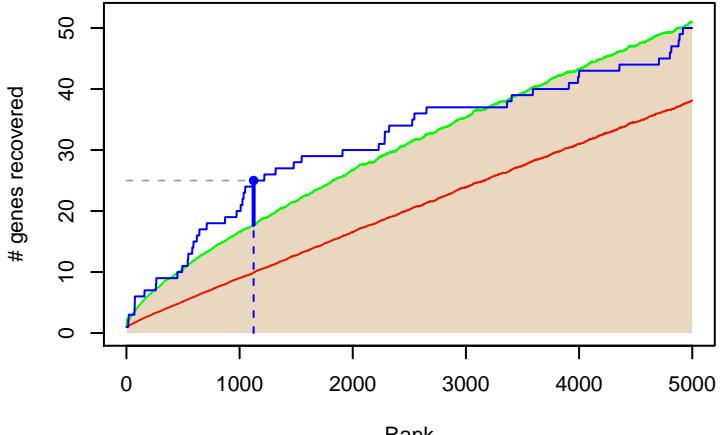
taipale_tf_pairs_E2F3_FOXI1_NNMCACCGCGCCMN_CAP_



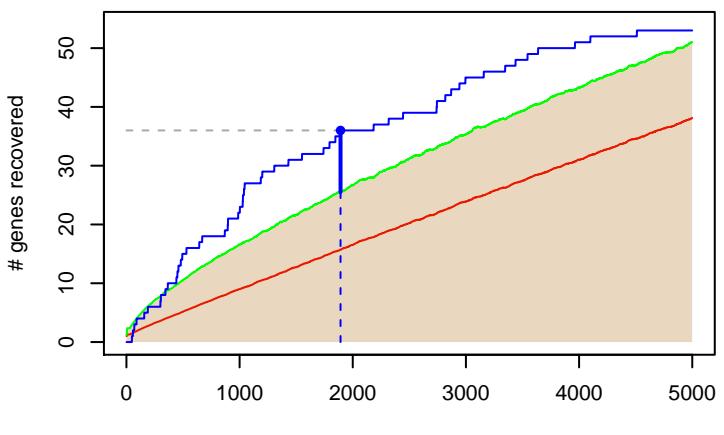
transfac_pro_M05538



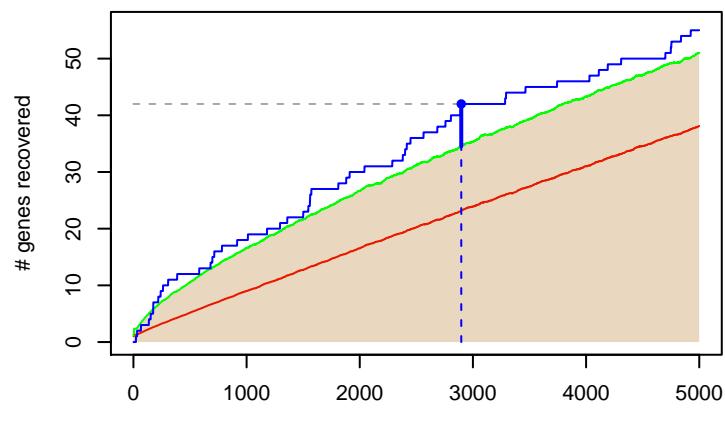
transfac_pro_M09090



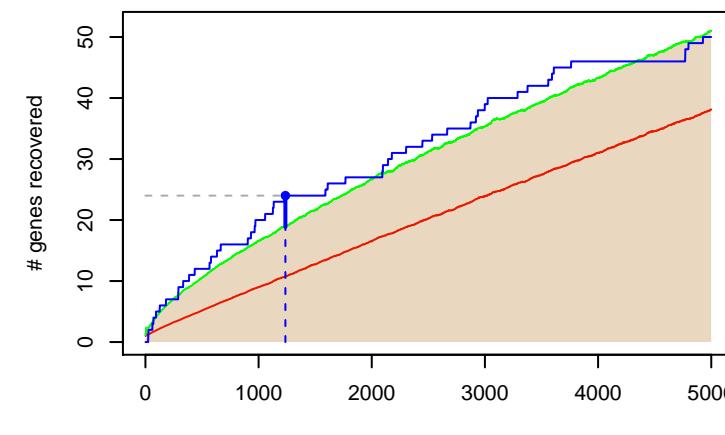
dbcrrdb_TAF1_ENCSR000BKS_1_m2



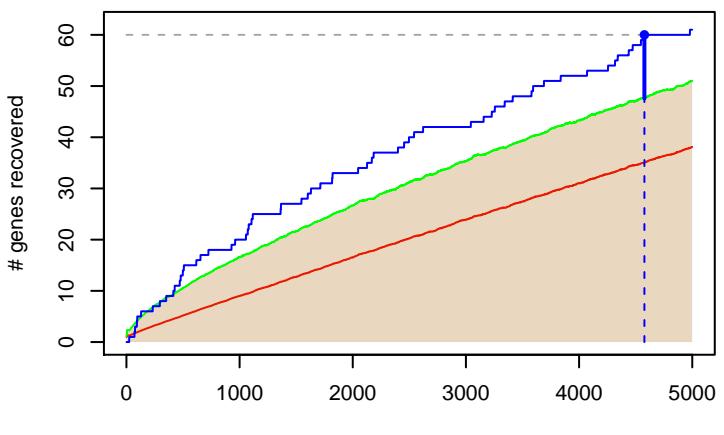
dbcrrdb_POLR2A_ENCSR000EAR_1_m1



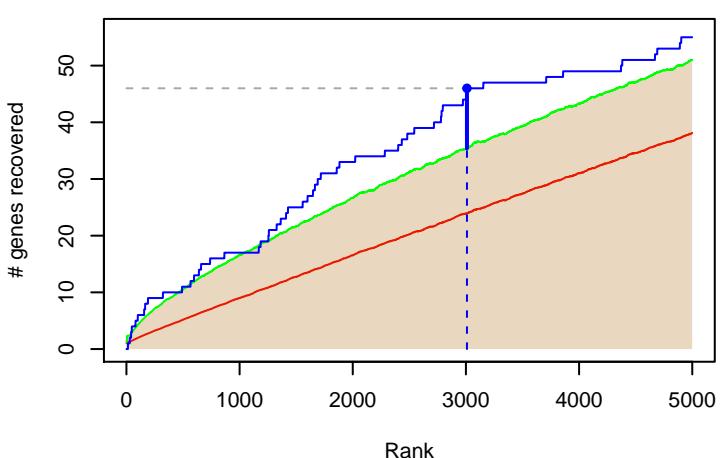
transfac_pro_M06352



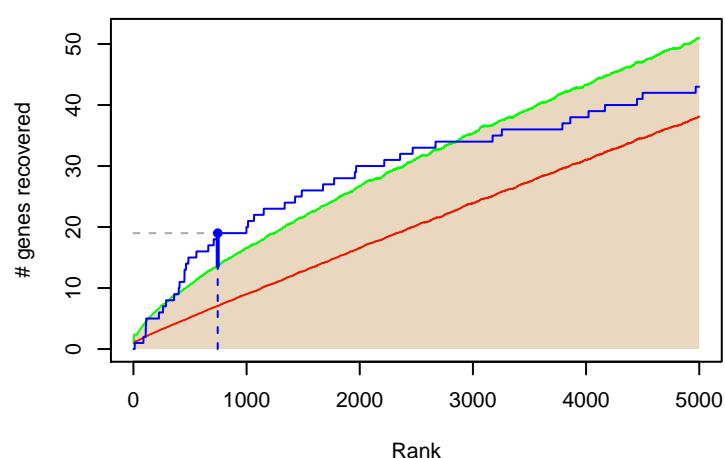
dbcrrdb_POLR2A_ENCSR000DKT_1_m1



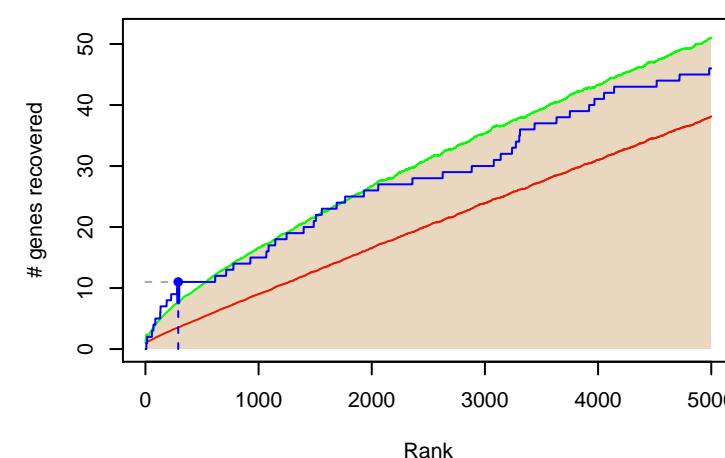
elemento_CGGCGGC



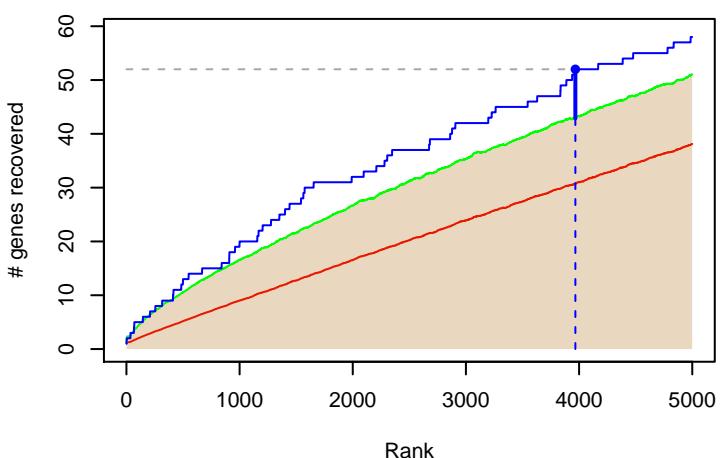
neph_UW.Motif.0390



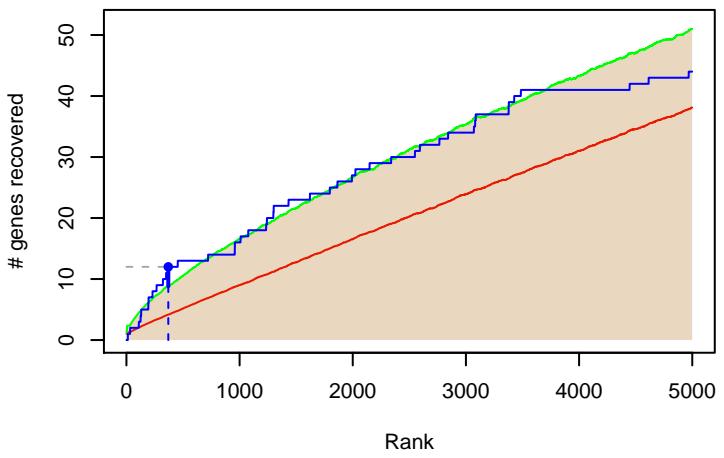
transfac_pro_M09028



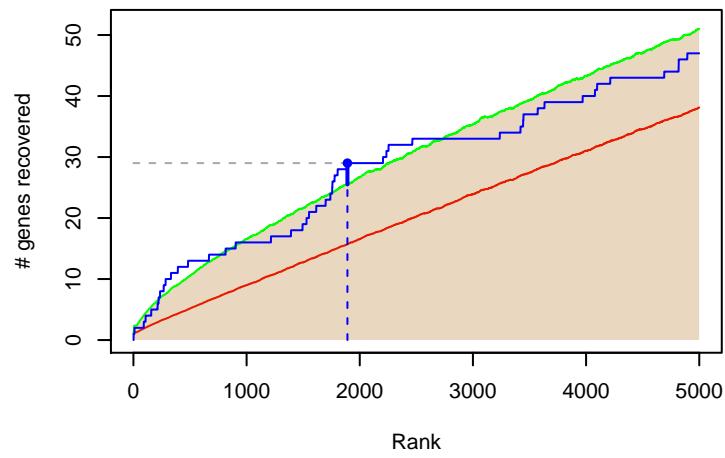
dbcrrdb_YY1_ENCSR000EWF_1_m2



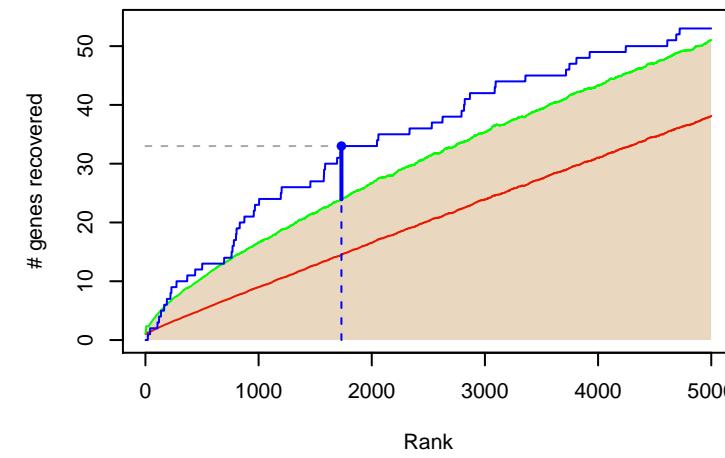
elemento_ACGACGA



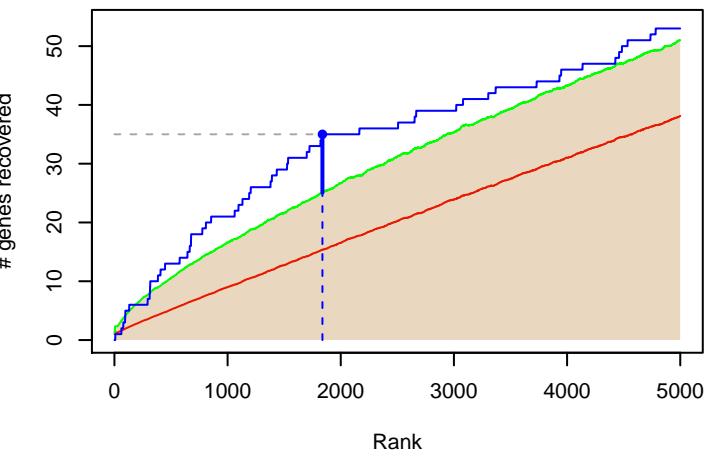
transfac_pro_M05712



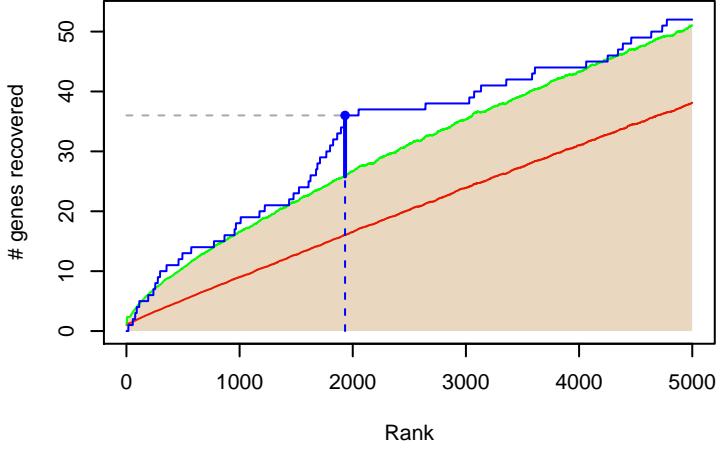
swissregulon_sacCer_ORC1



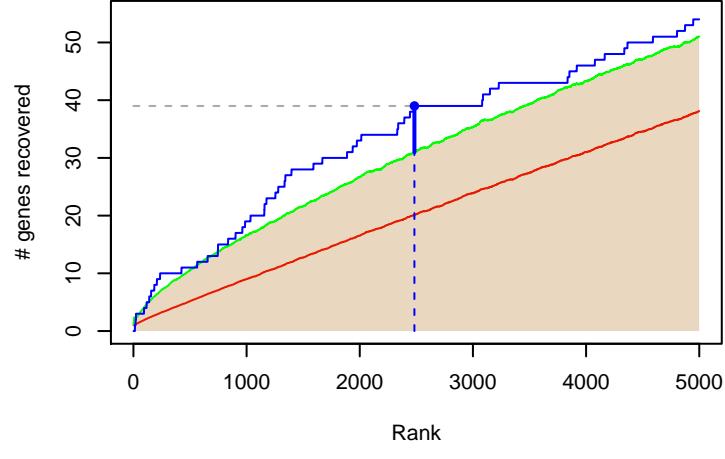
jaspar_MA1052.1



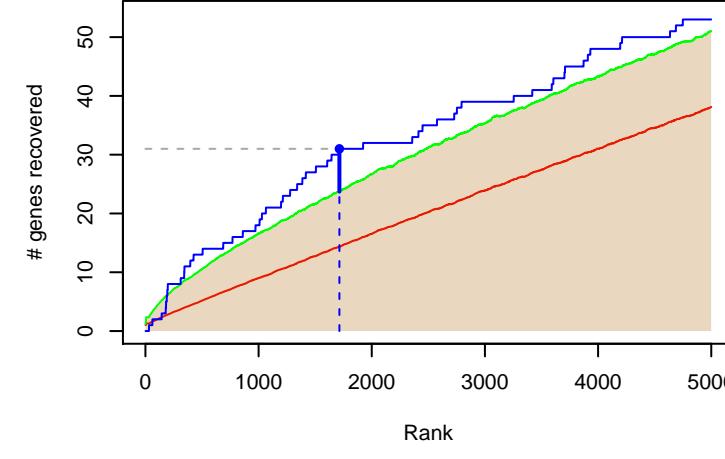
tfdimers_MD00454



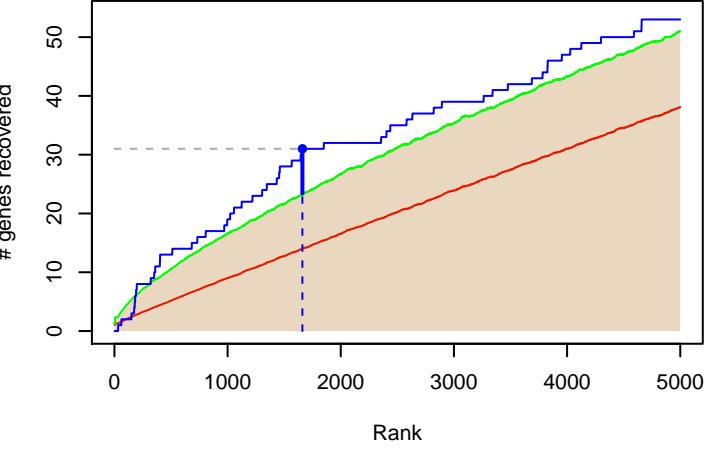
hdpi_NANOS1



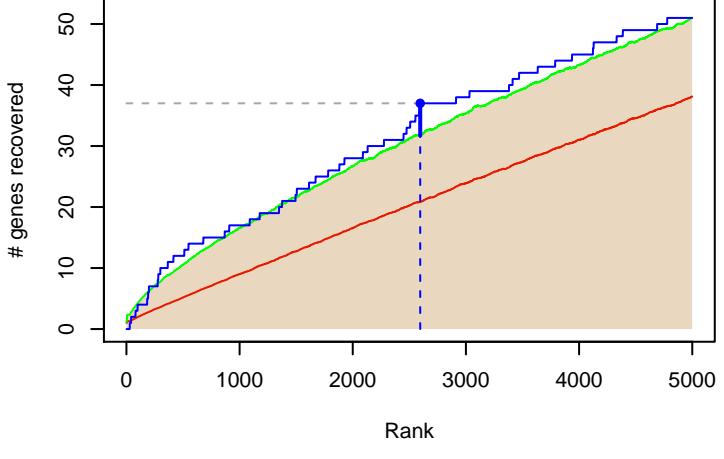
flyfactorsurvey_hb_FlyReg_FBgn0001180



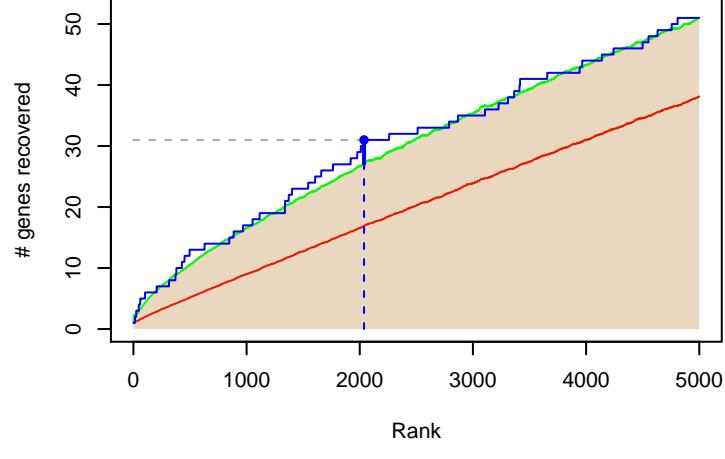
cisbp_M4997



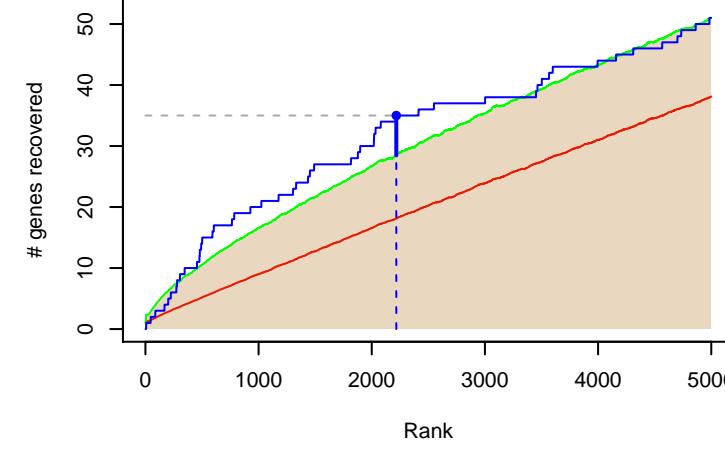
tfdimers_MD00356



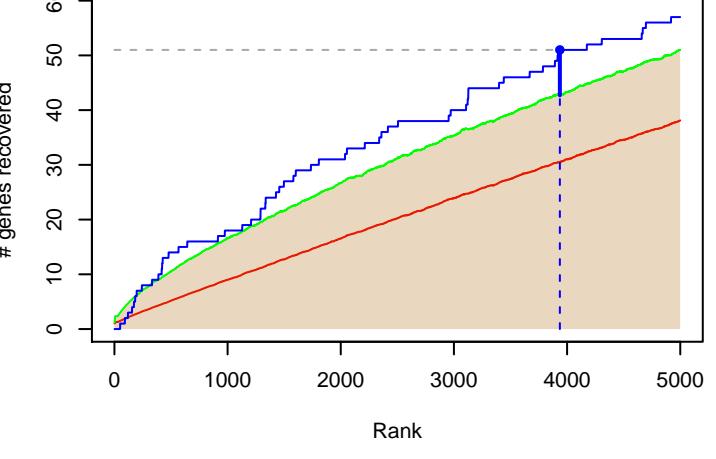
transfac_pro_M09089



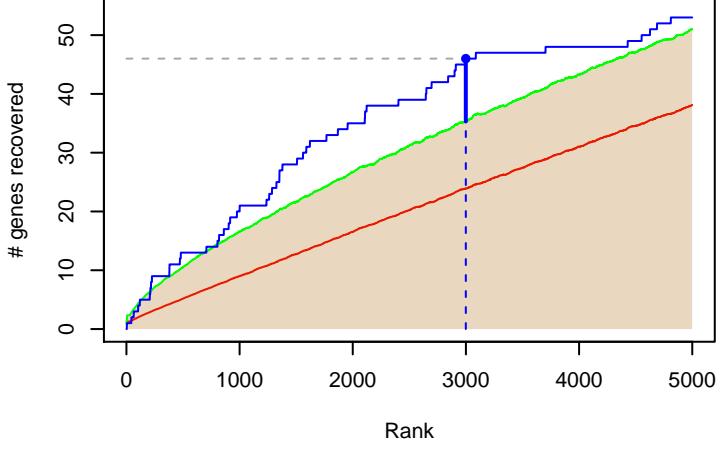
homer_CAAAGATGGCGGC_YY1



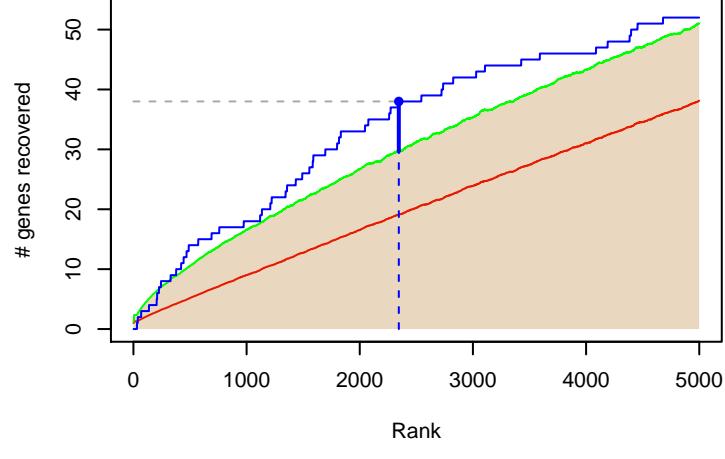
transfac_pro_M09167



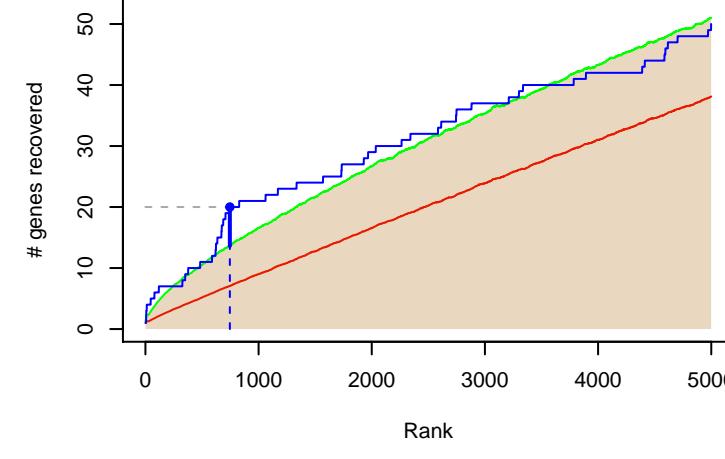
dbcrrdb_POLR2A_ENCSR000FAJ_1_m2



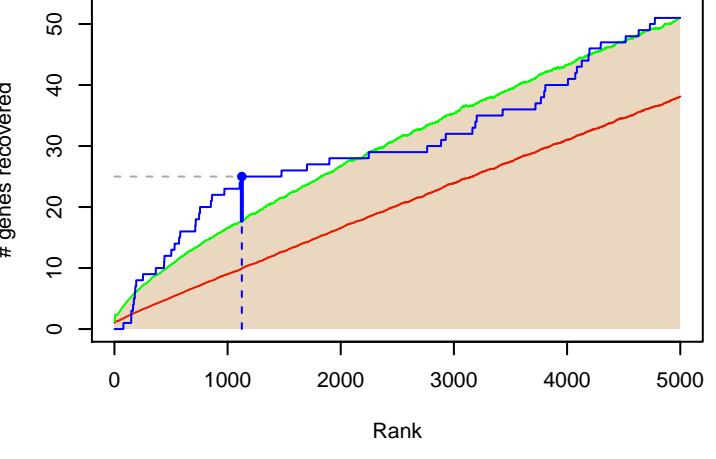
dbcrrdb_POLR2A_ENCSR000EAJ_1_m1

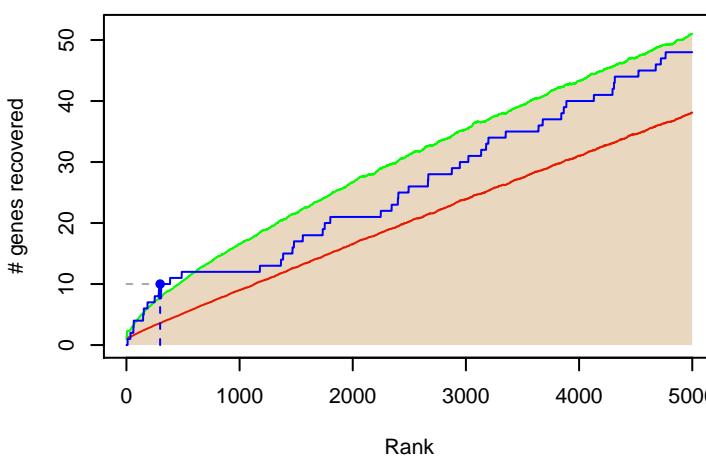
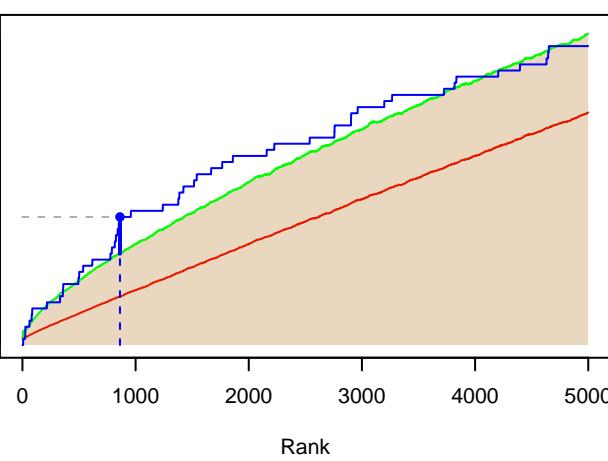
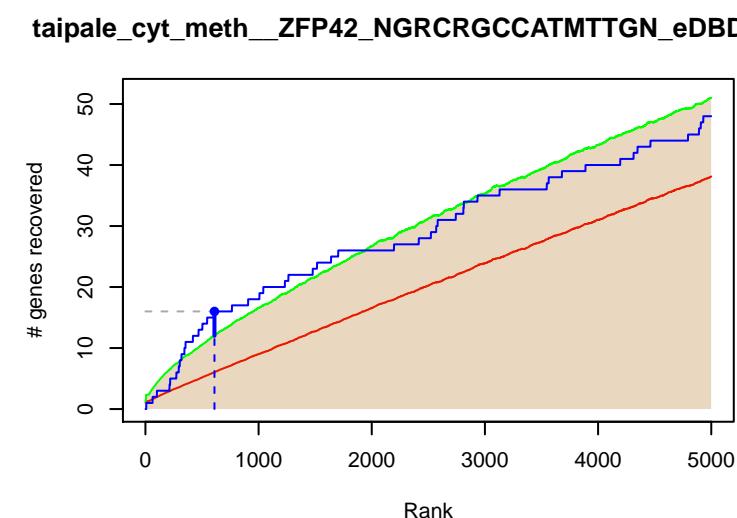
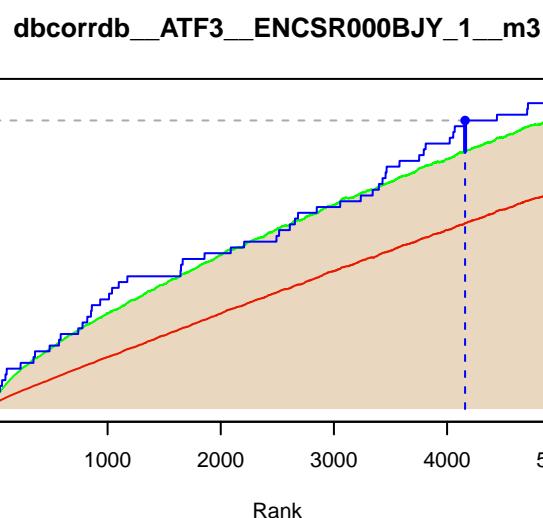
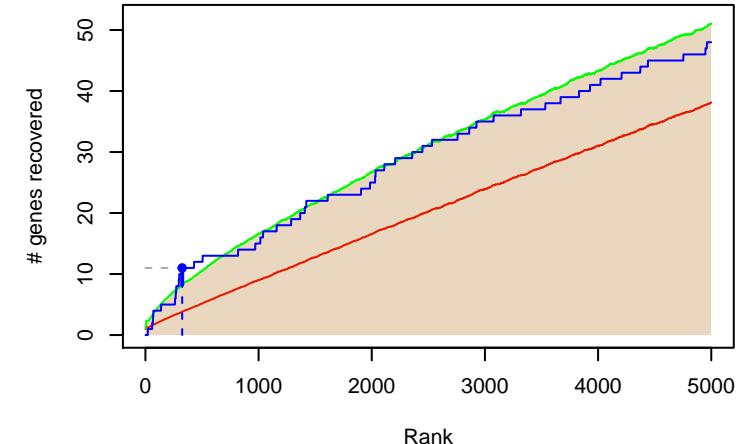
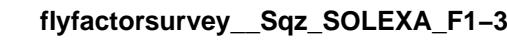
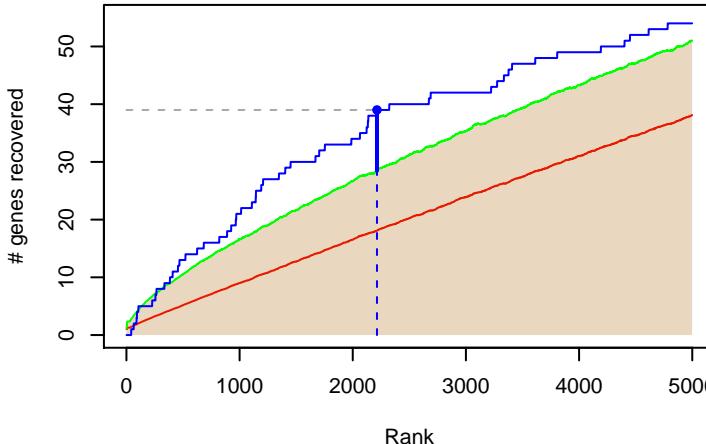
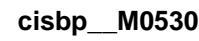
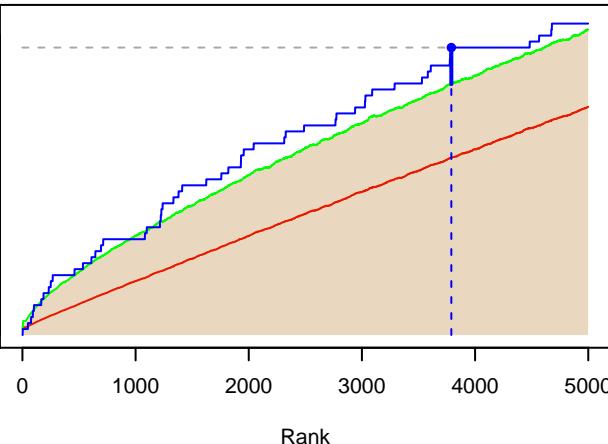
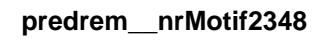
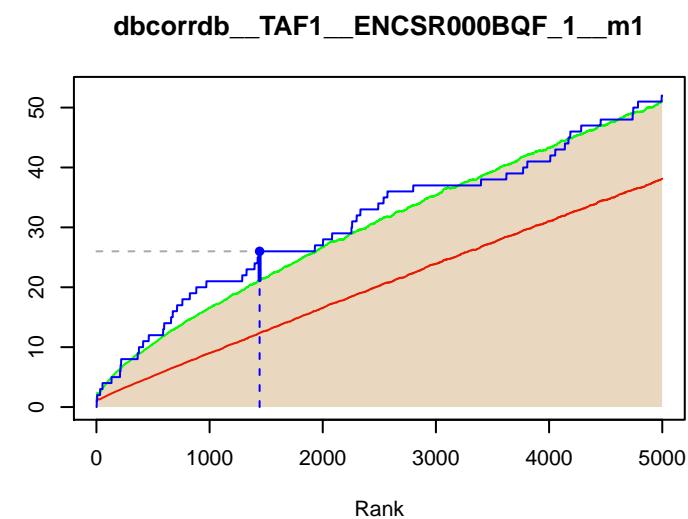
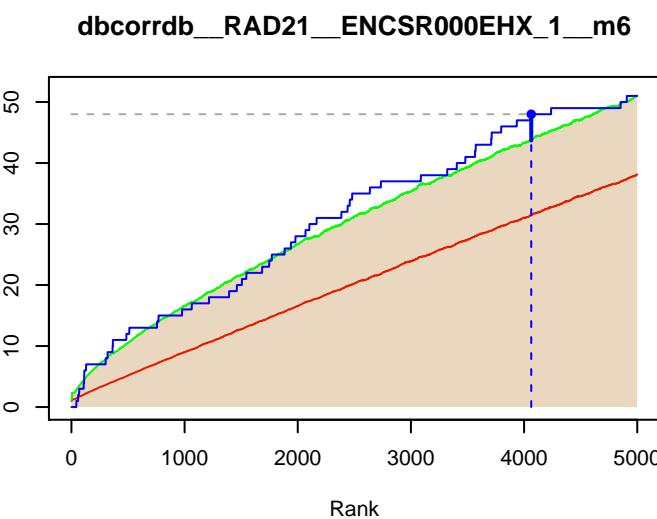
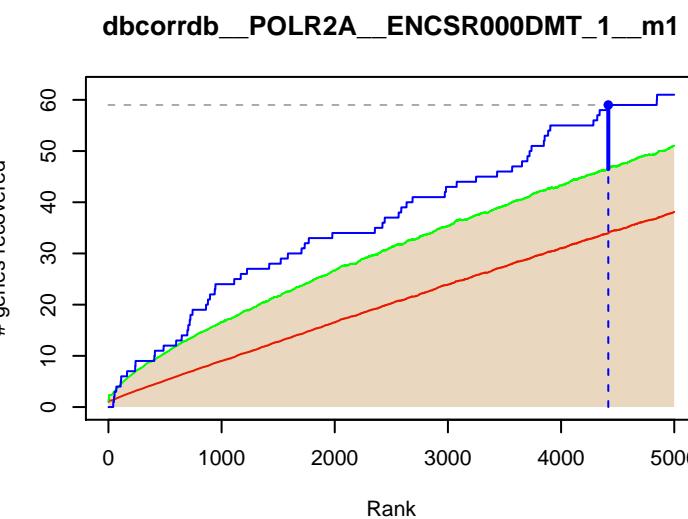
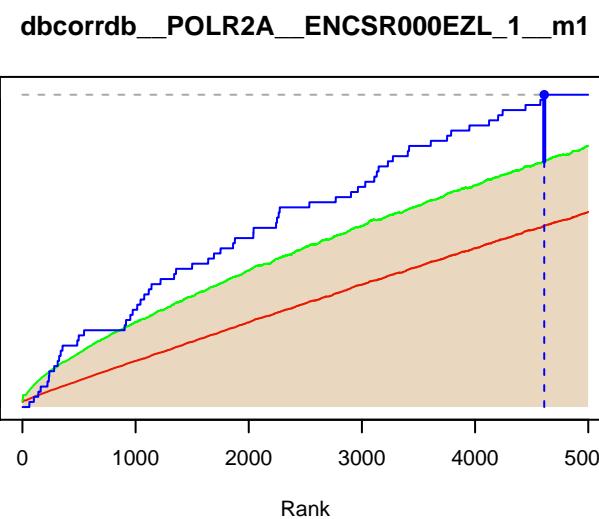


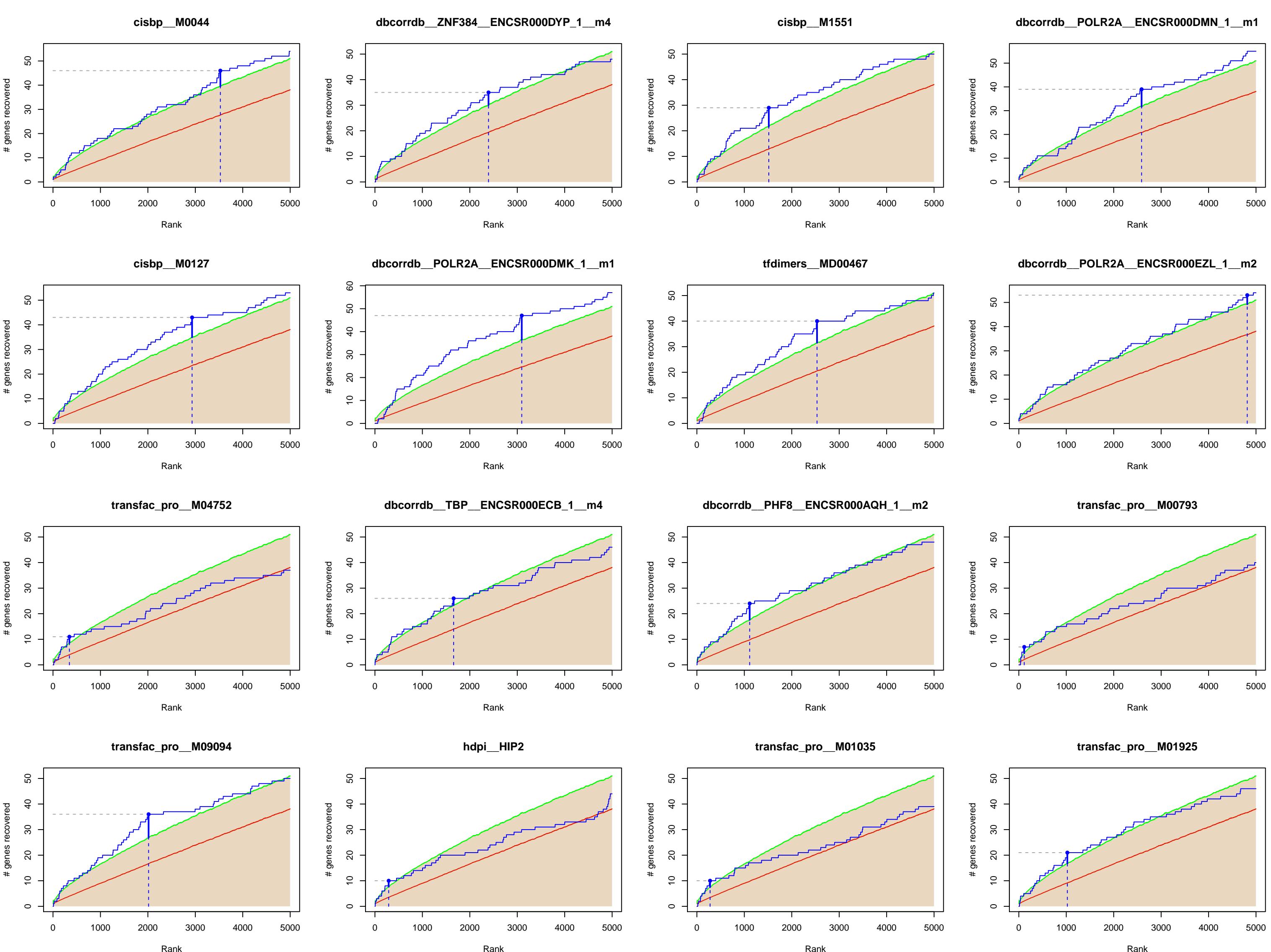
dbcrrdb_POLR2A_ENCSR000BQB_1_m2

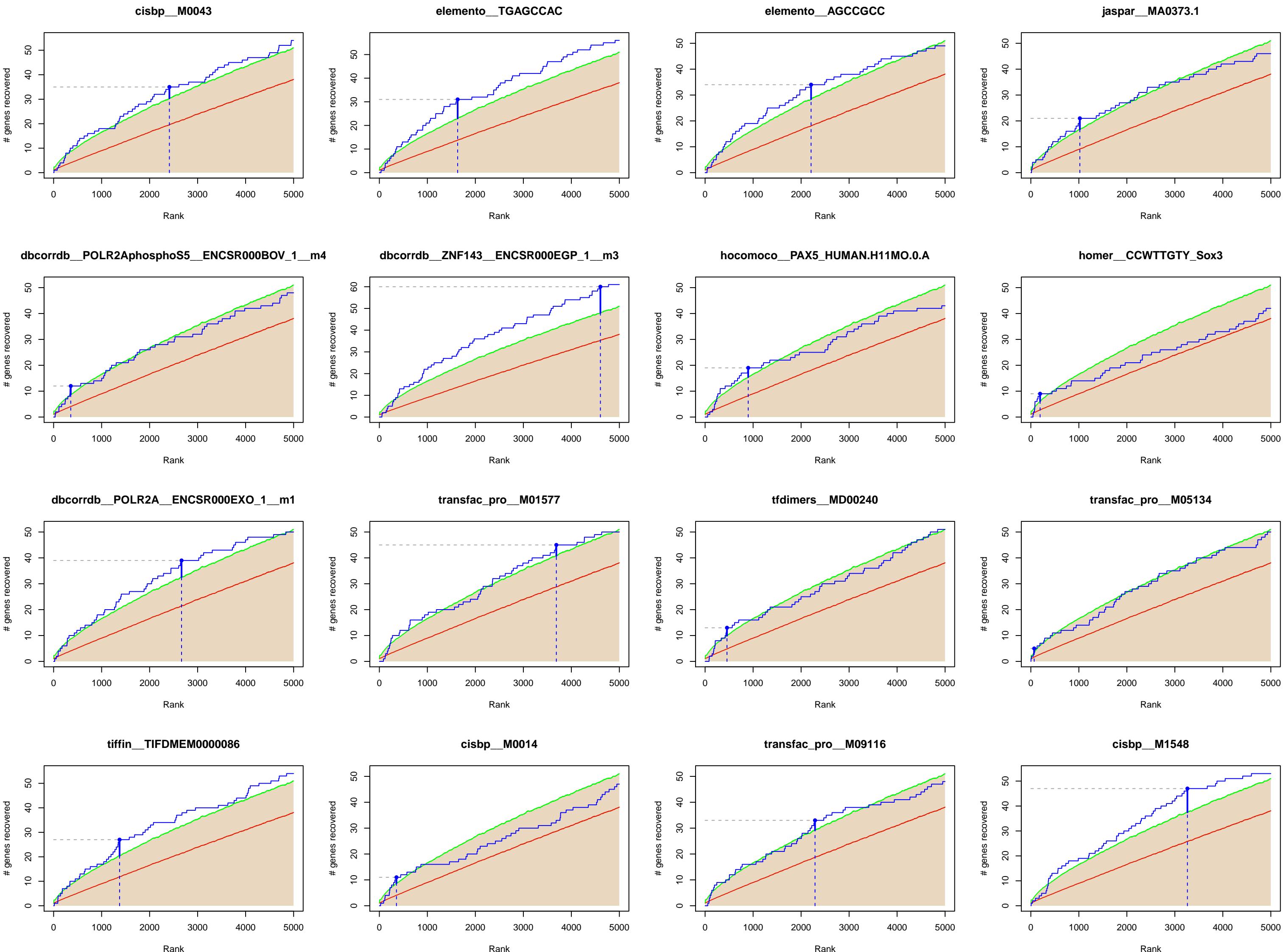


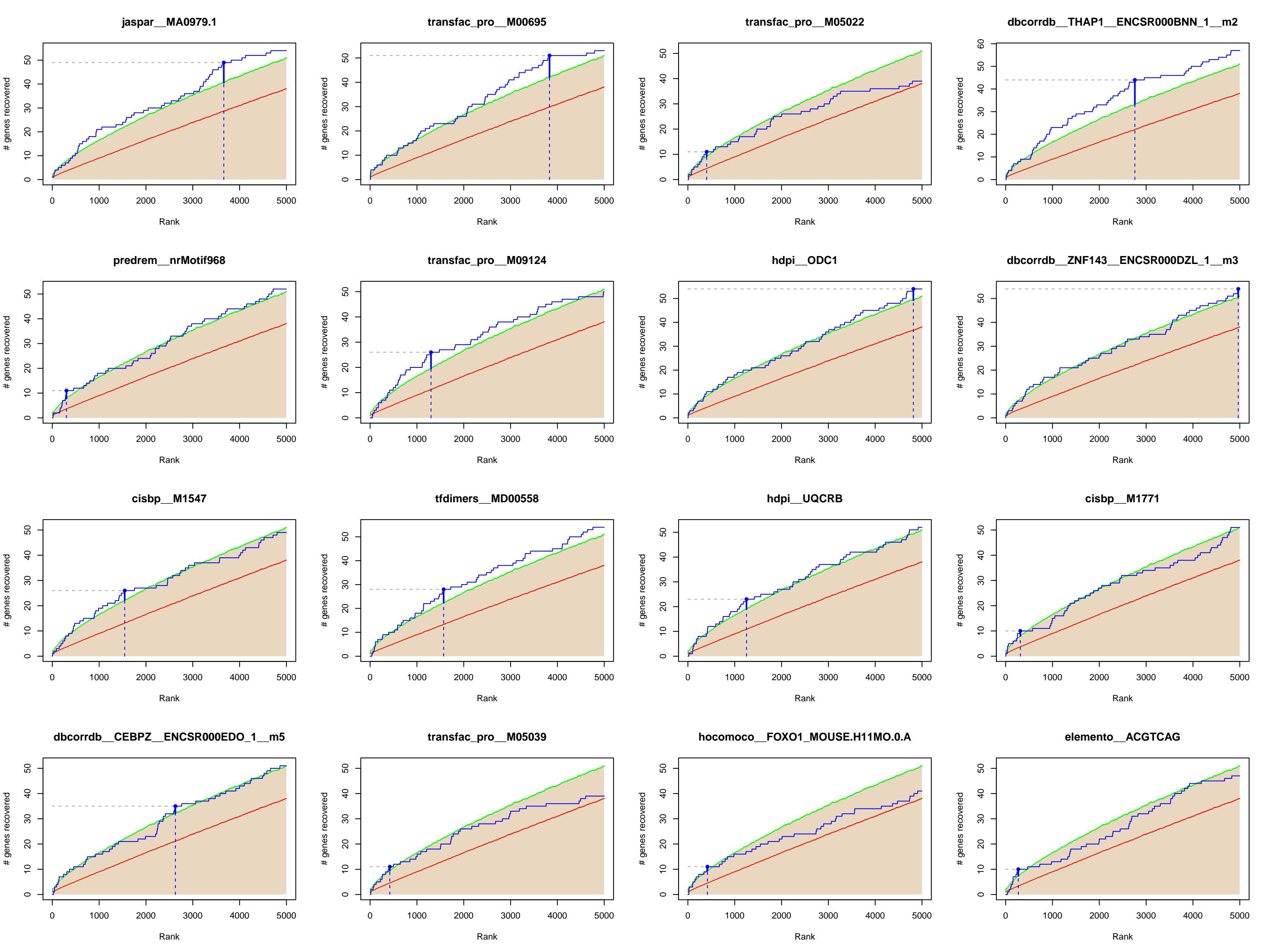
transfac_pro_M07942



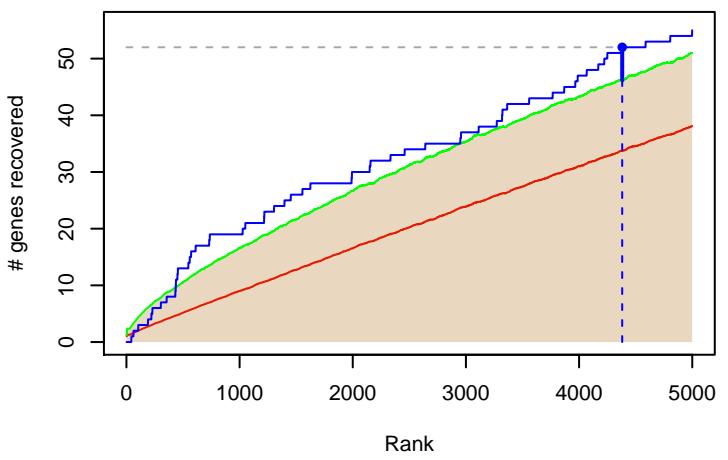




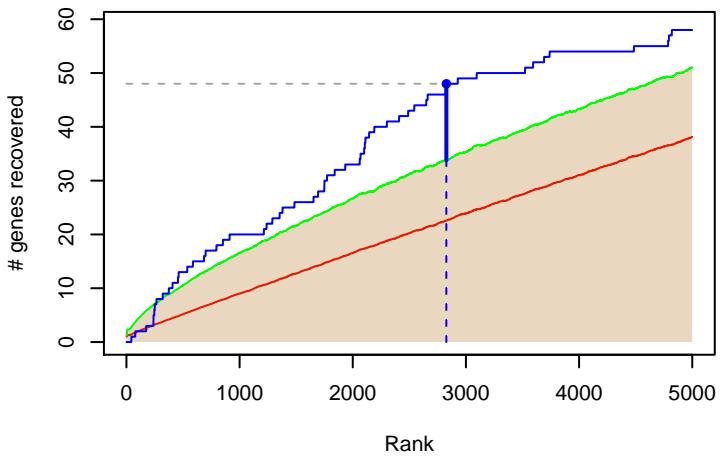




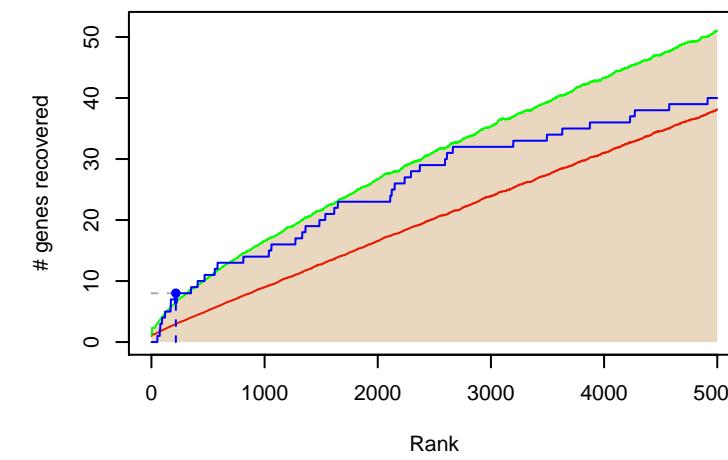
elemento_TCGAACTC



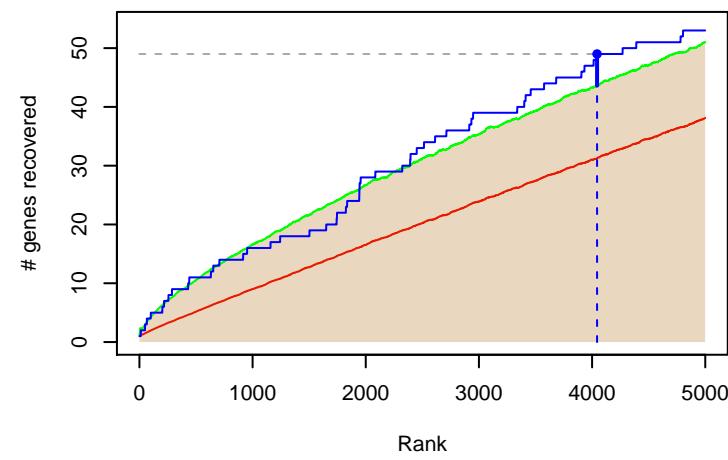
dbcordb_POLR2AphosphoS5_ENCSR000BML_1_m1



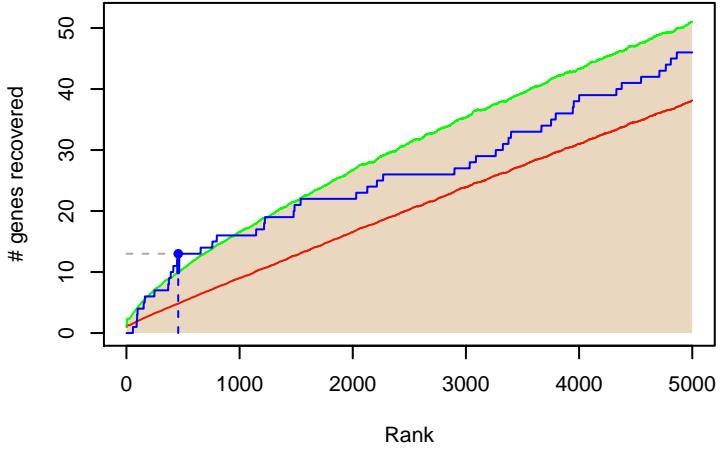
dbcordb_SAP30_ENCSR000AQJ_1_m1



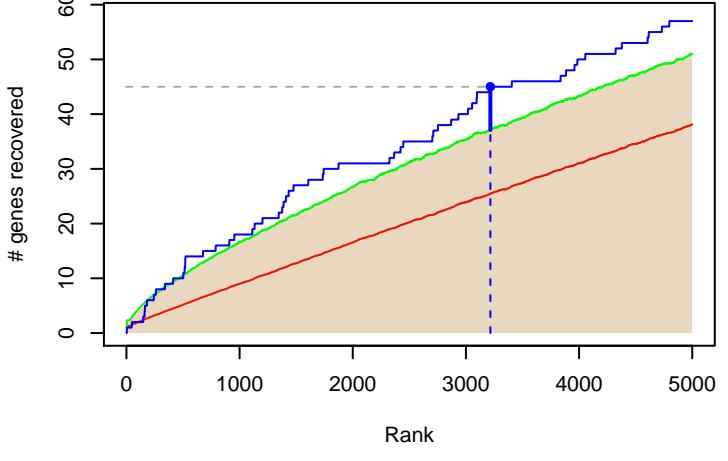
cisbp_M0609



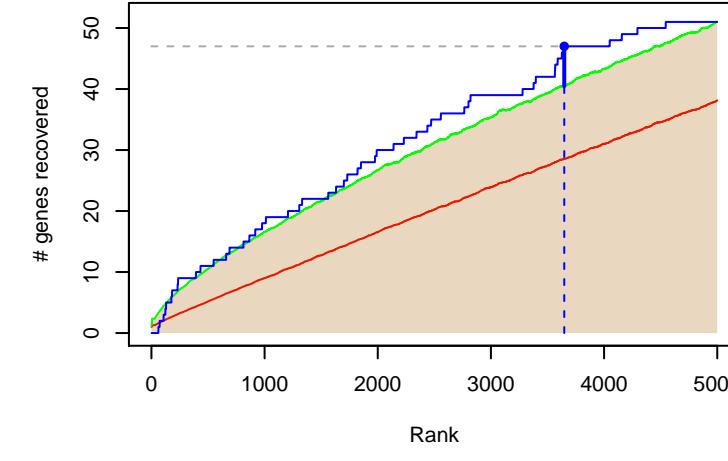
dbcordb_ATF1_ENCSR000DNZ_1_m2



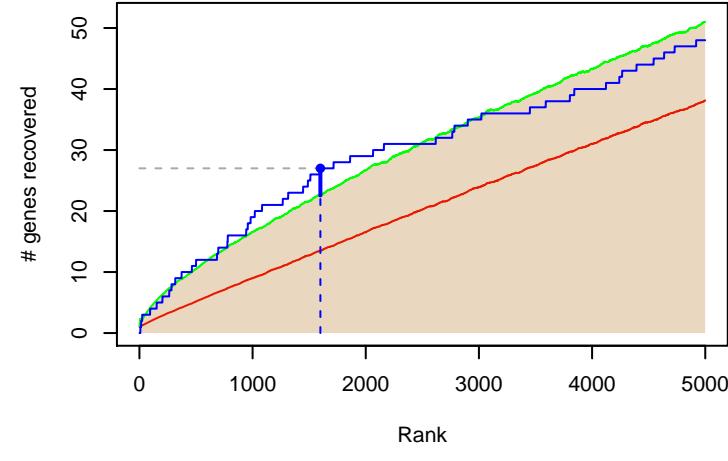
dbcordb_GTF2F1_ENCSR000ECZ_1_m1



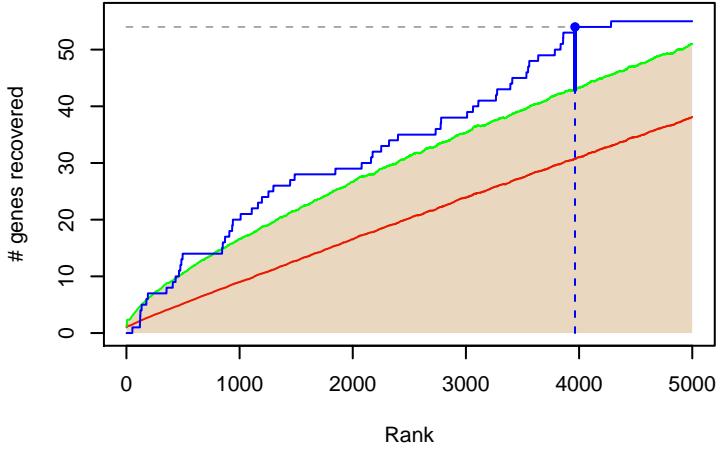
flyfactorsurvey_rn_SOLEXA_F2-4



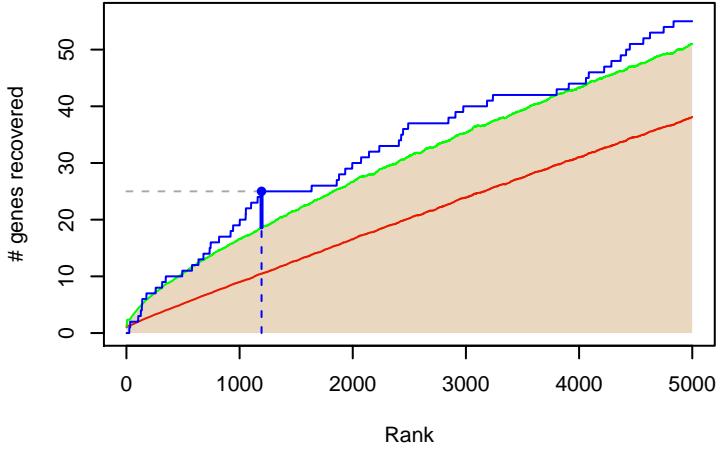
pale_tf_pairs_E2F3_FOXO6_NMATGACACGCCMNN_CAG



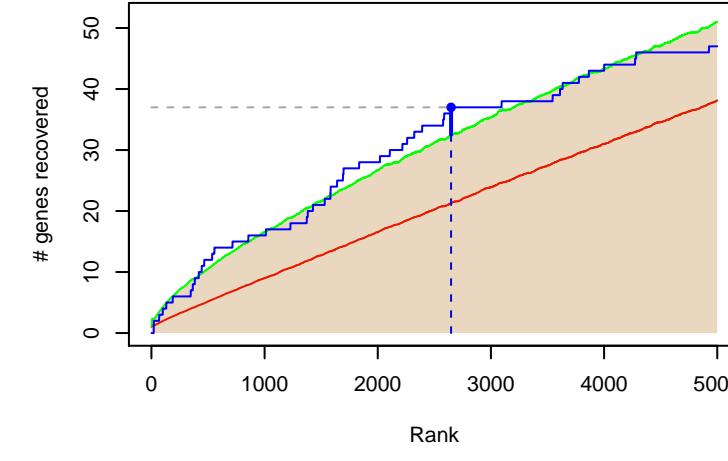
dbcordb_POLR2A_ENCSR000EAM_1_m1



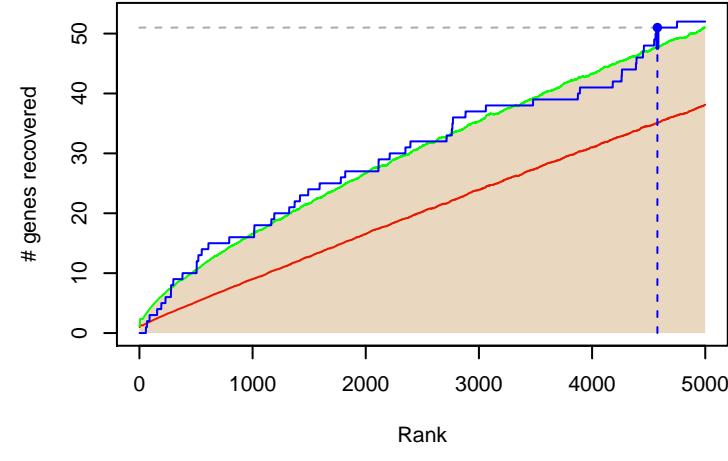
dbcordb_EZH2_ENCSR000ASW_1_m4



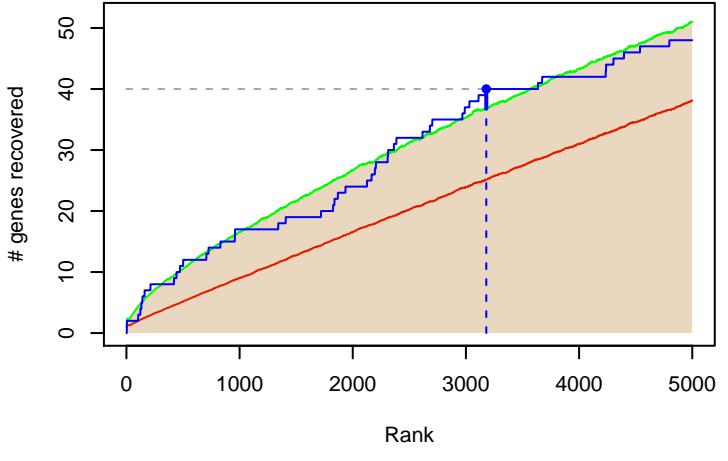
predrem_nrMotif50



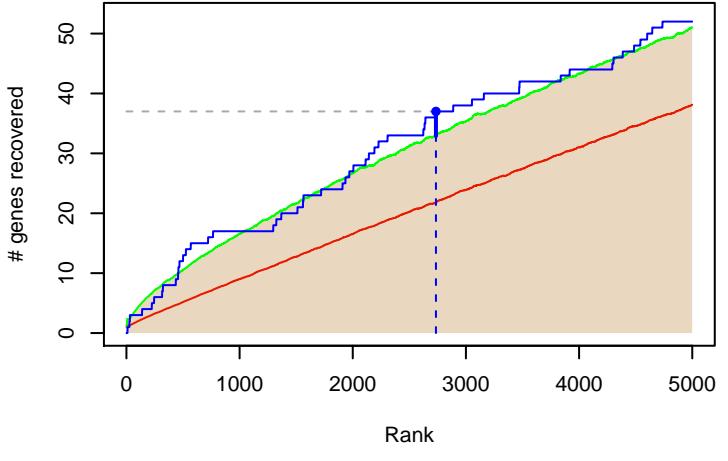
dbcordb_IRF3_ENCSR000DZX_1_m4



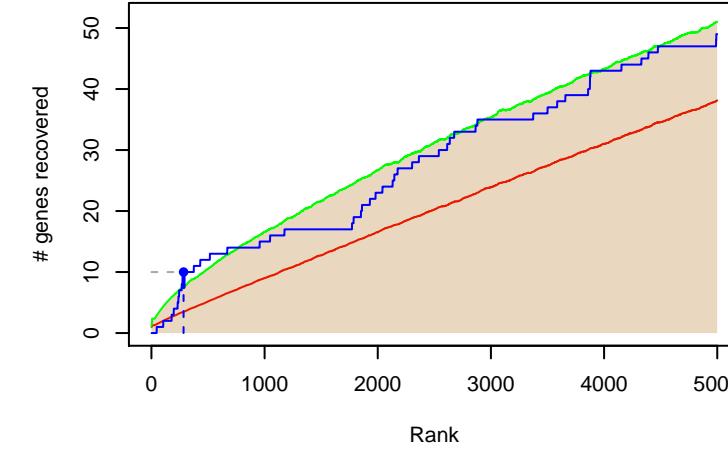
dbcordb_RBBP5_ENCSR000AQC_1_m2



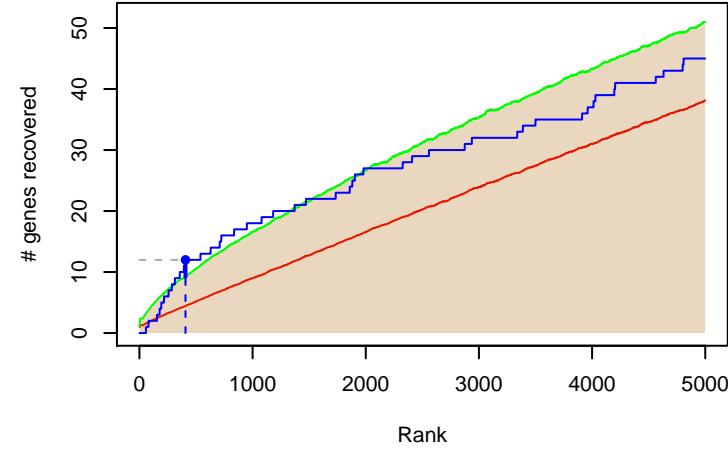
cisbp_M0017



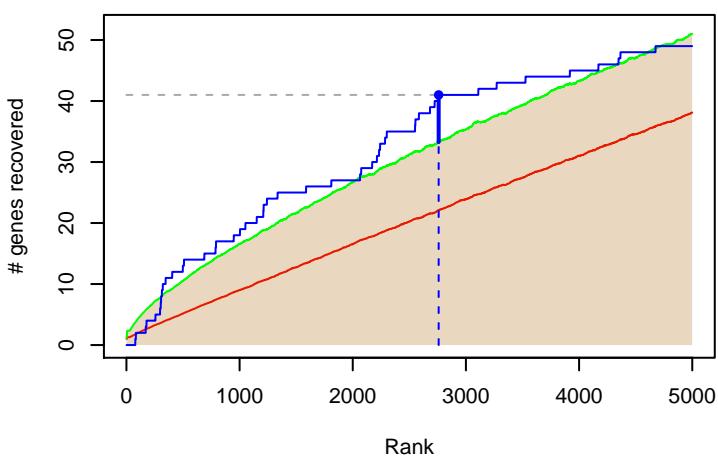
cisbp_M5020



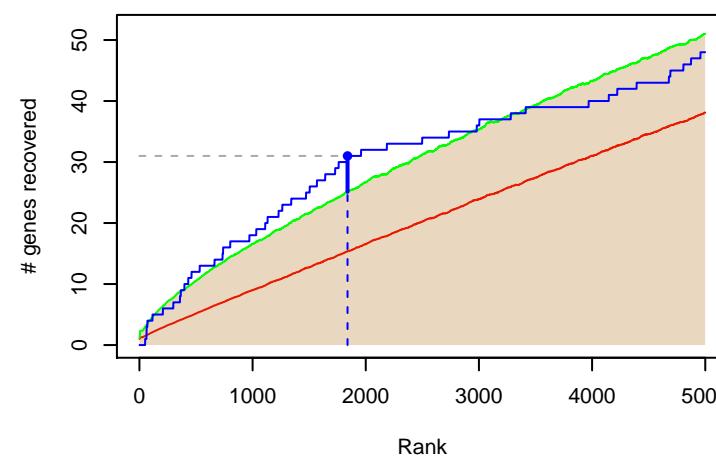
elemento_AGGCCGC



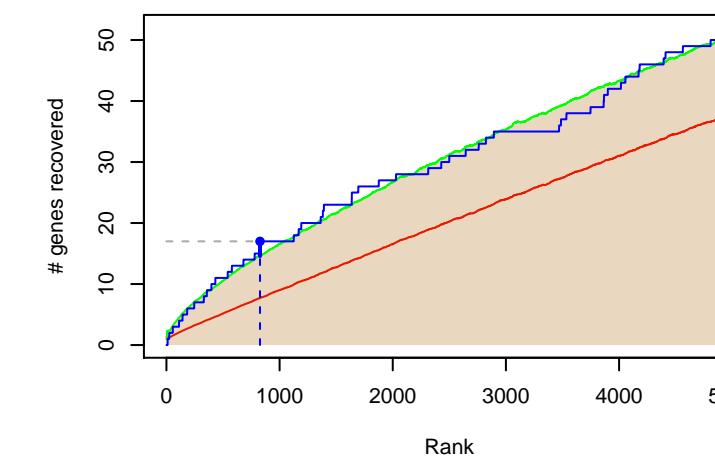
dbcrrdb__POLR2A__ENCSR000EAY_1_m1



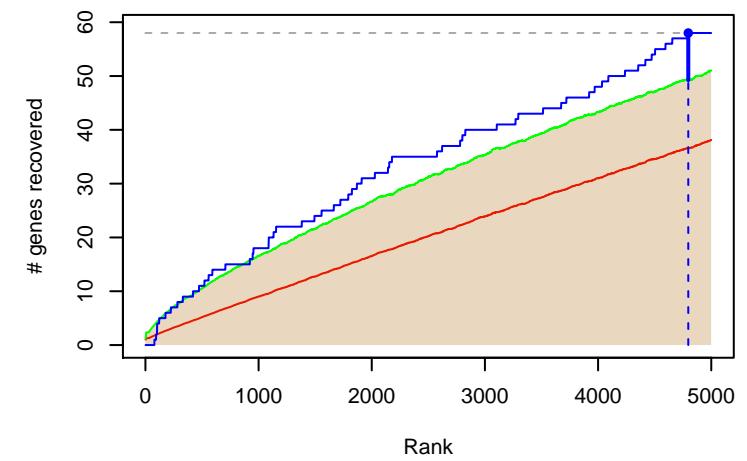
flyfactorsurvey__jigr1_SANGER_5_FBgn0039350



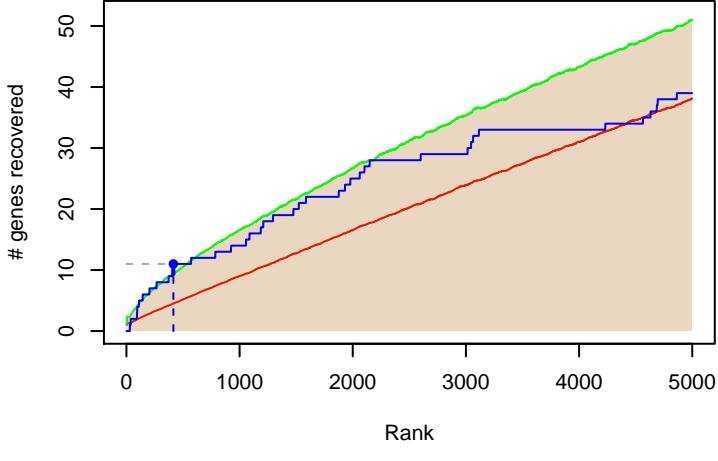
cisbp__M1543



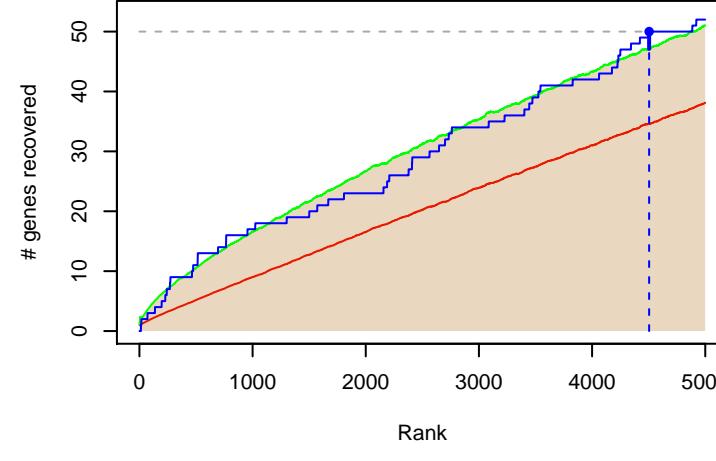
elemento__CCGCCCCG



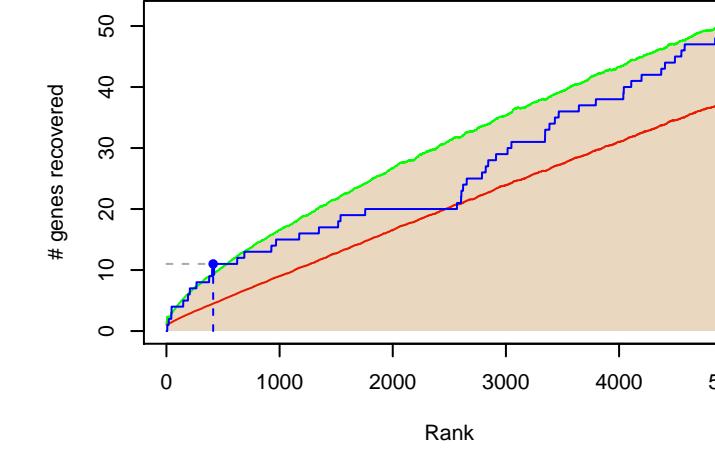
elemento__ATGGCGGC



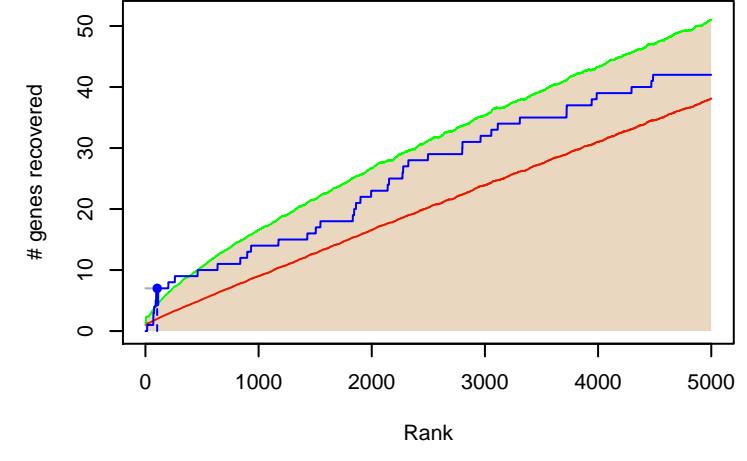
cisbp__M1462



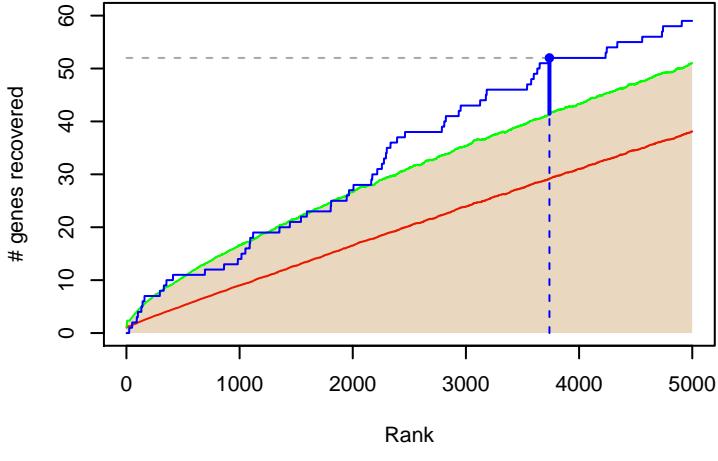
transfac_pro__M02114



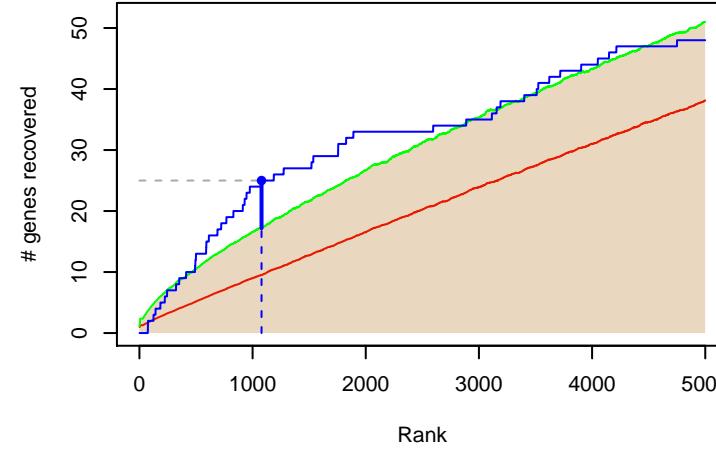
scertf__macisaac.AZF1



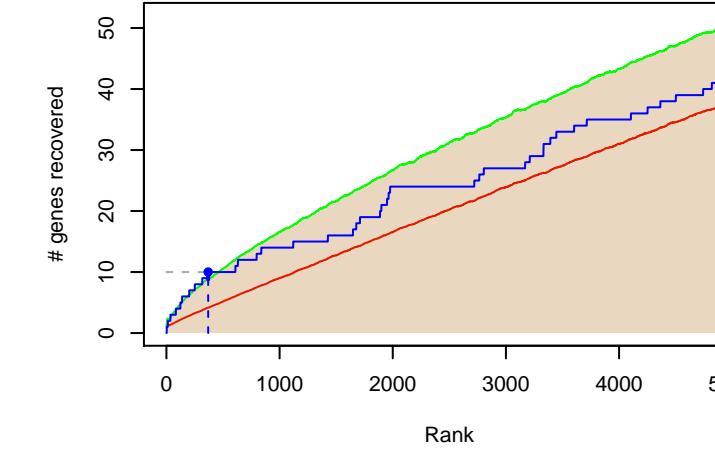
cisbp__M0608



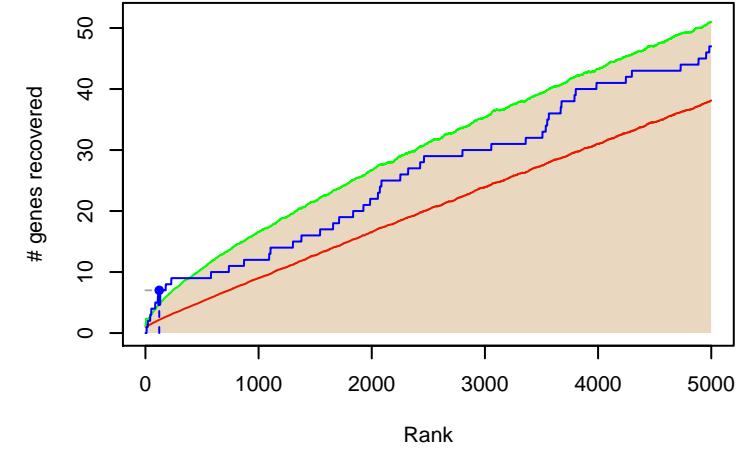
homer__WTTTTCYYTTTT_RLR1_



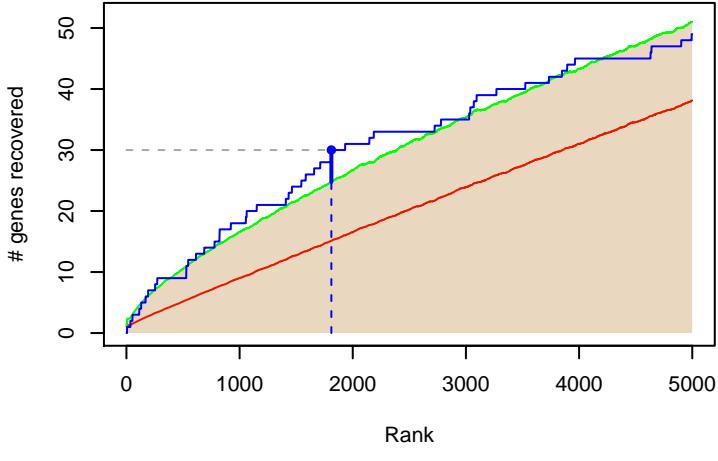
idmmpmm__Mad



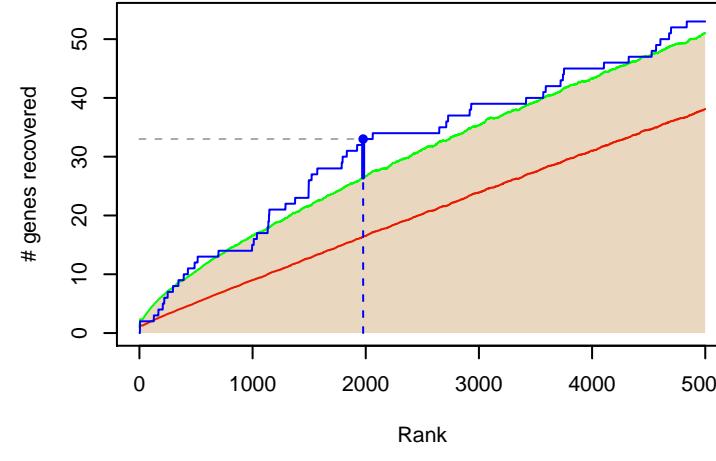
swissregulon__hs__CREB1.p2



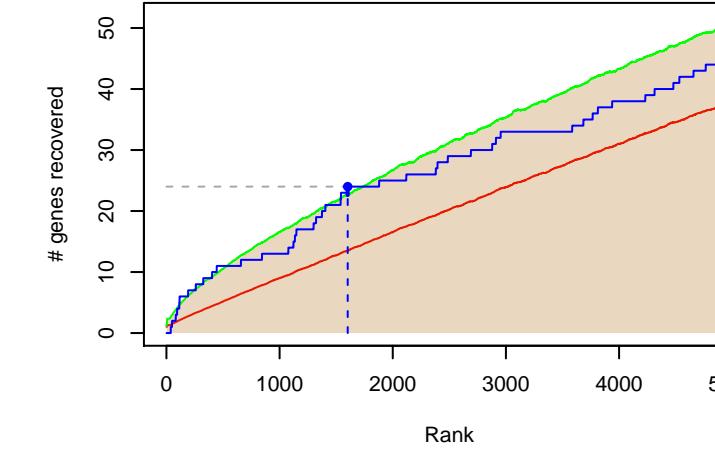
cisbp__M1553



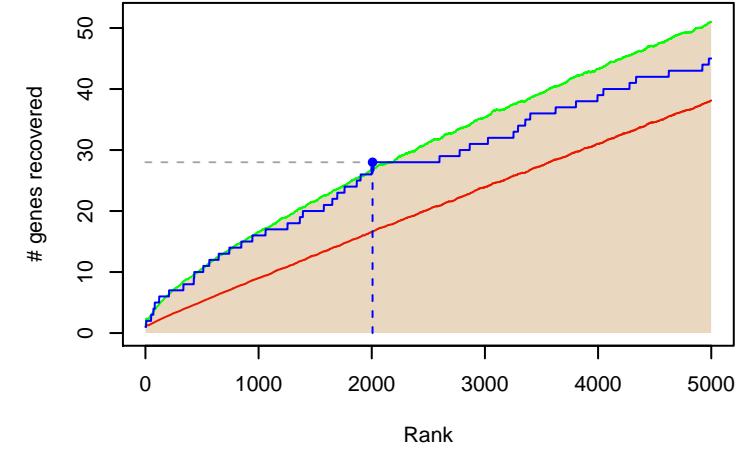
transfac_pro__M04916

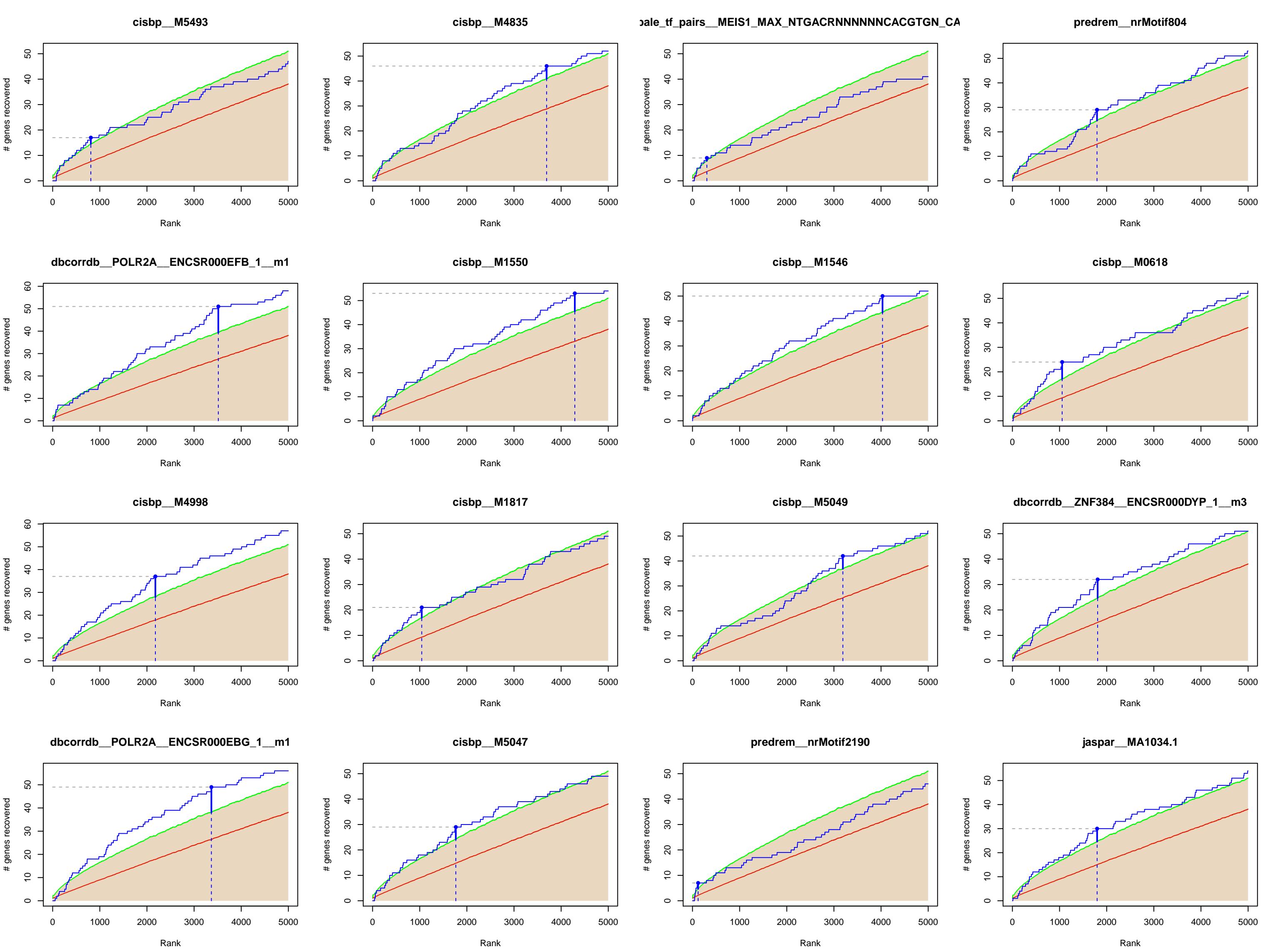


transfac_pro__M09283

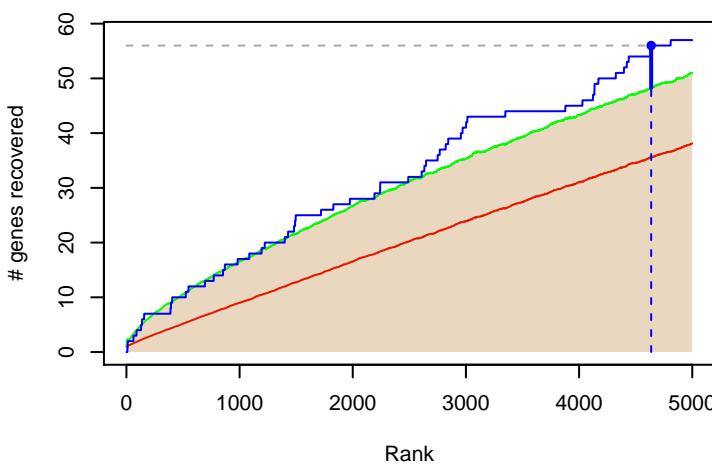


dbcrrdb__ZNF384__ENCSR000EFP_1_m3

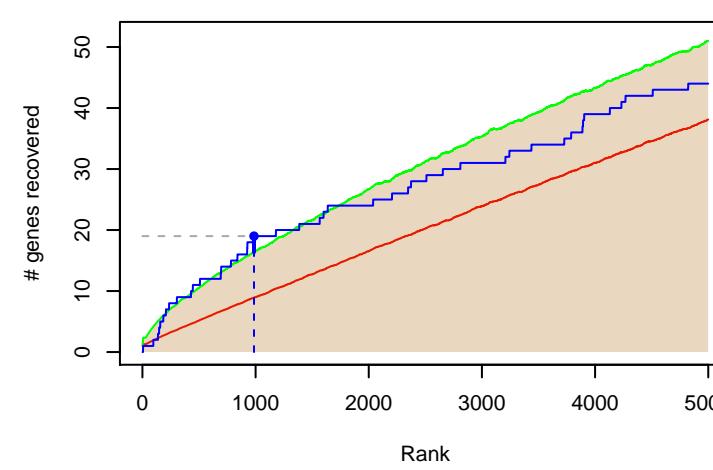




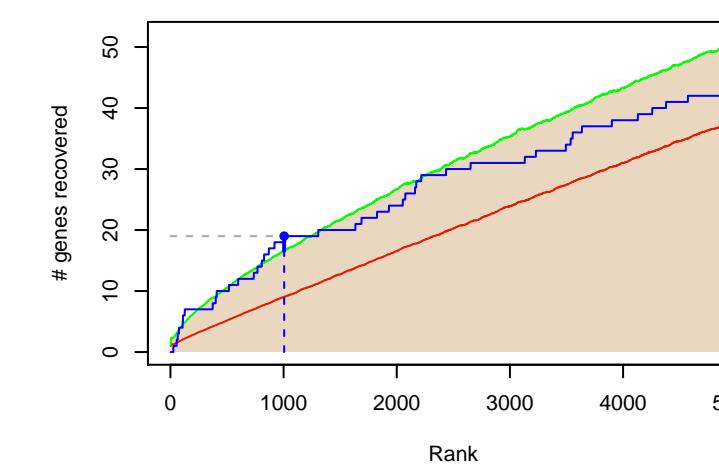
transfac_pro_M07533



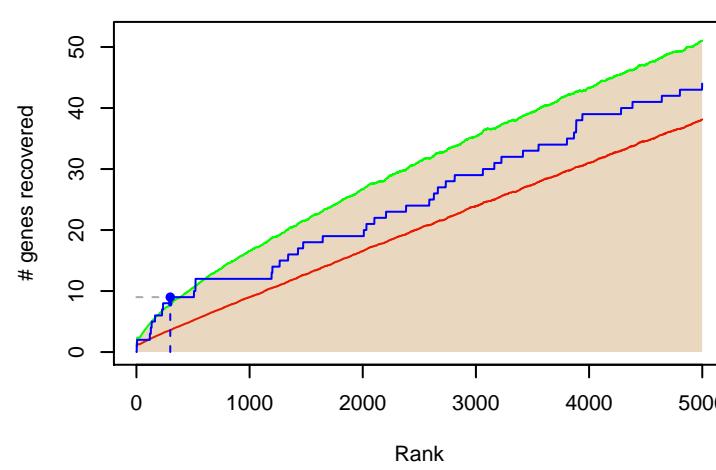
transfac_pro_M02109



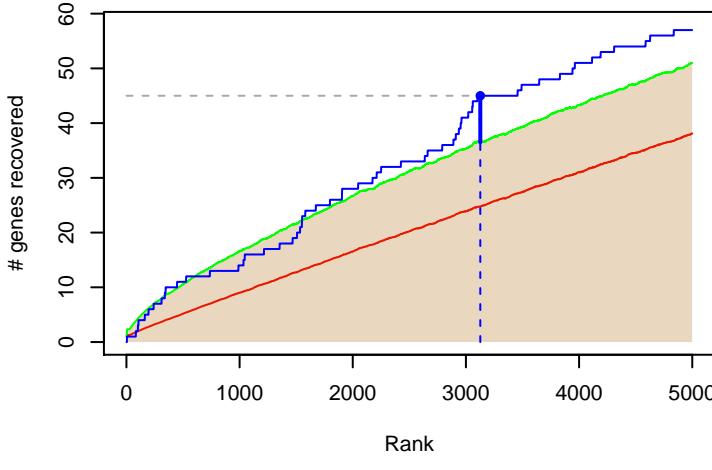
yetfasco_YOR032C_1498



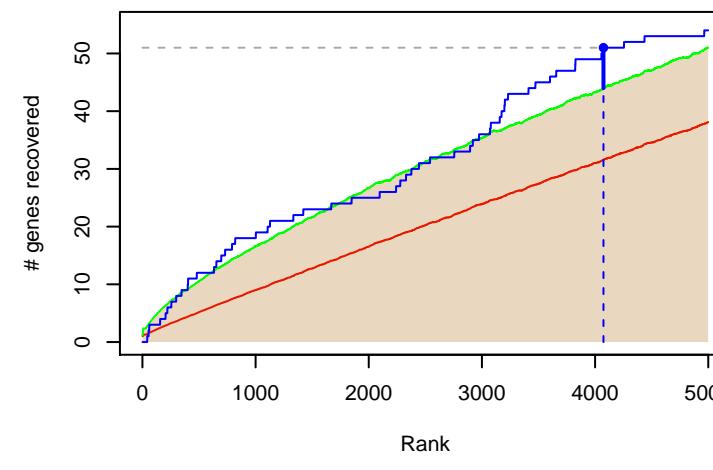
jaspar_MA0604.1



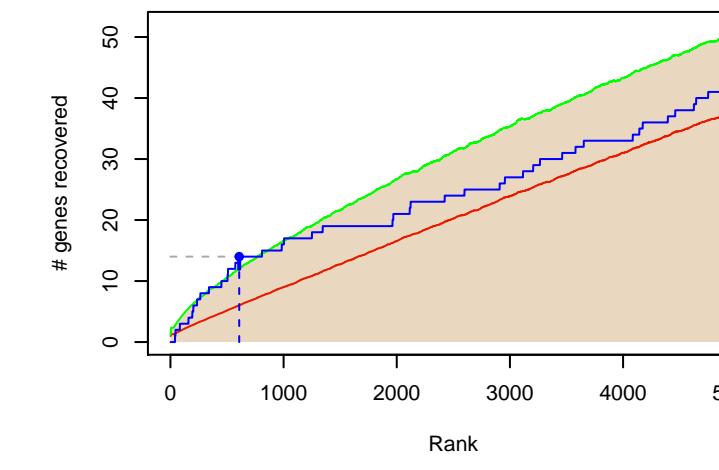
dbcorrdb_NR3C1_ENCSR000EEV_1_m1



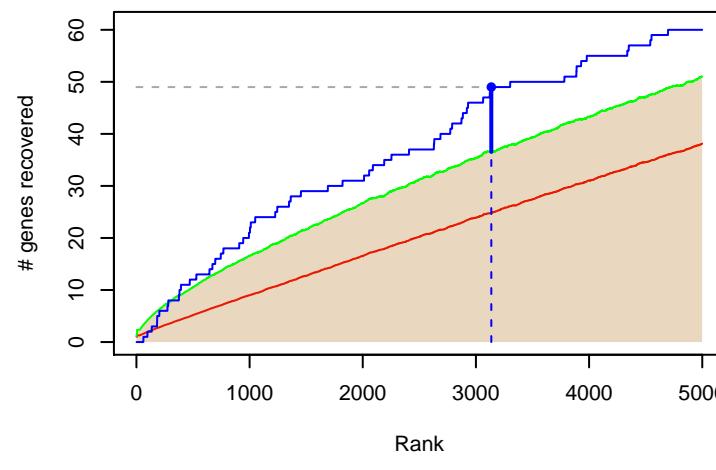
tfdimers_MD00285



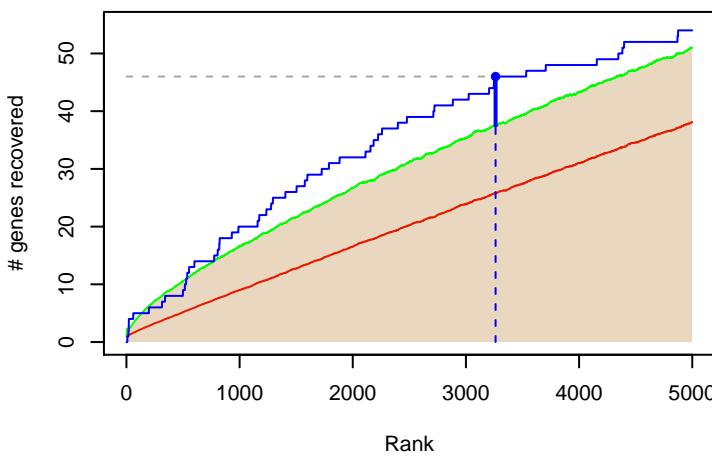
fantom_motif110_AGTCGCAGCG



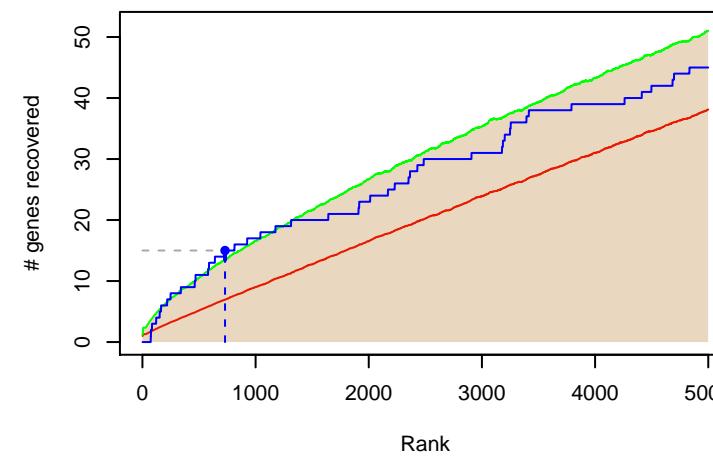
dbcorrdb_POLR2A_ENCSR000FAL_1_m1



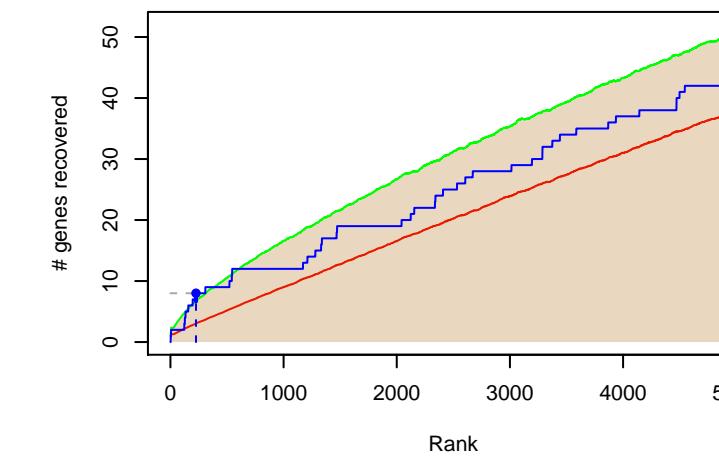
scertf_harbison.SUT1



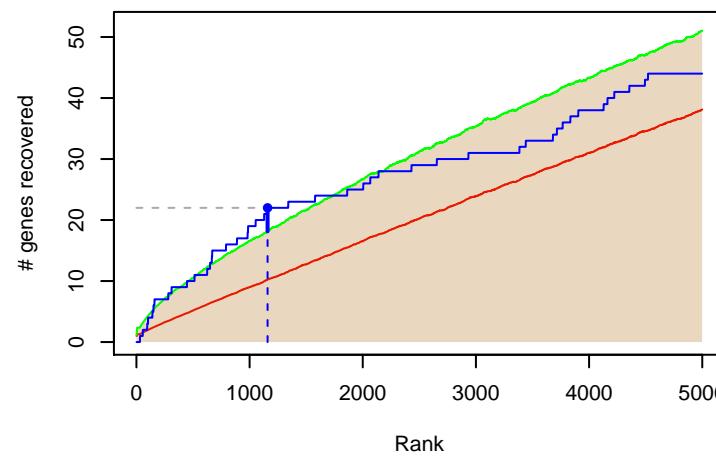
taipale_GMEB2_DBD_TTACGYAM



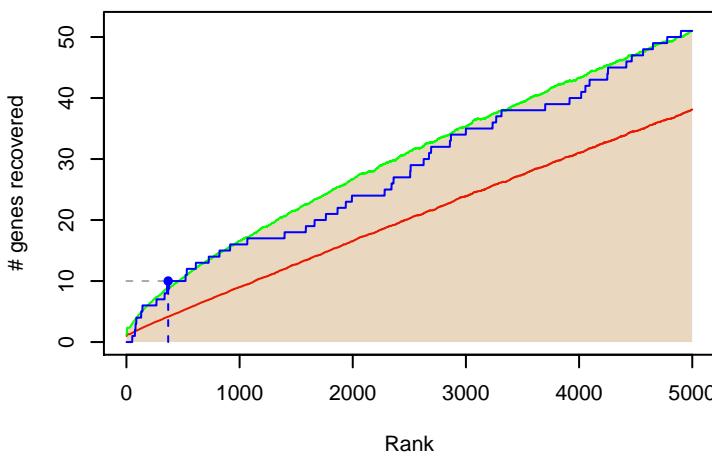
cisbp_M0295



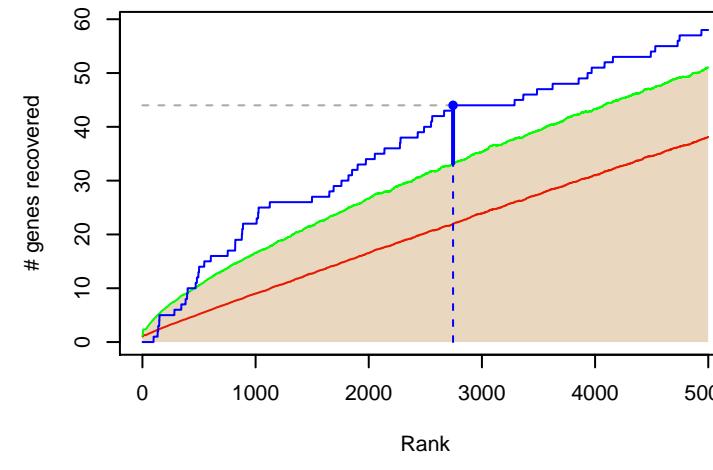
transfac_pro_M09131



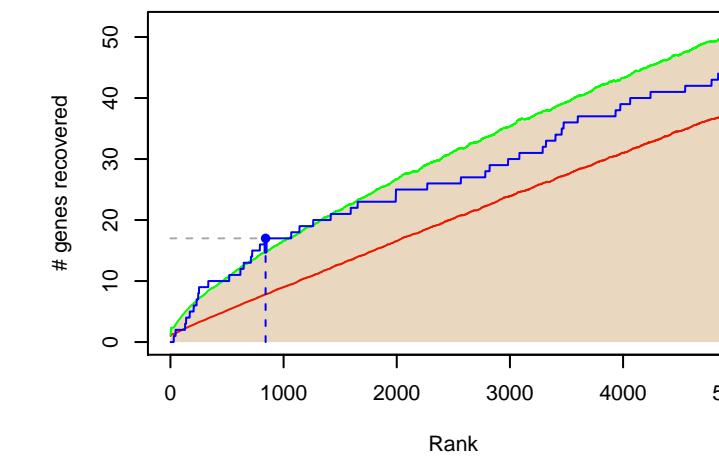
transfac_pro_M00987



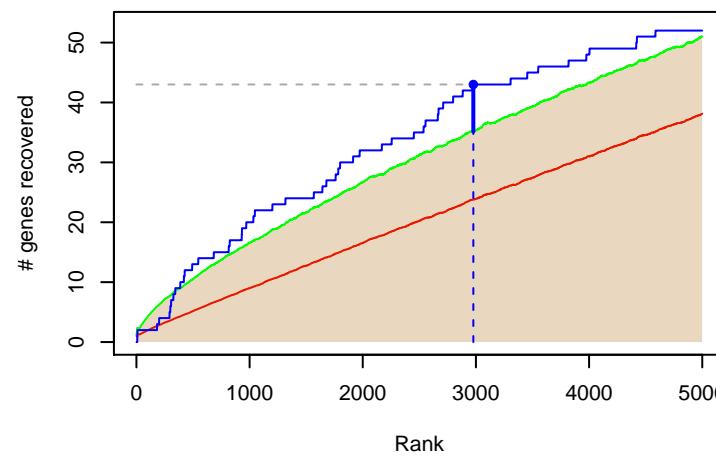
dbcorrdb_POLR2AphosphoS5_ENCSR000BIA_1_m1



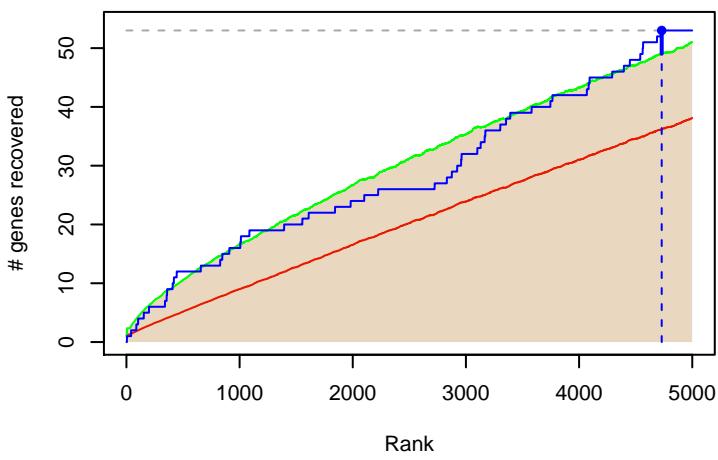
transfac_pro_M04744



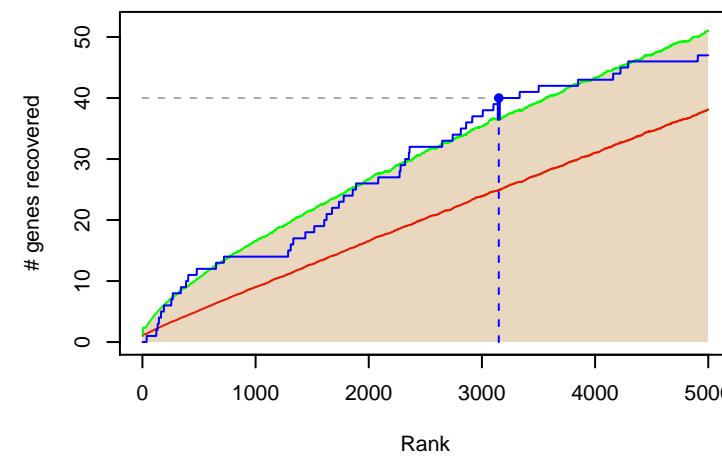
cisbp_M1545



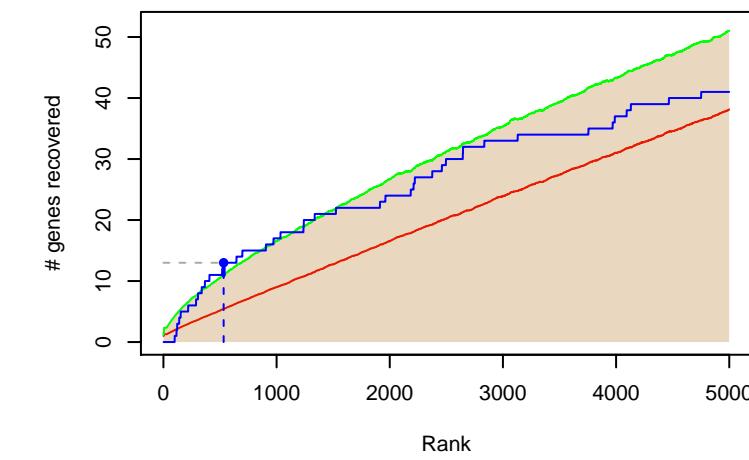
dbcrrdb_KDM5A_ENCSR000AQL_1_m1



flyfactorsurvey(CG2052_SANGER_2.5_FBgn0039905)



hocomoco_MLX_HUMAN.H11MO.0.D



predrem_nrMotif583

