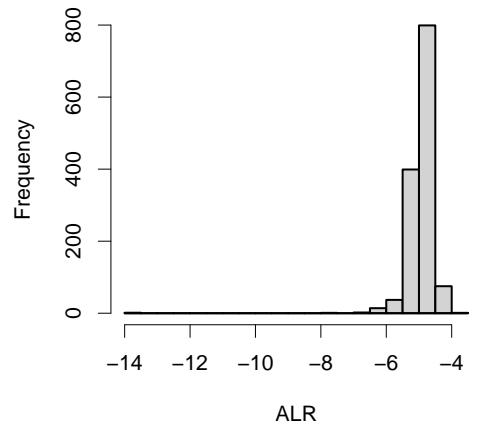
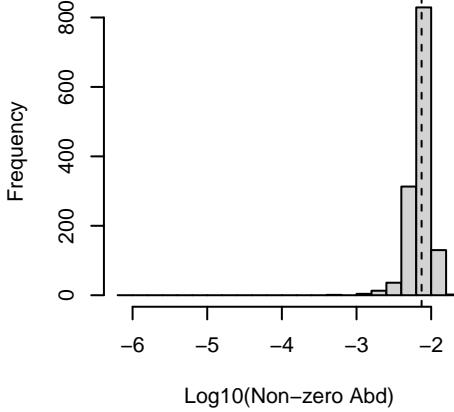
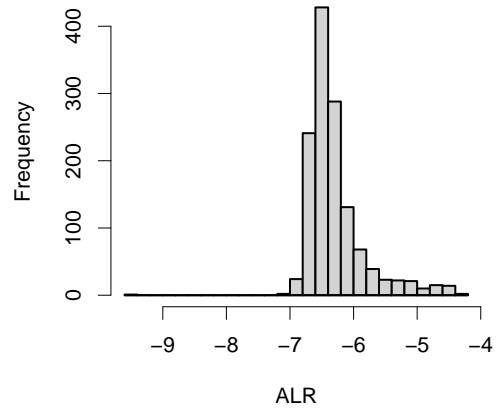
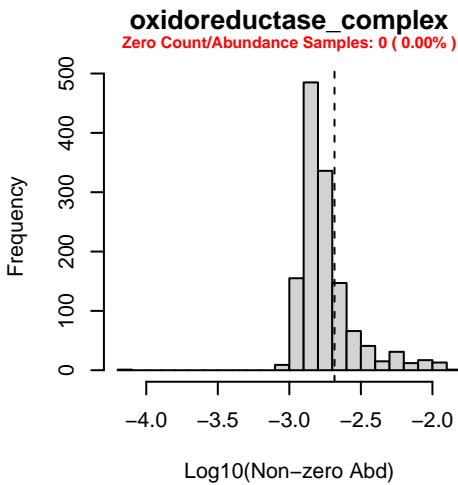
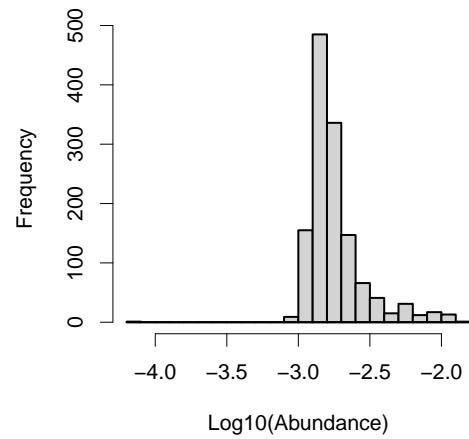
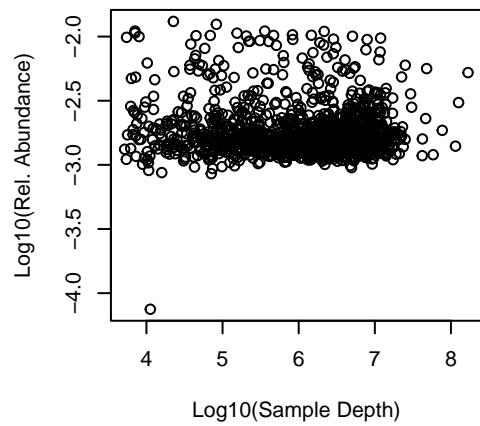
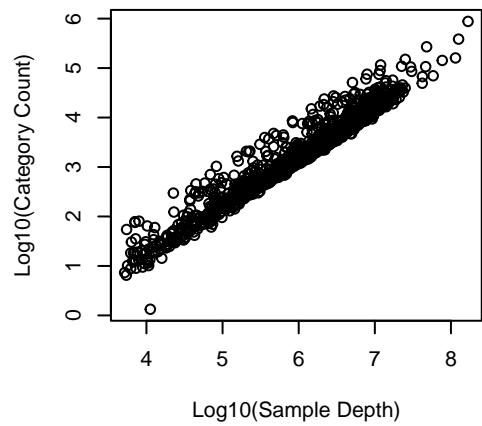
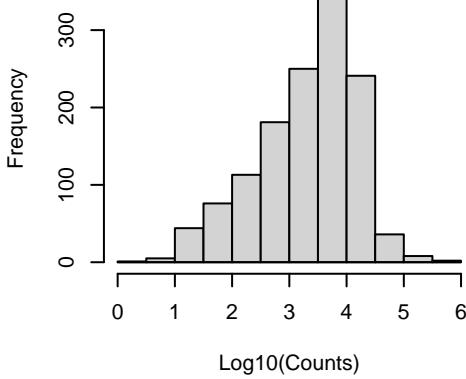
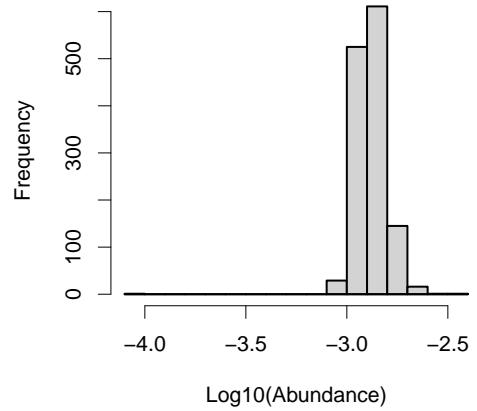
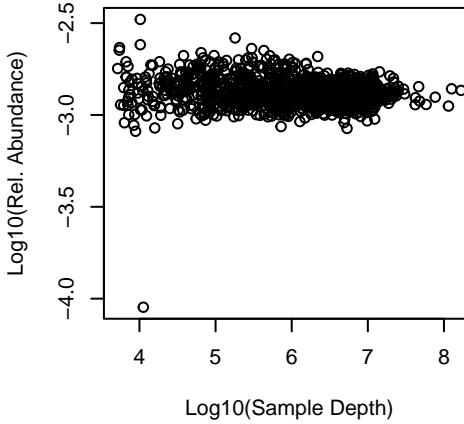
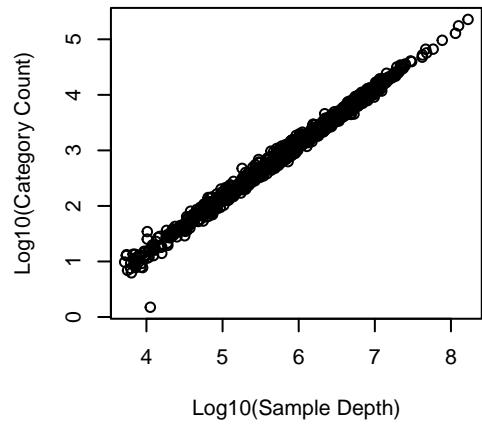
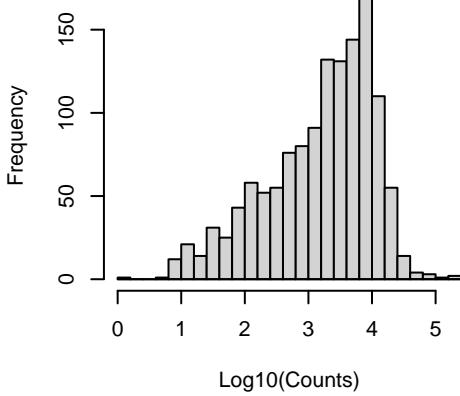


beta_galactosidase_complex

Zero Count/Abundance Samples: 1 (0.08%)

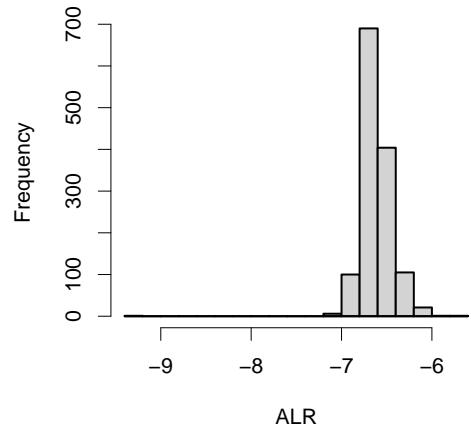
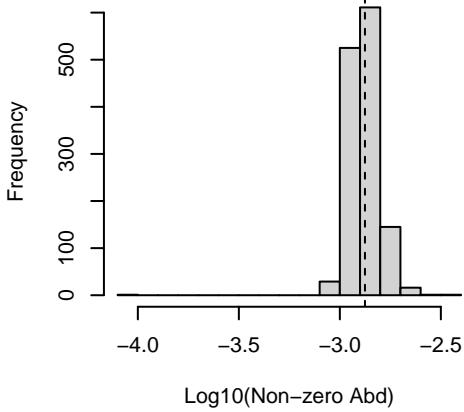


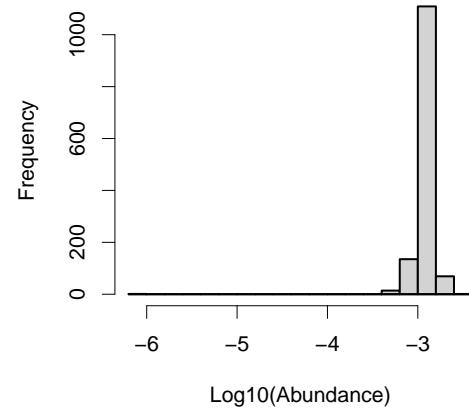
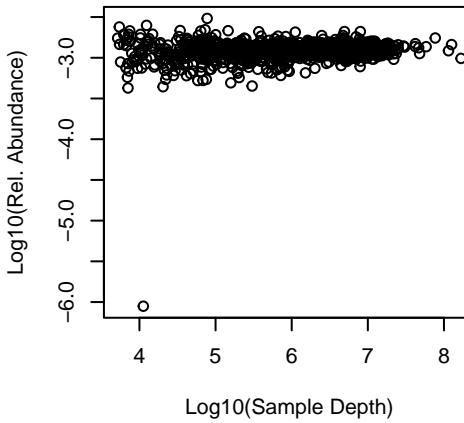
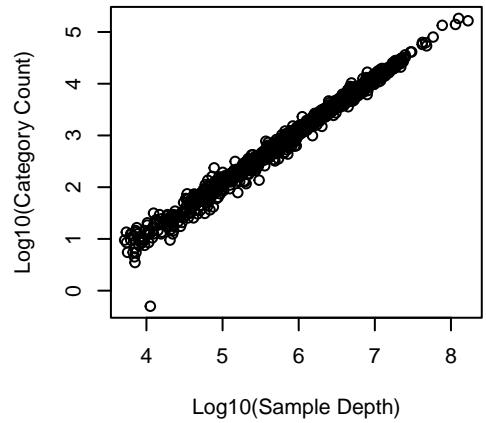
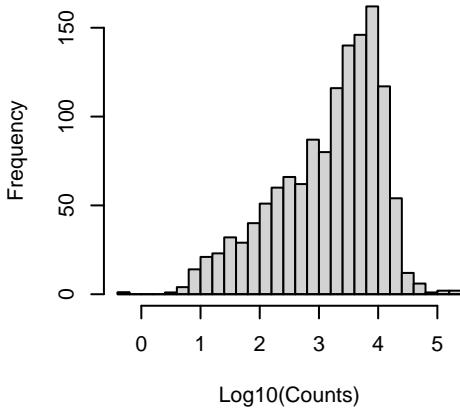




endodeoxyribonuclease complex

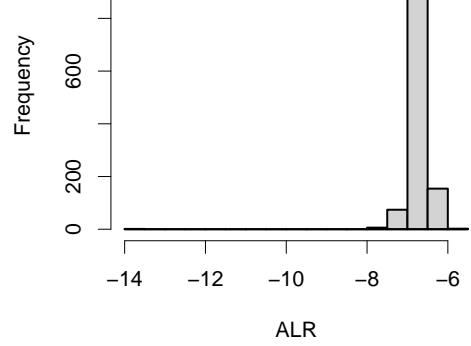
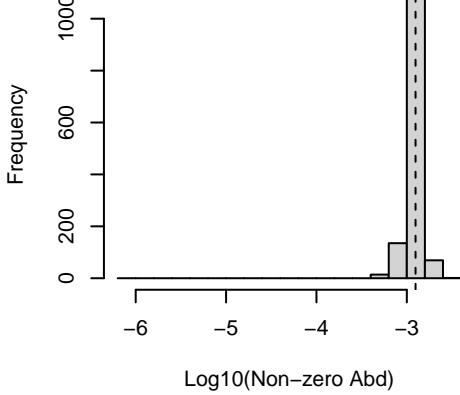
Zero Count/Abundance Samples: 0 (0.00%)

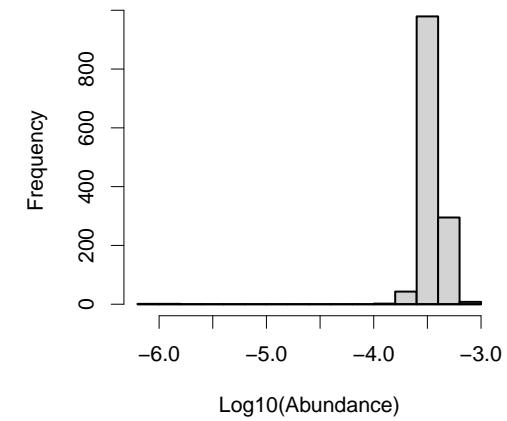
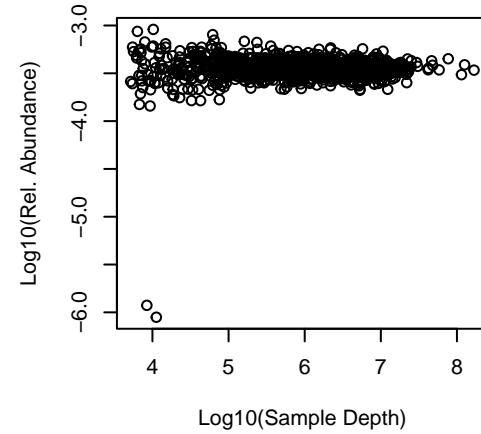
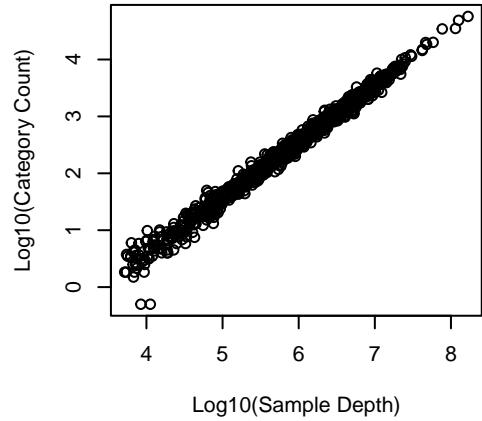
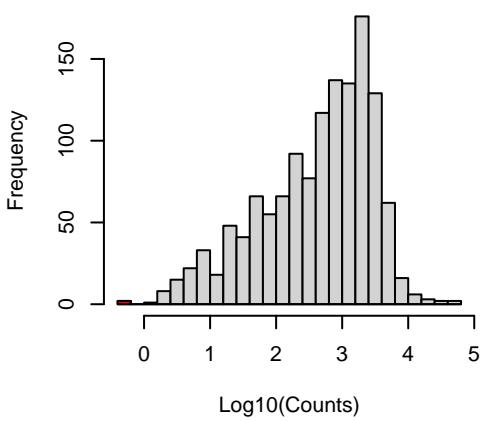




acetyl_CoA_carboxylase_complex

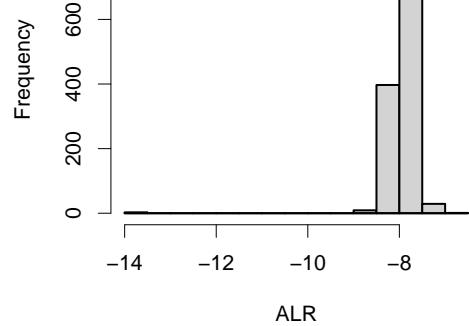
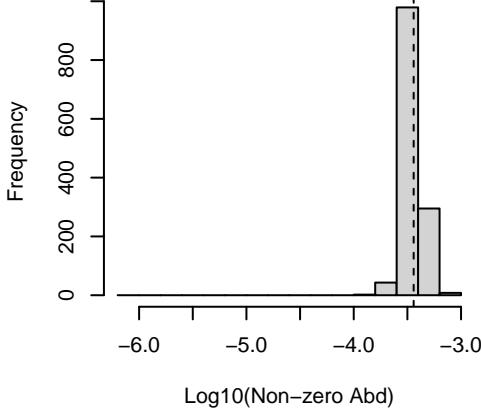
Zero Count/Abundance Samples: 1 (0.08%)

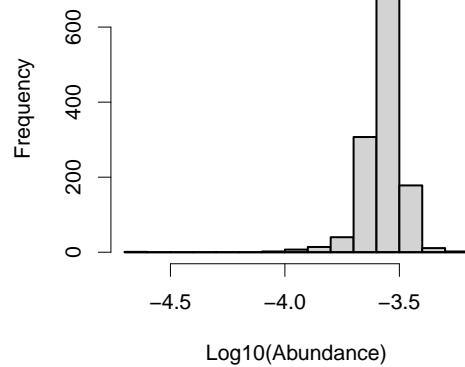
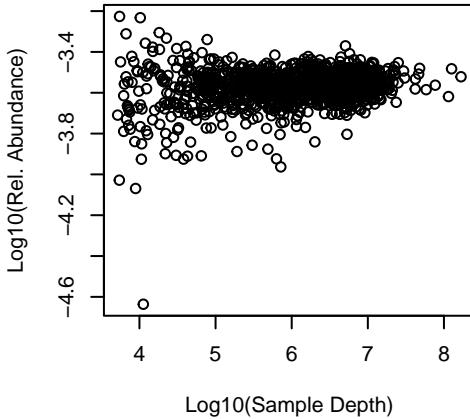
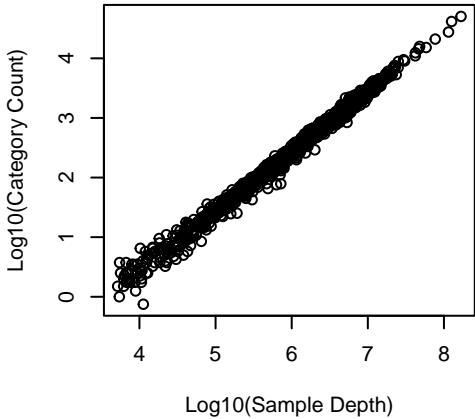
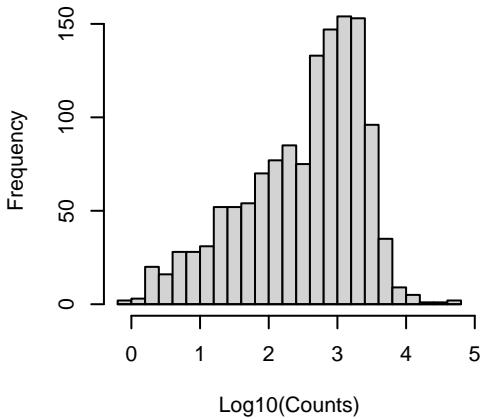




phosphopyruvate_hydratase_complex

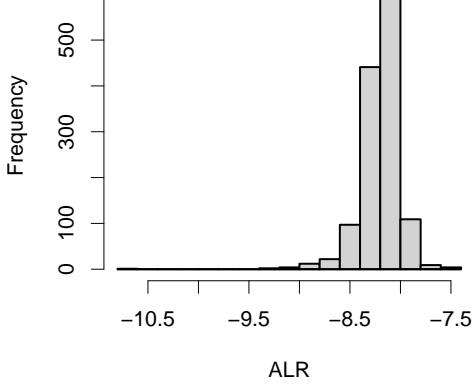
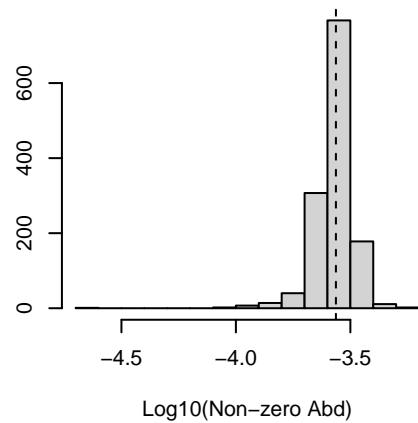
Zero Count/Abundance Samples: 2 (0.15%)

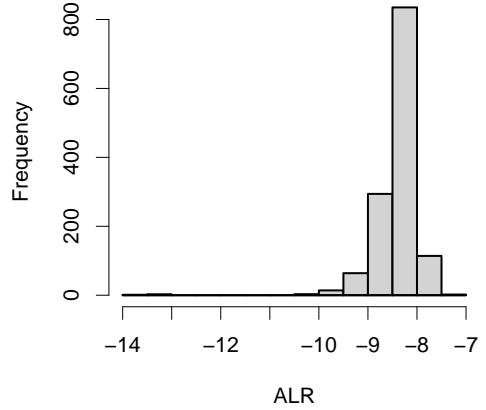
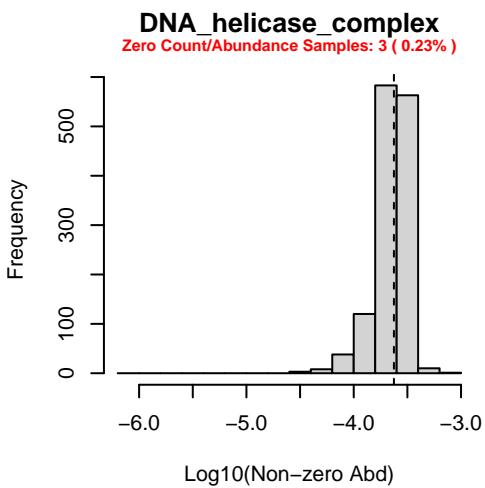
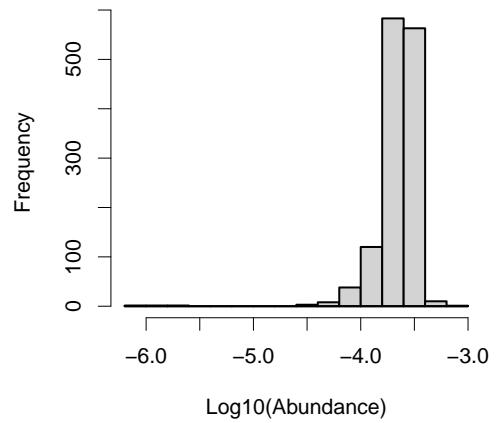
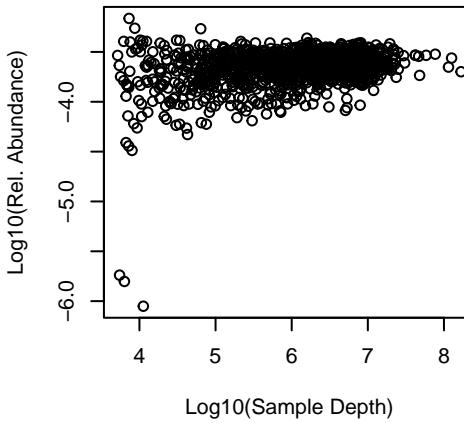
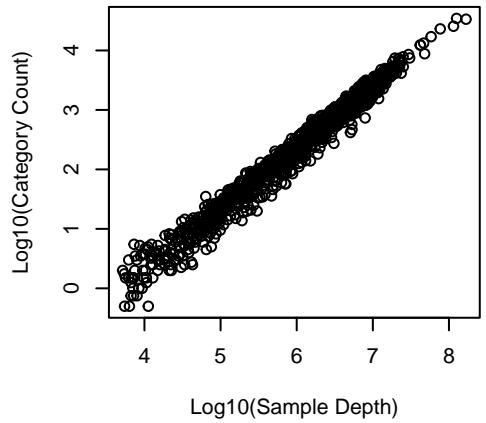
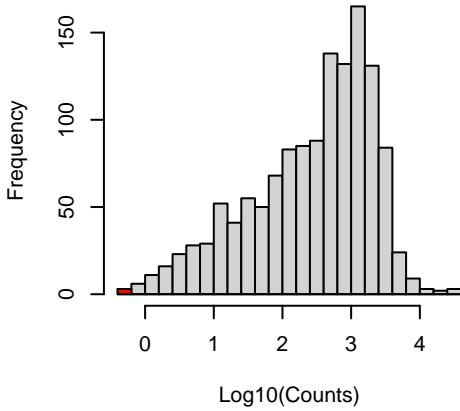


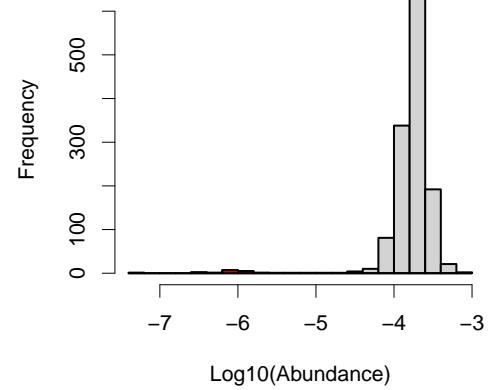
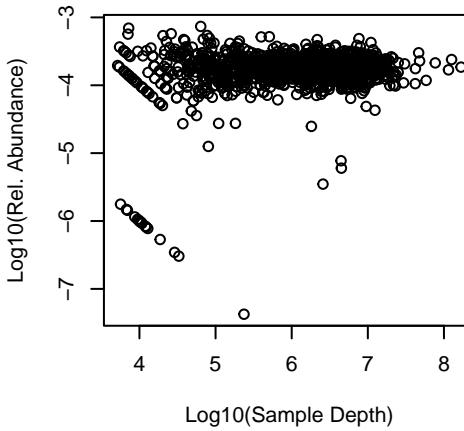
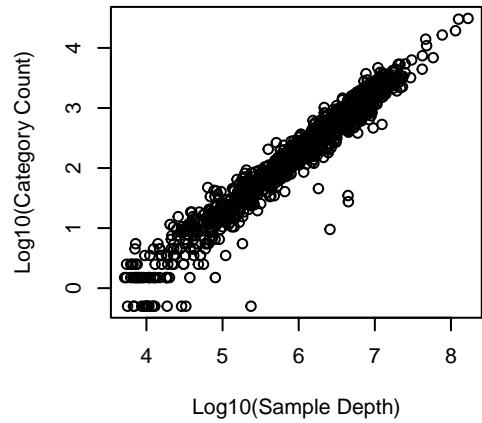
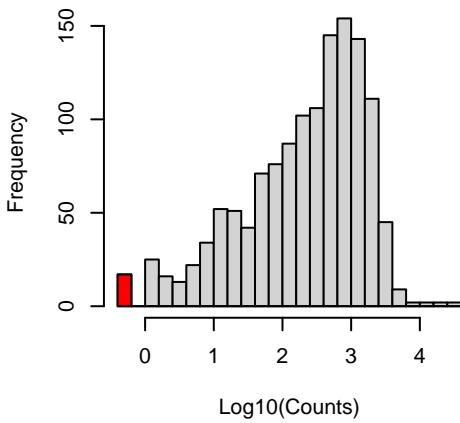


exodeoxyribonuclease_VII_complex

Zero Count/Abundance Samples: 0 (0.00%)

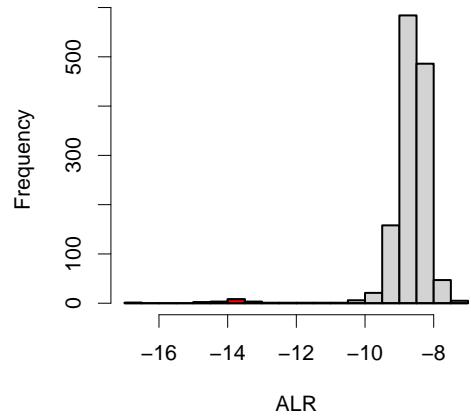
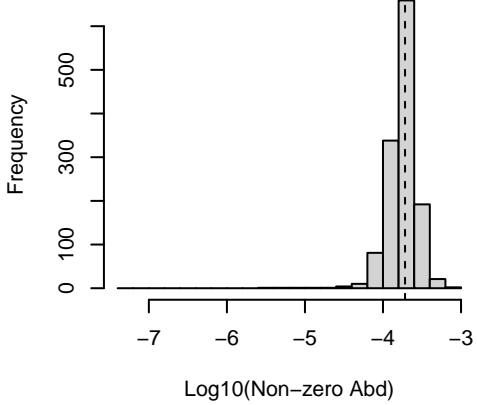


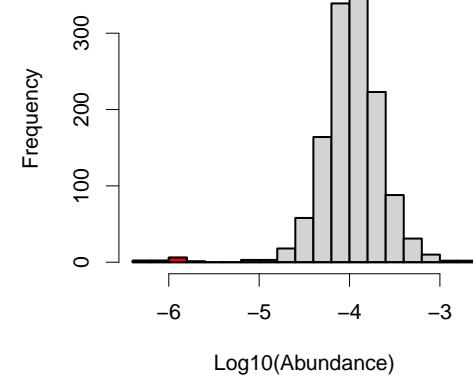
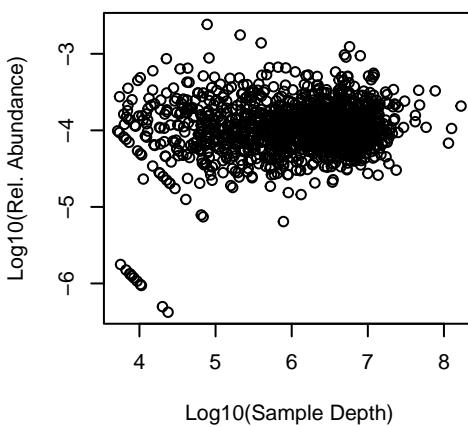
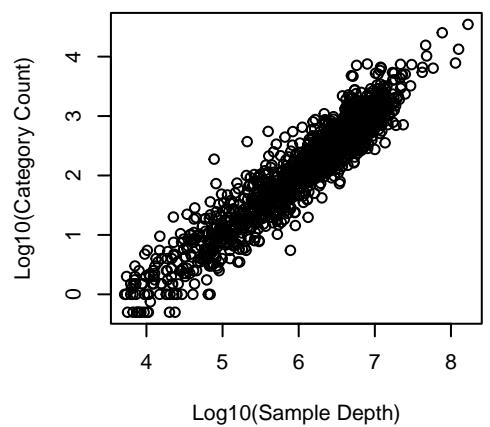
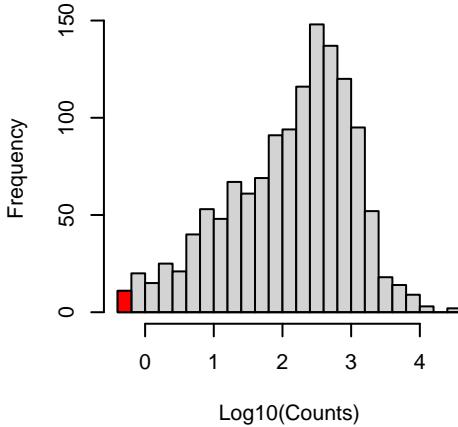




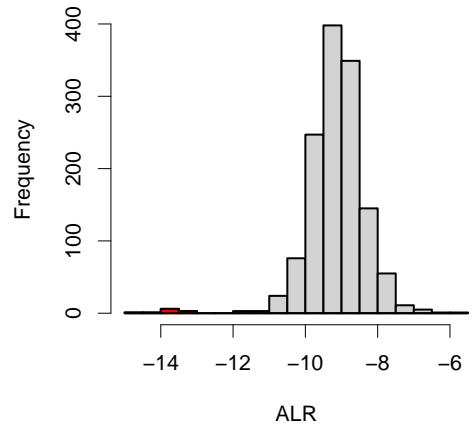
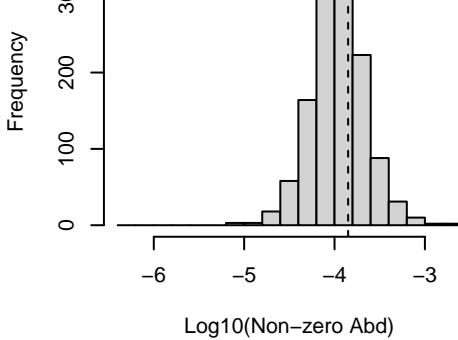
X3_isopropylmalate_dehydratase_comp

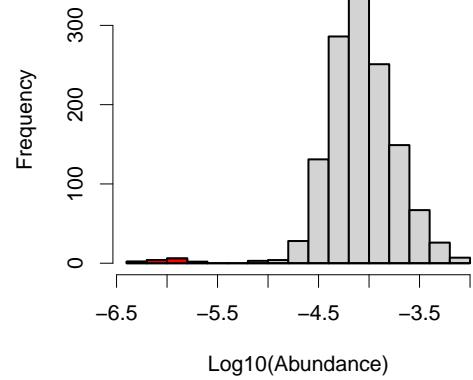
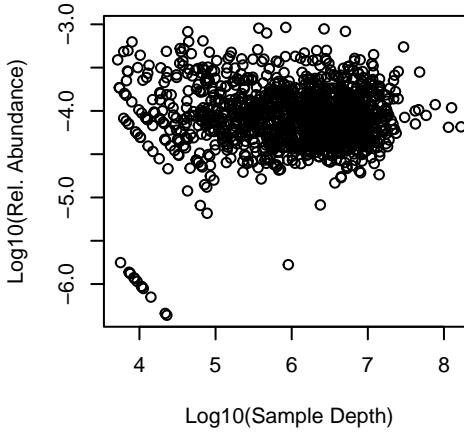
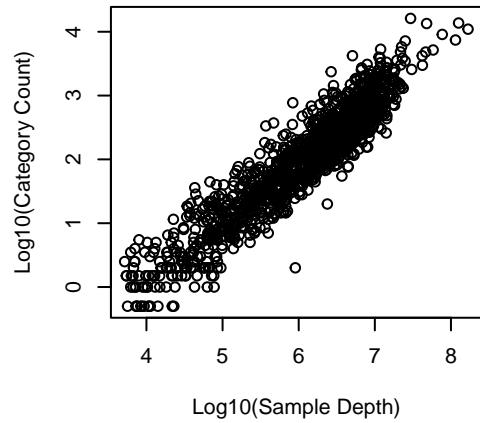
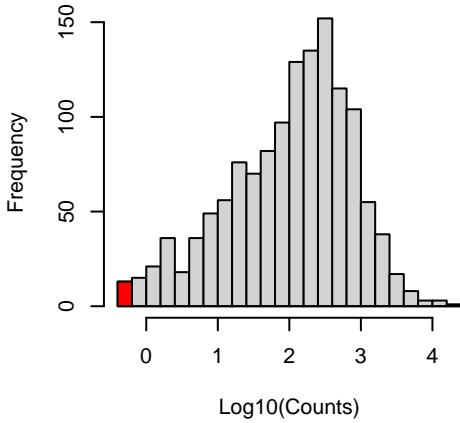
Zero Count/Abundance Samples: 17 (1.28%)





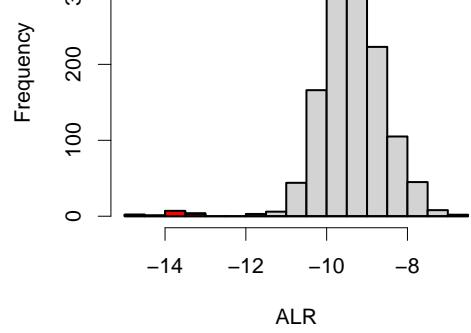
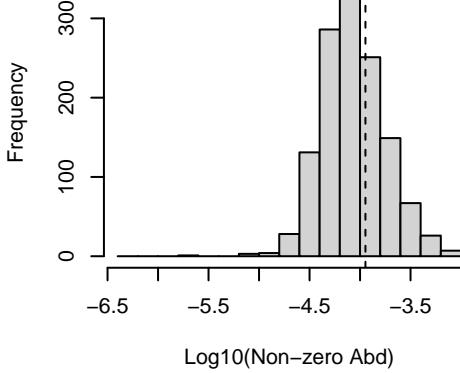
peptidase_complex
Zero Count/Abundance Samples: 11 (0.83%)

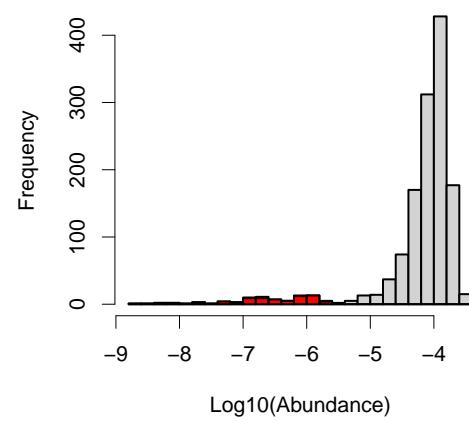
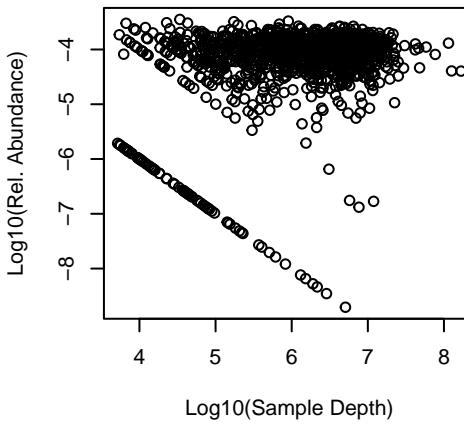
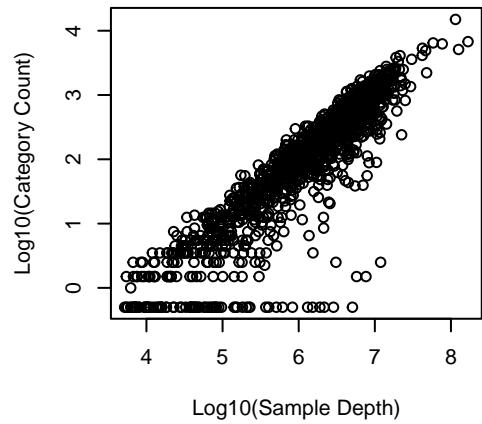
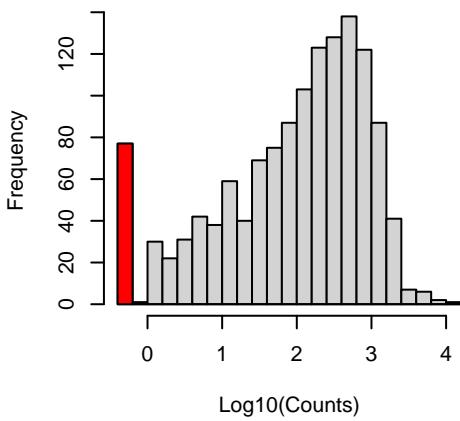




ATP_independent_citrate_lyase_complex

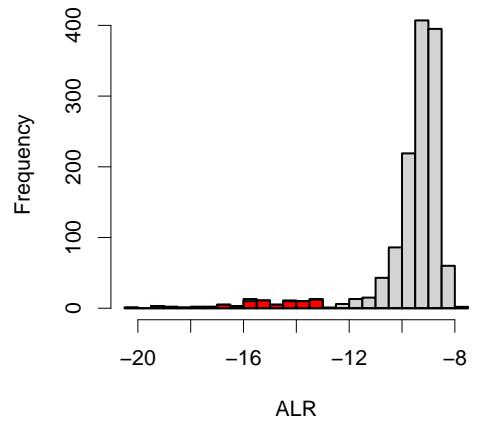
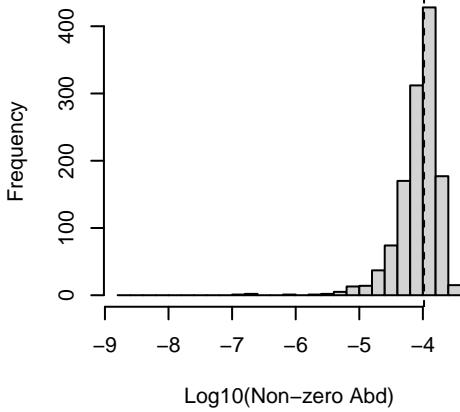
Zero Count/Abundance Samples: 13 (0.98%)

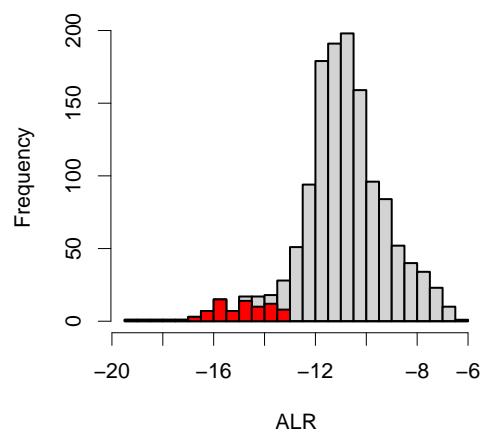
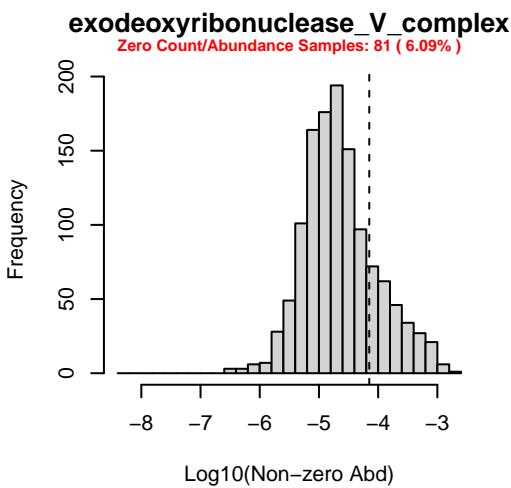
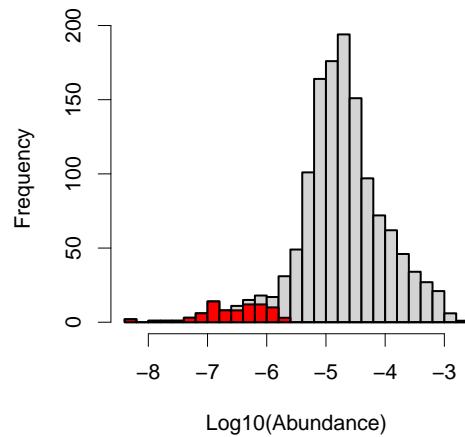
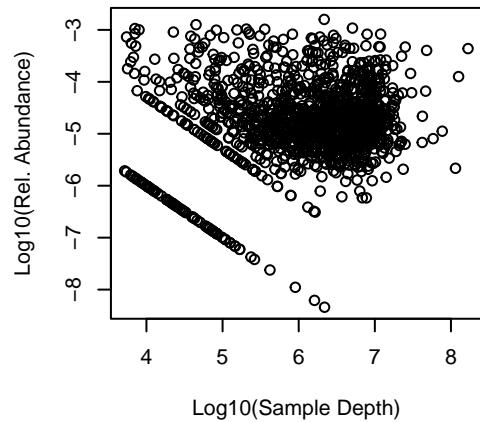
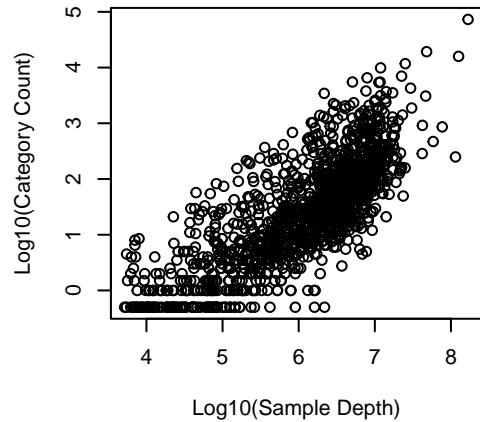
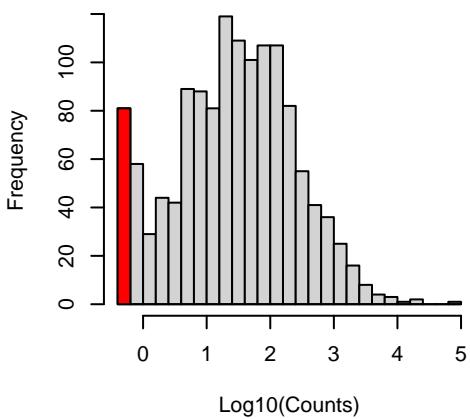


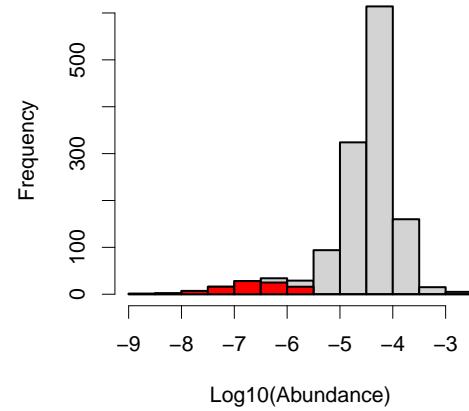
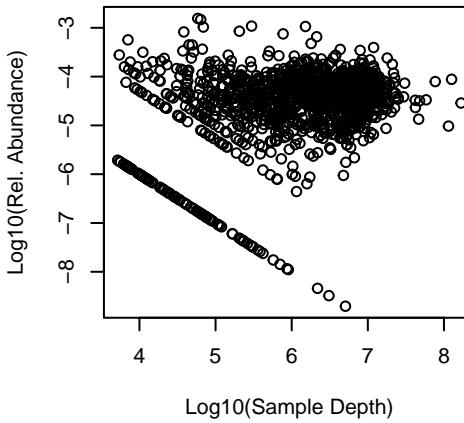
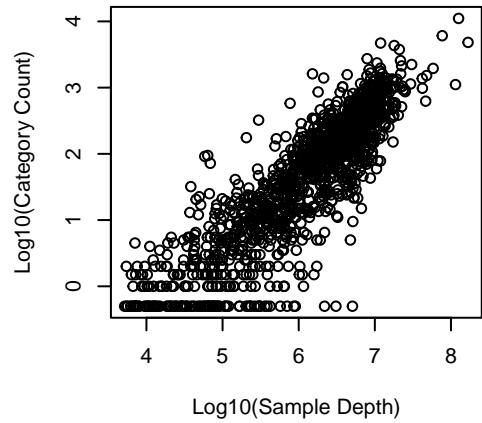
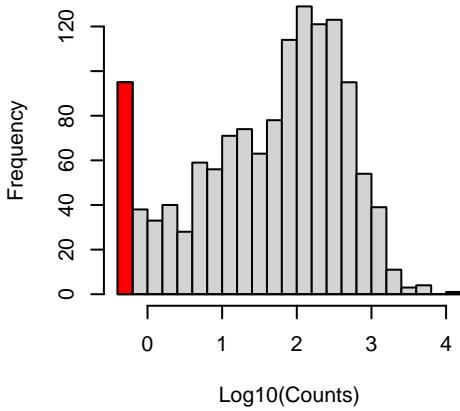


phopantethenoylcysteine_decarboxylase

Zero Count/Abundance Samples: 77 (5.79%)

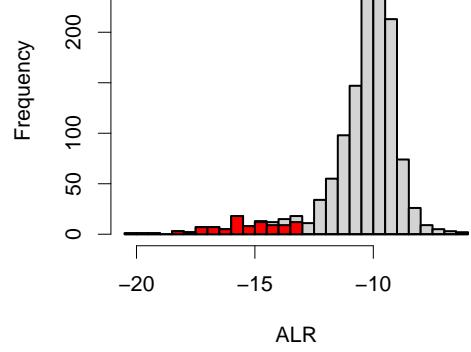
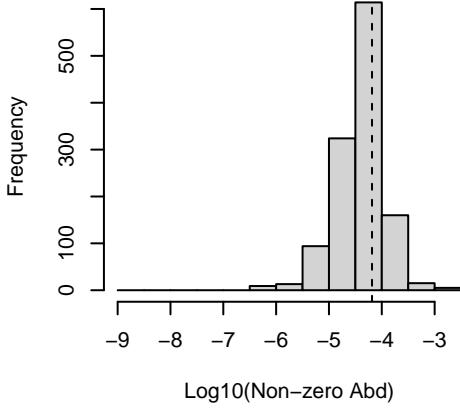


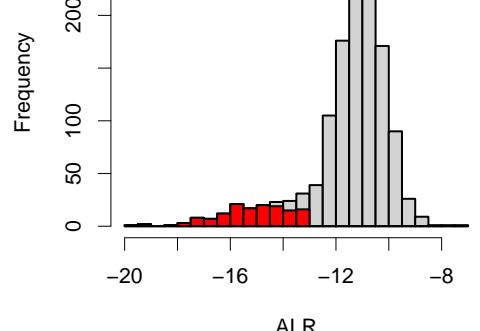
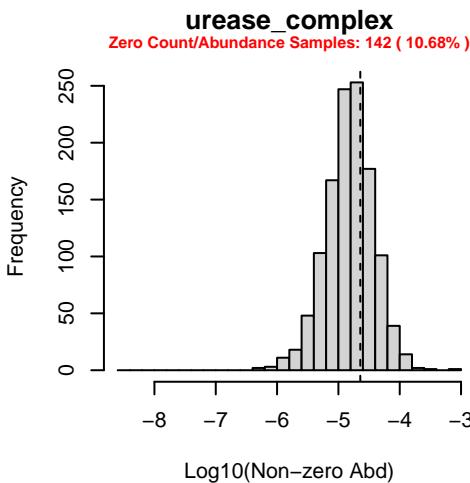
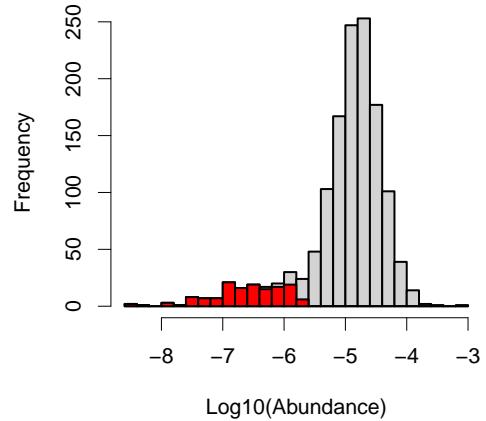
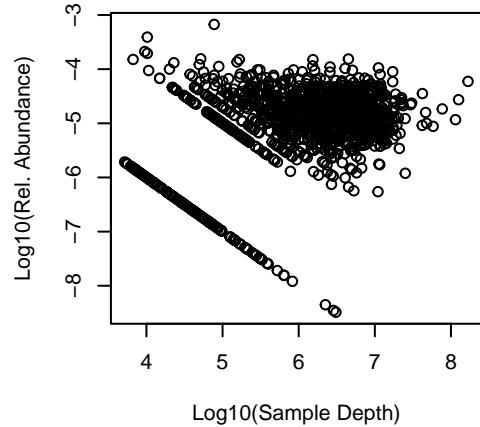
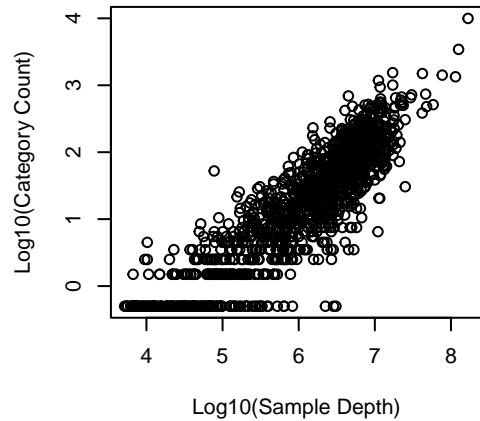
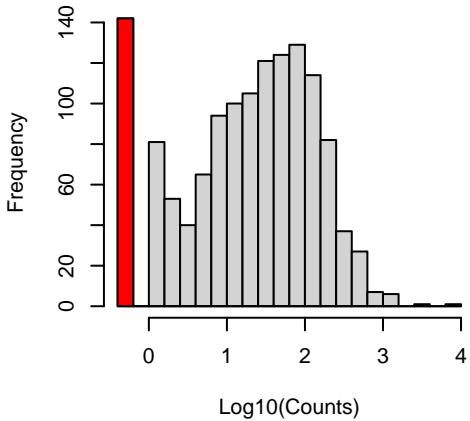


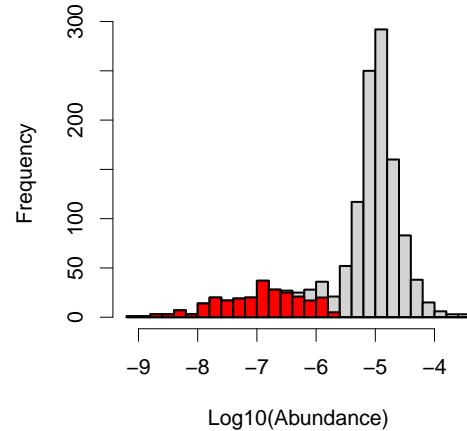
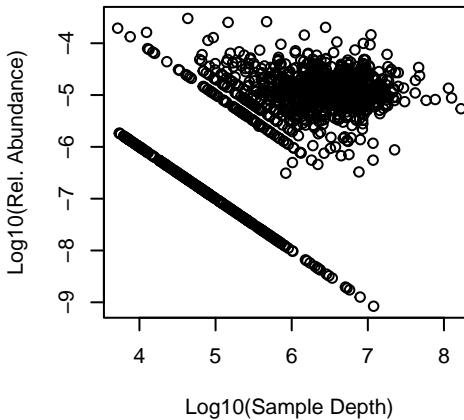
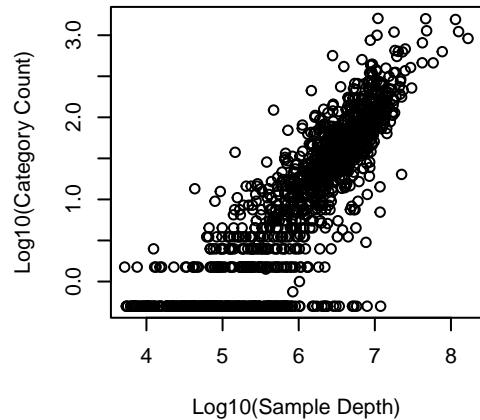
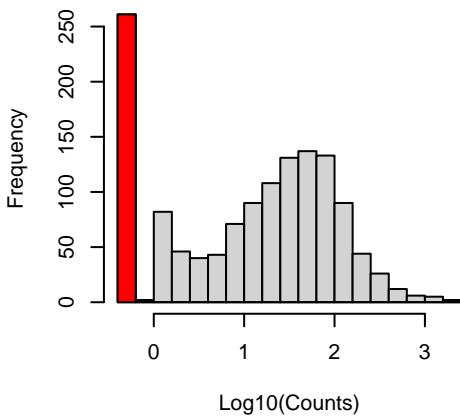


viral_terminase_large_subunit

Zero Count/Abundance Samples: 95 (7.15%)

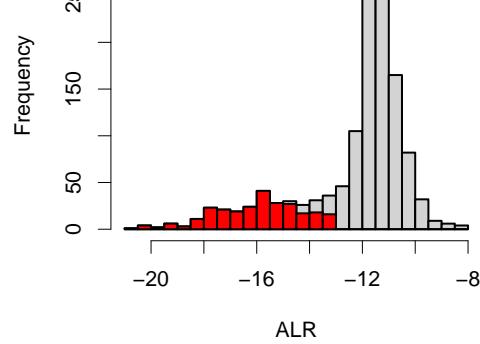
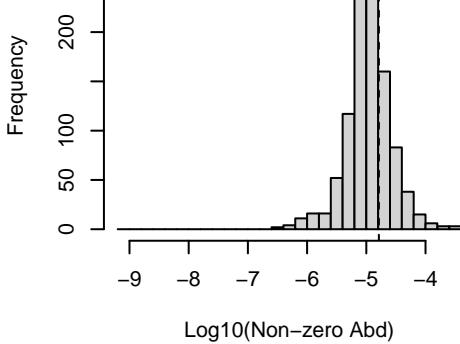


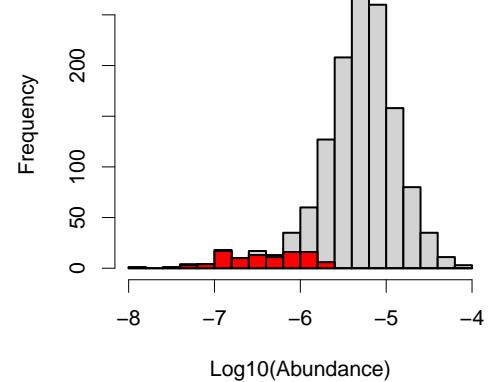
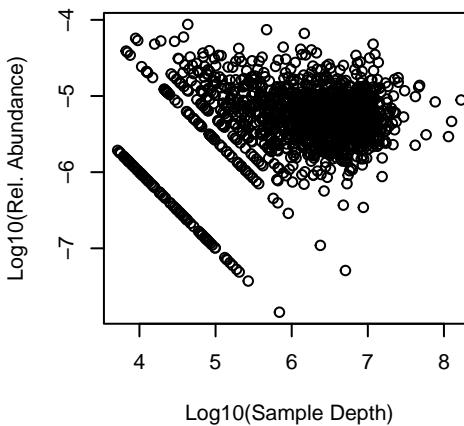
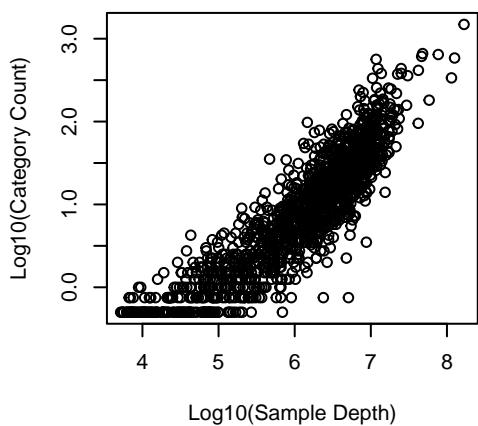
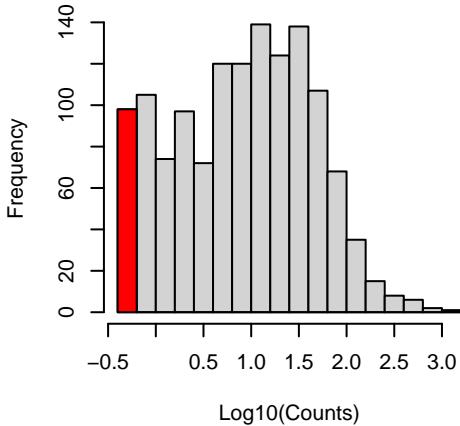




methylcrotonoyl_CoA_carboxylase_compl

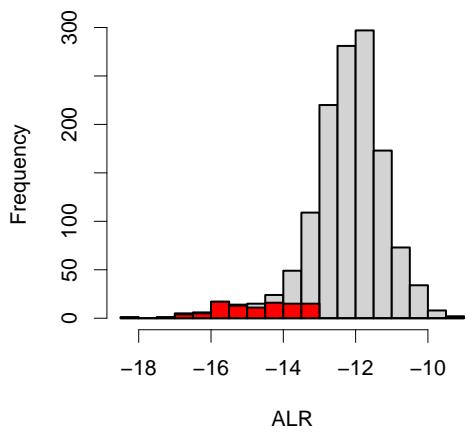
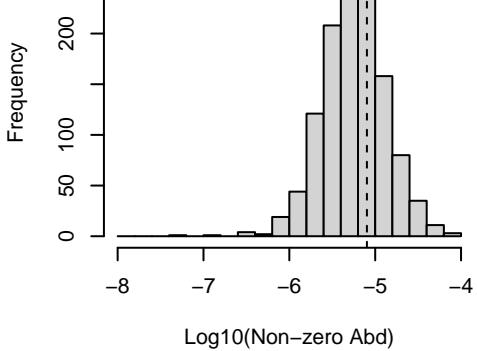
Zero Count/Abundance Samples: 261 (19.64%)

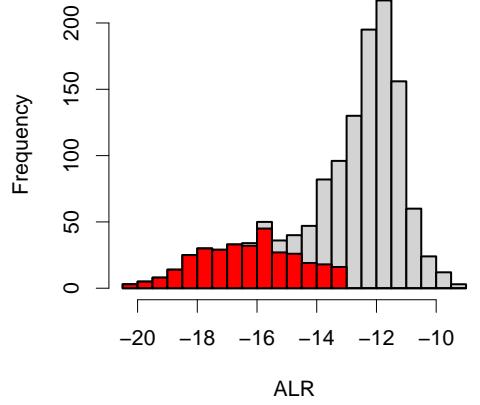
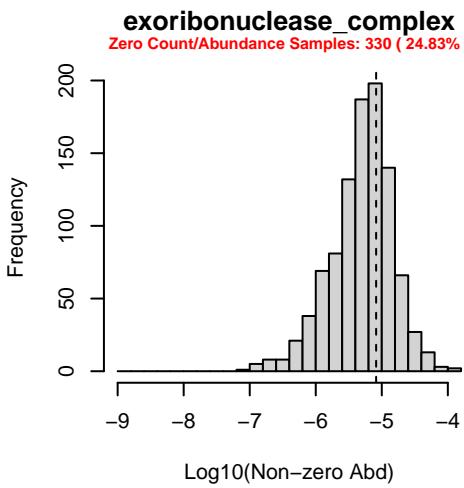
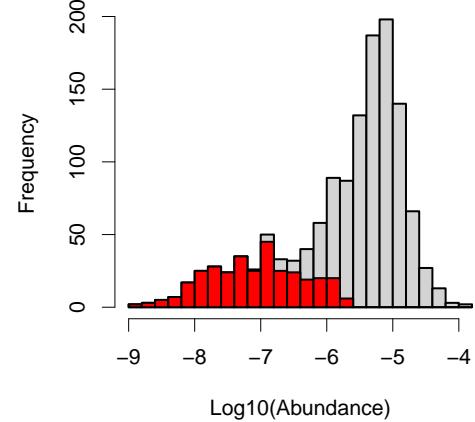
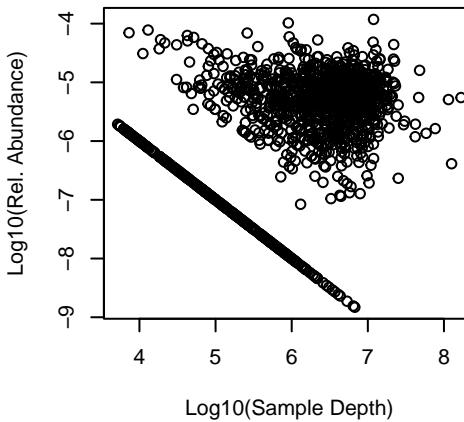
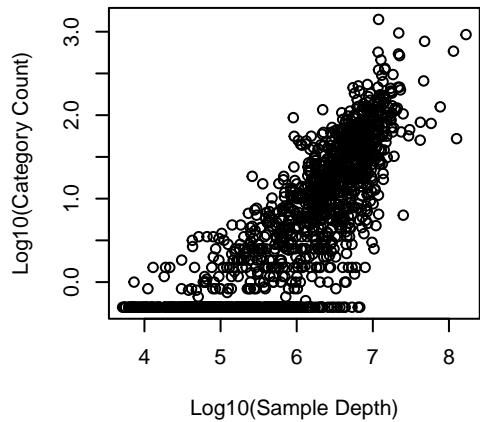
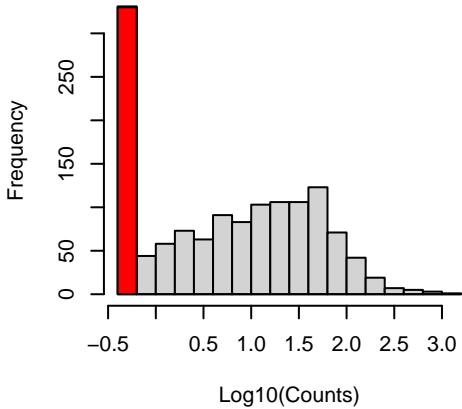


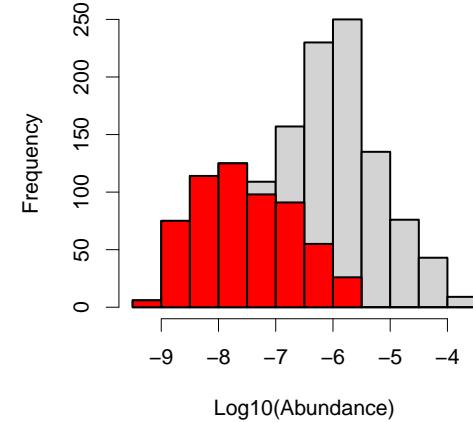
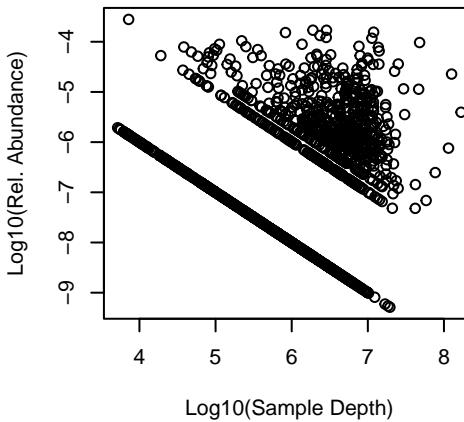
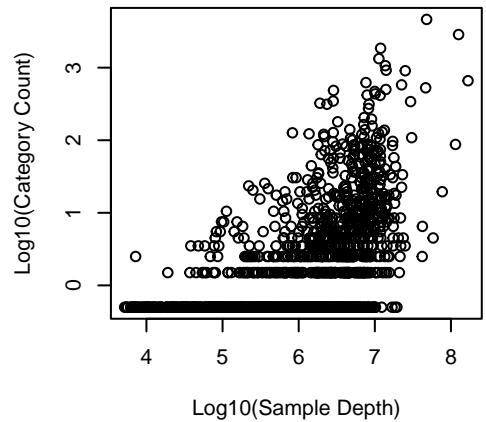
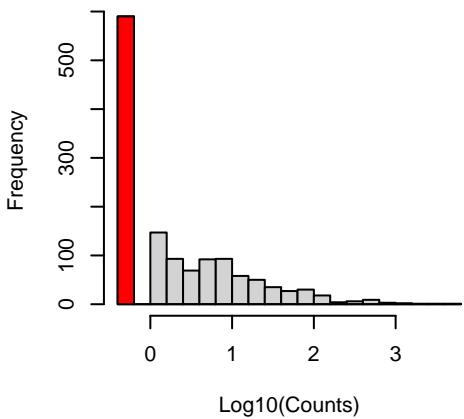


ethanolamine_ammonia_lyase_complex

Zero Count/Abundance Samples: 98 (7.37%)

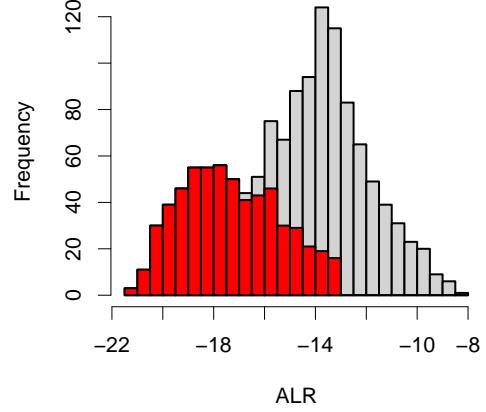
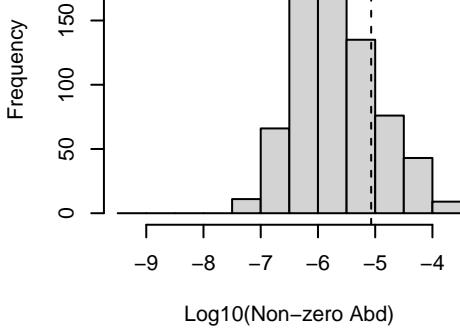


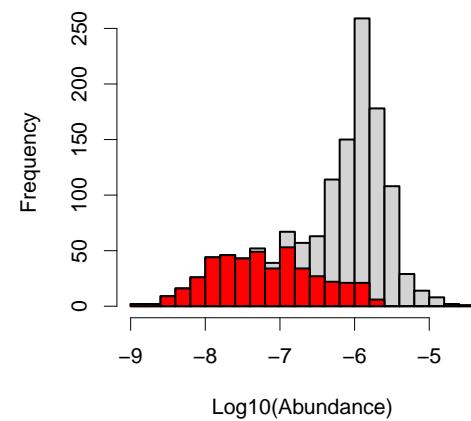
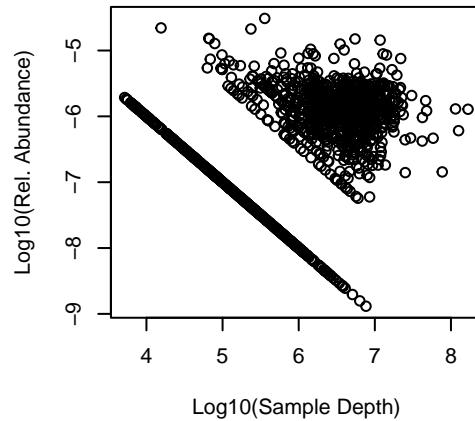
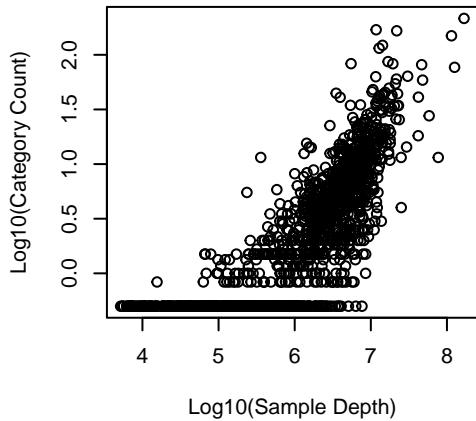
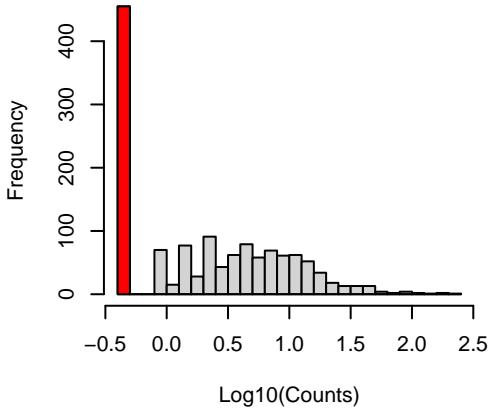




anthranilate_synthase_complex

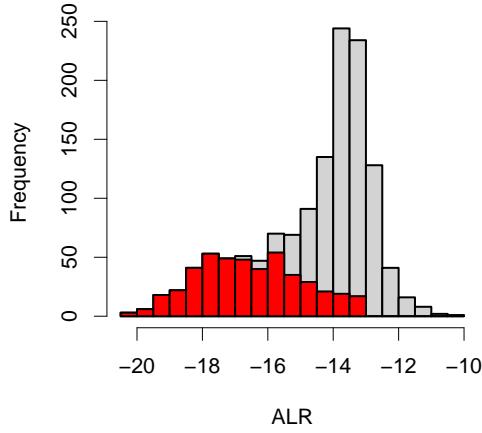
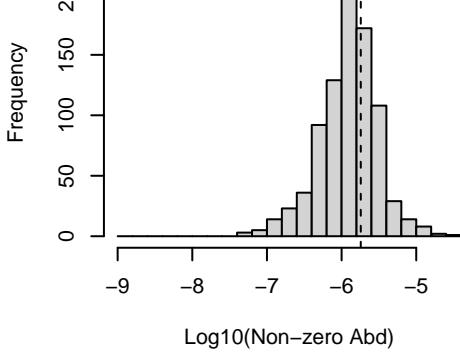
Zero Count/Abundance Samples: 590 (44.39%)

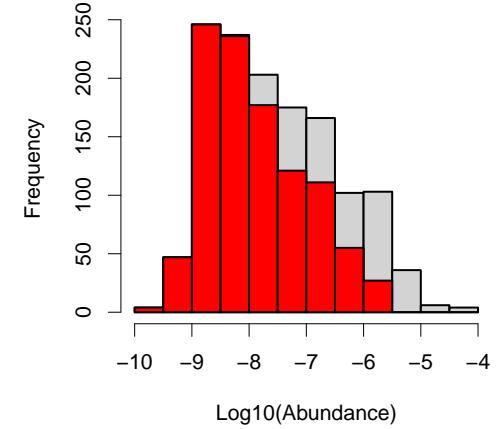
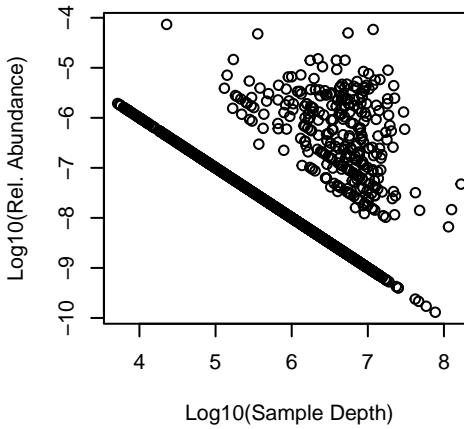
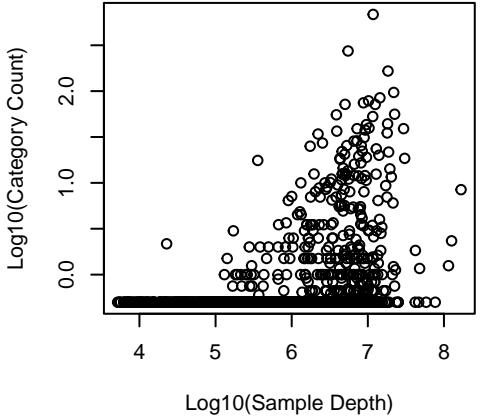
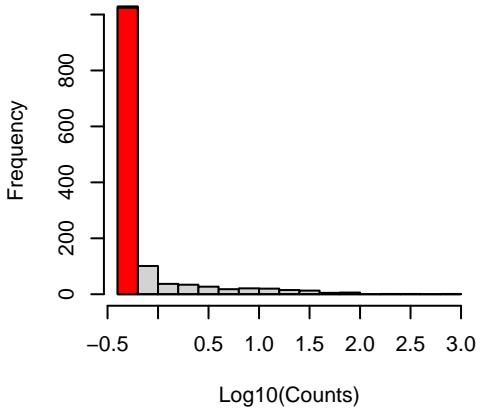




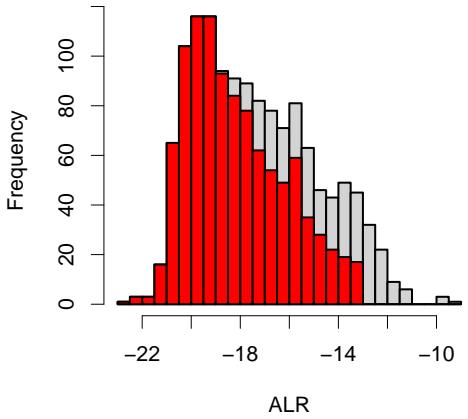
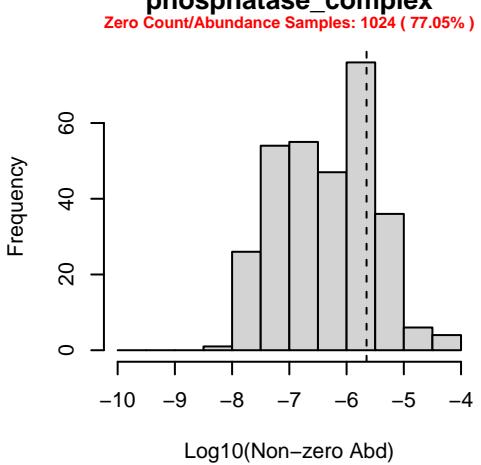
carbamoyl_phosphate_synthase_compl

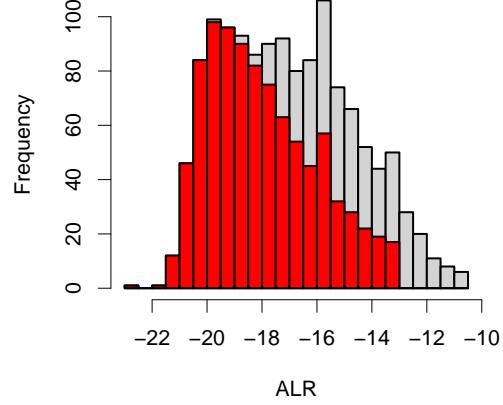
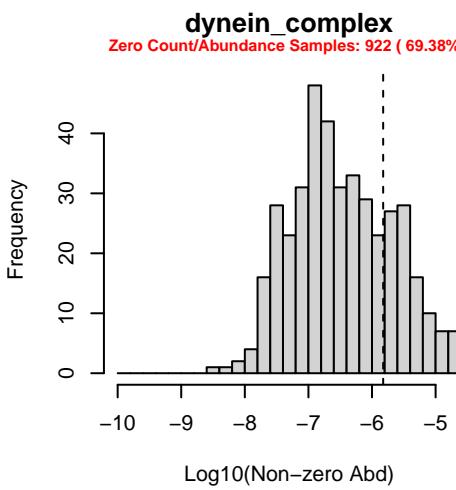
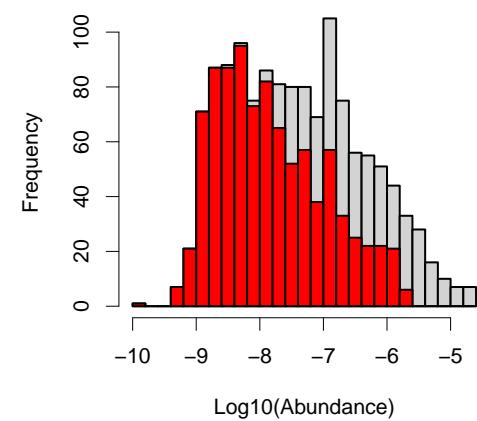
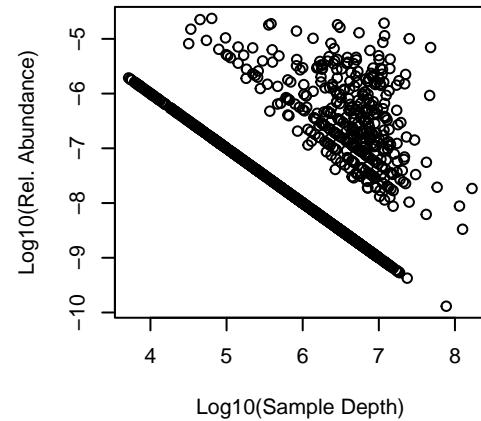
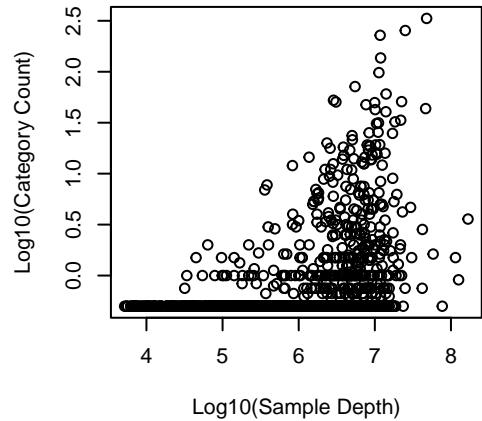
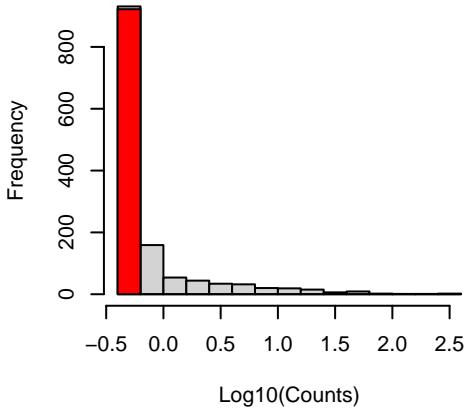
Zero Count/Abundance Samples: 455 (34.24%)

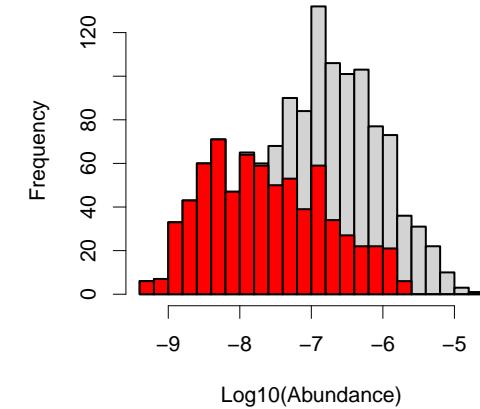
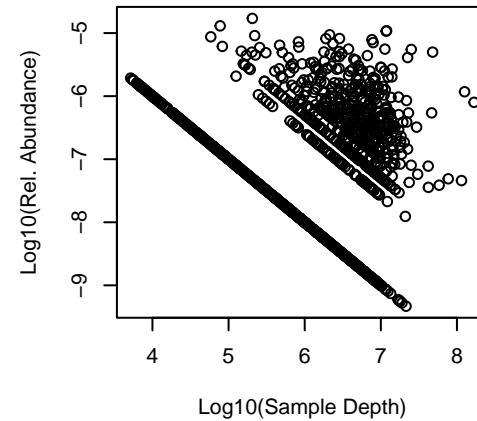
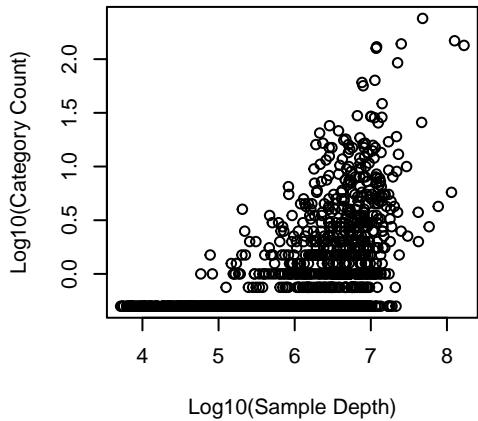
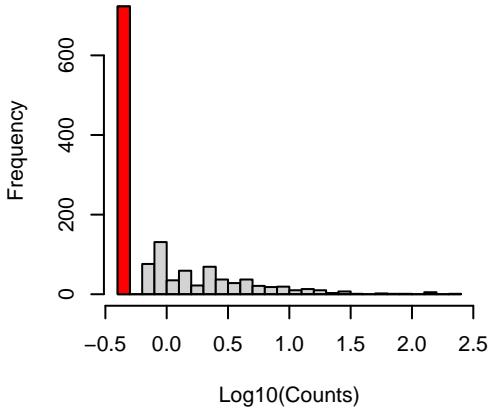




phosphatase_complex
Zero Count/Abundance Samples: 1024 (77.05%)

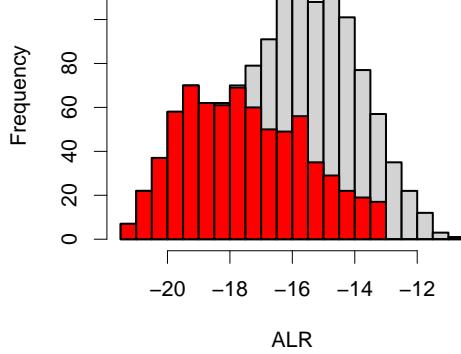
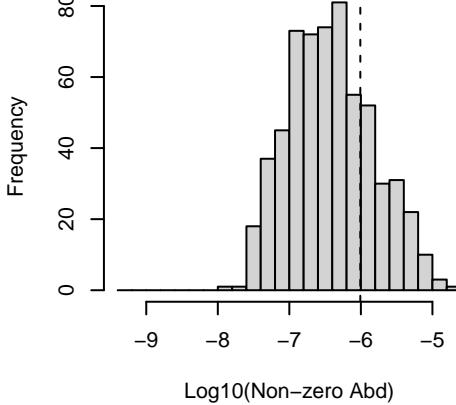


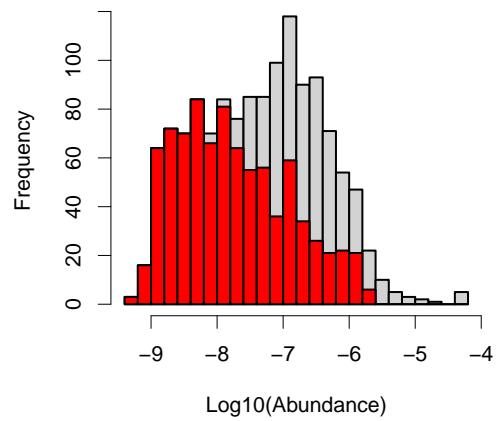
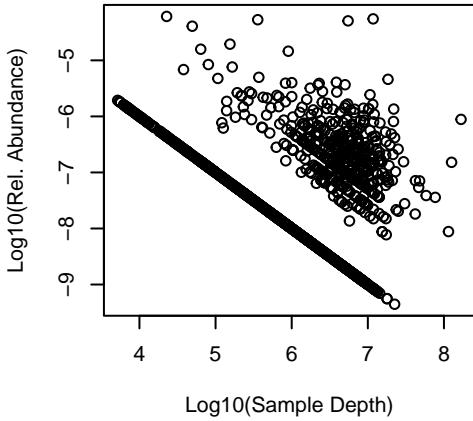
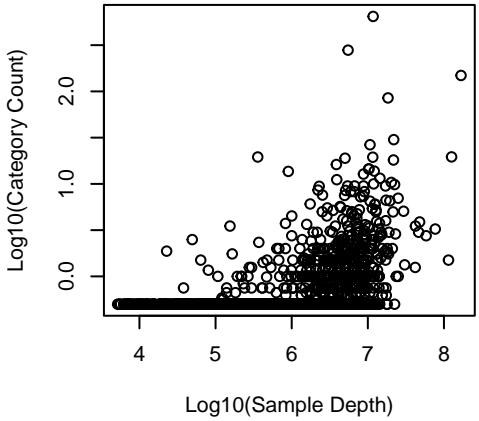
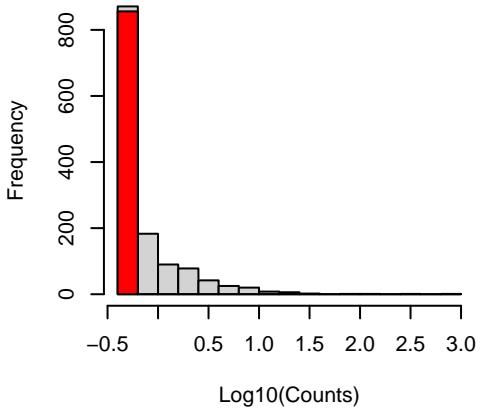




succinate_CoA_ligase_complex

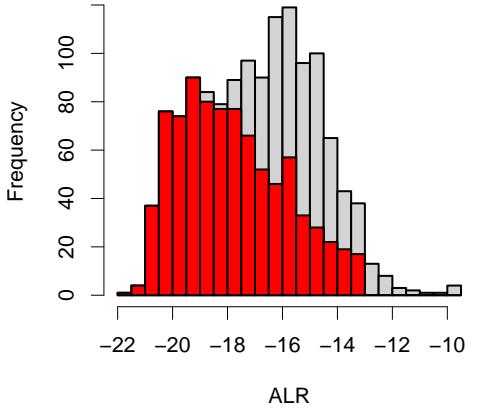
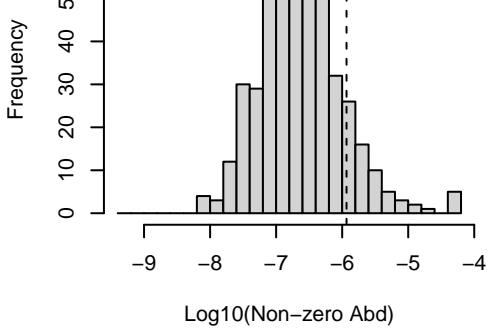
Zero Count/Abundance Samples: 723 (54.40%)

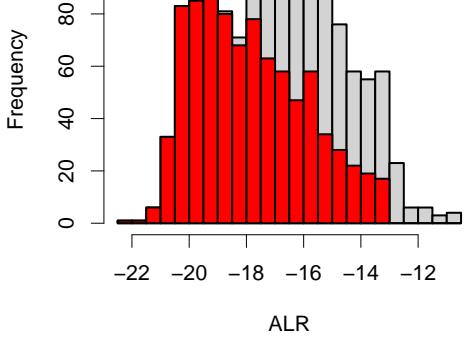
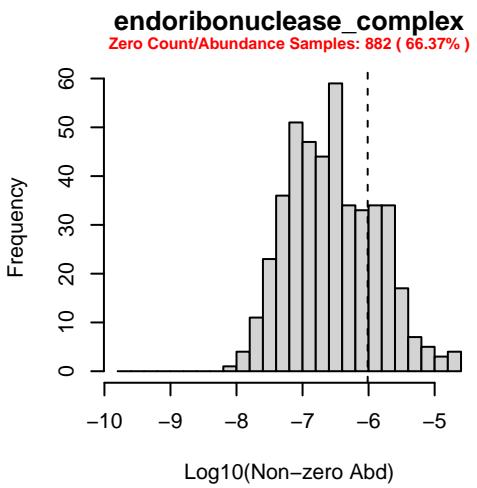
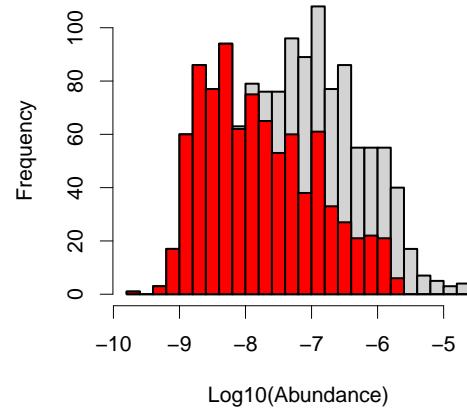
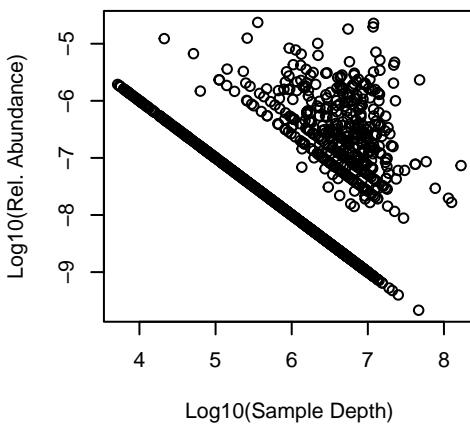
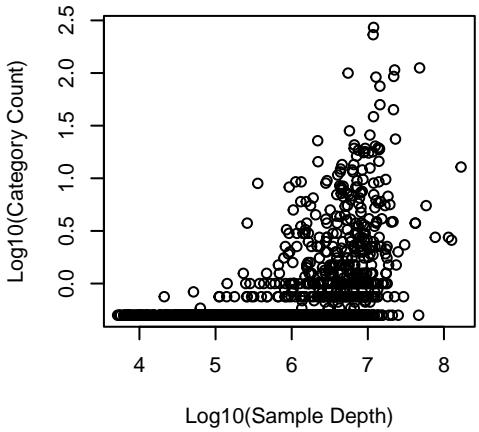
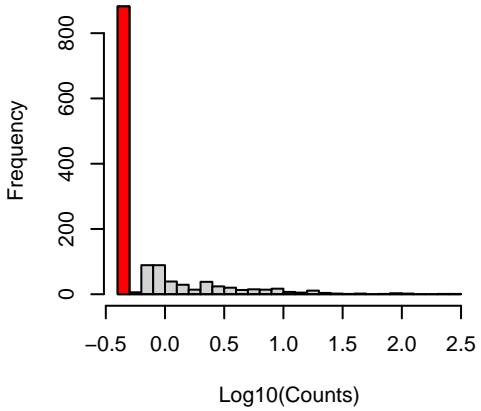


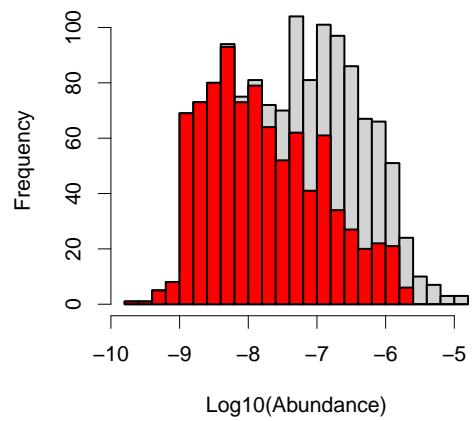
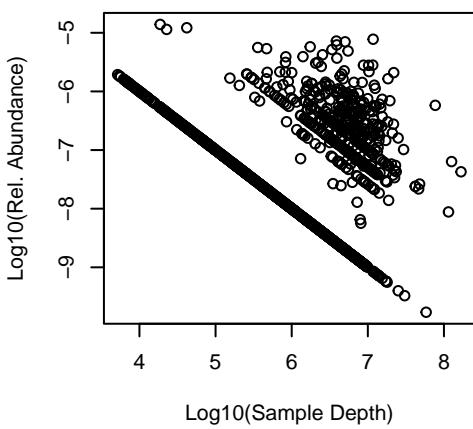
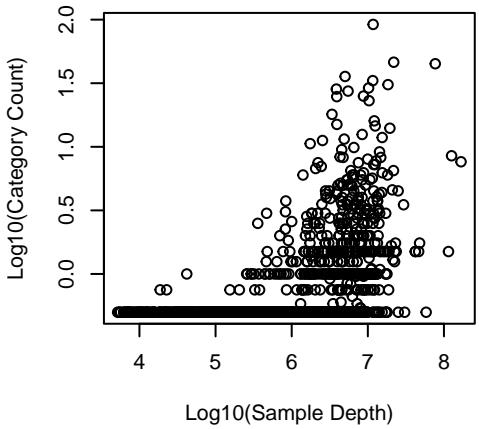
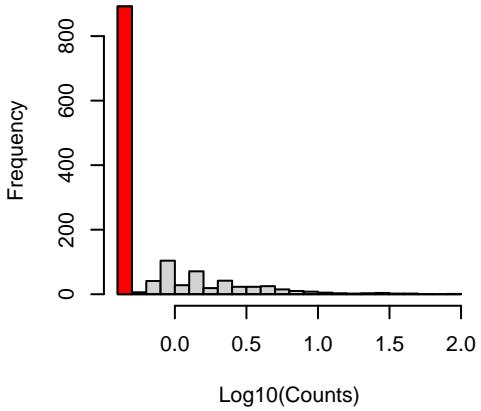


ATPase_complex

Zero Count/Abundance Samples: 856 (64.41%)

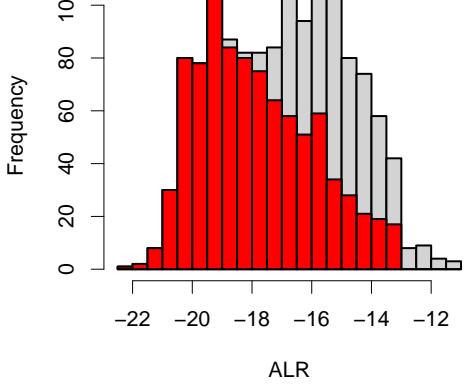
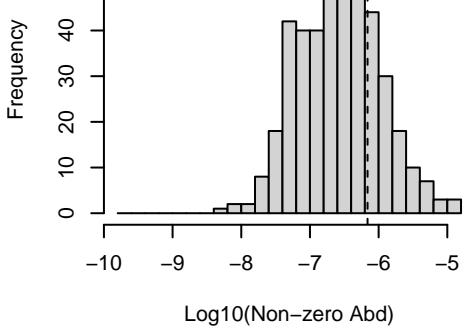


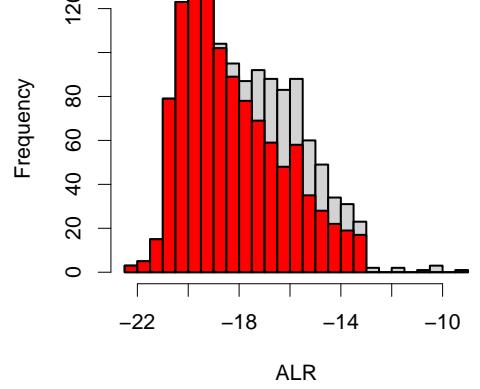
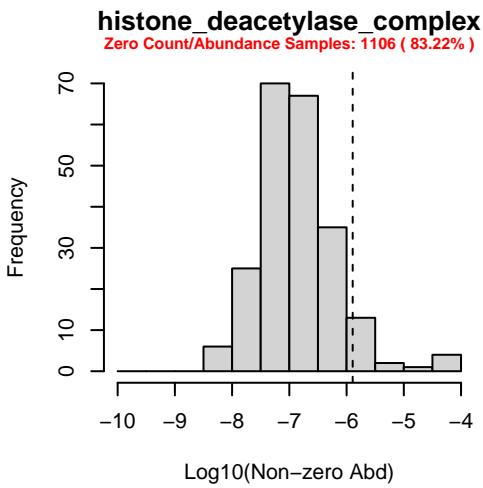
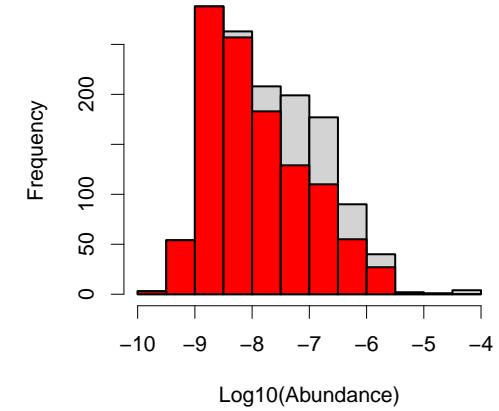
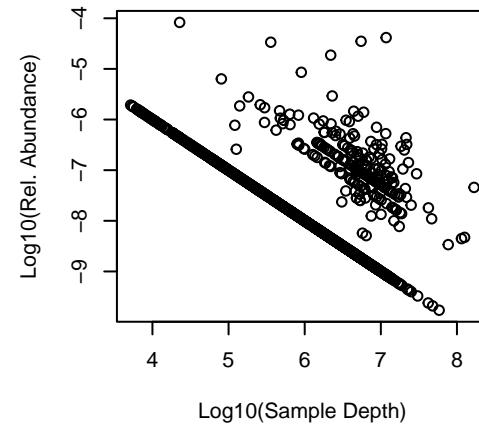
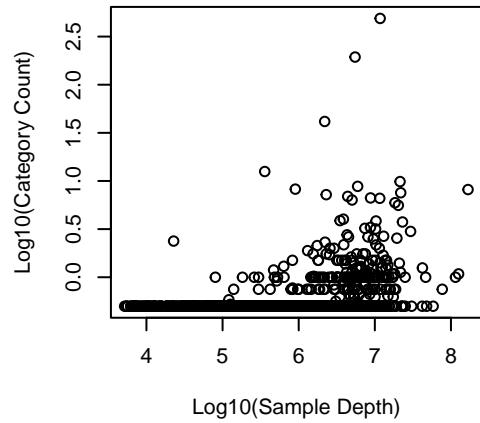
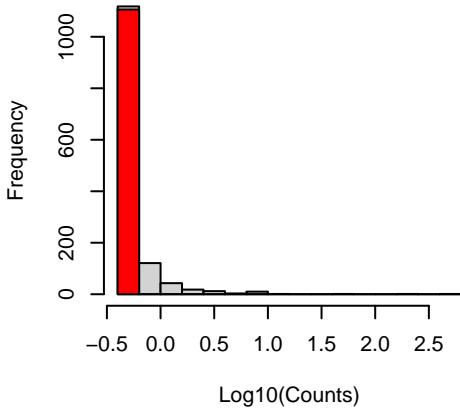


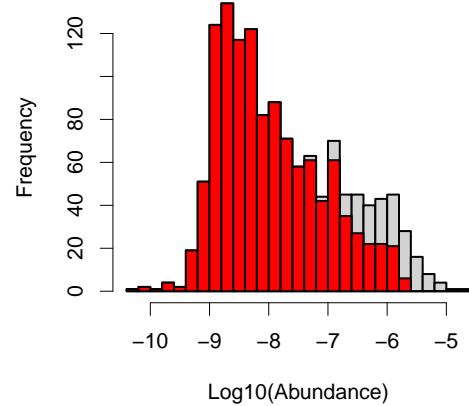
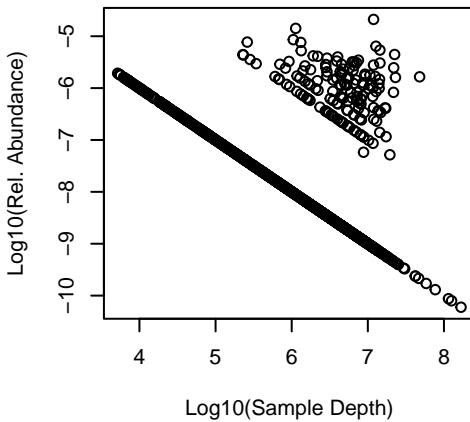
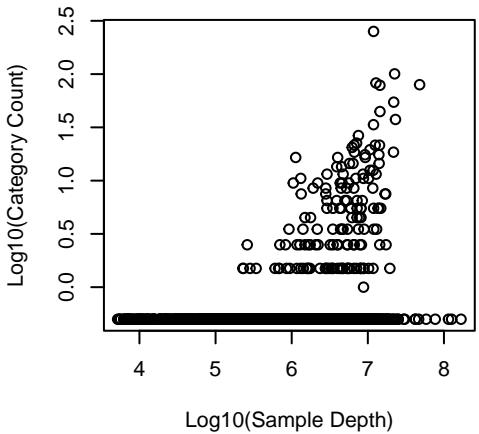
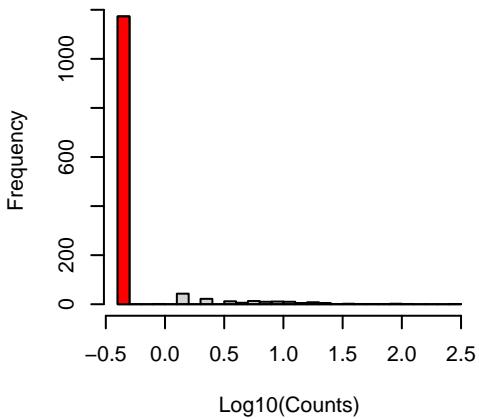


catalytic_step_2_spliceosome

Zero Count/Abundance Samples: 892 (67.12%)

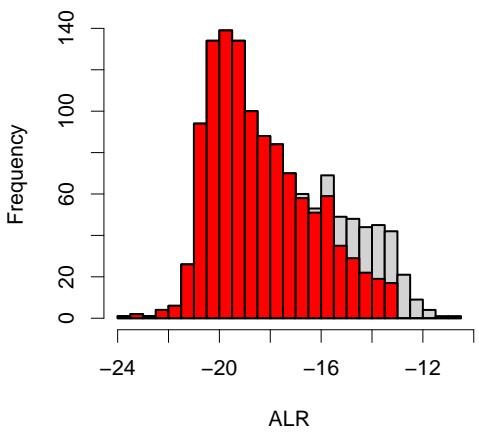
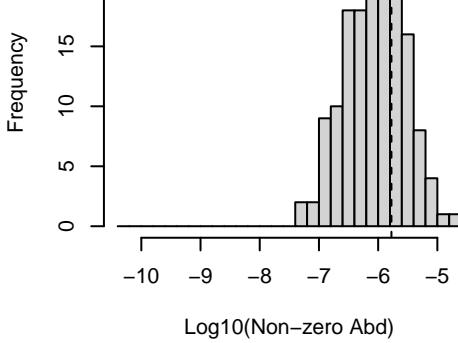


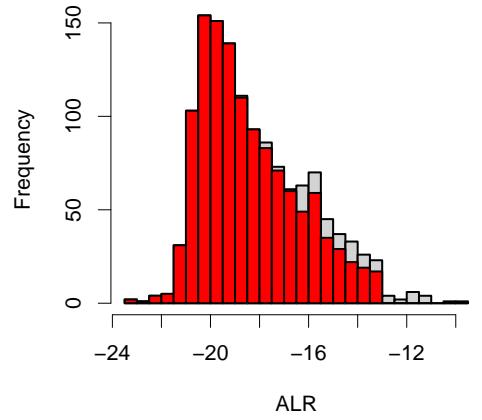
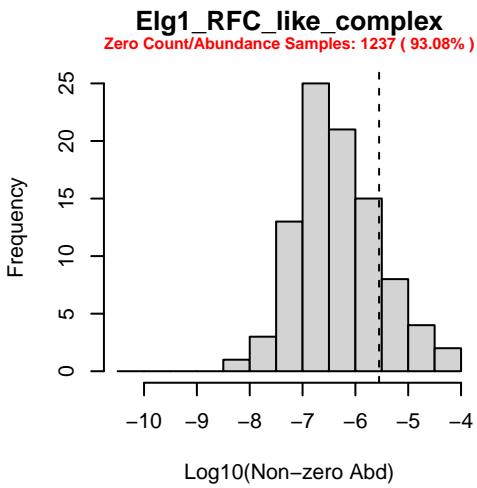
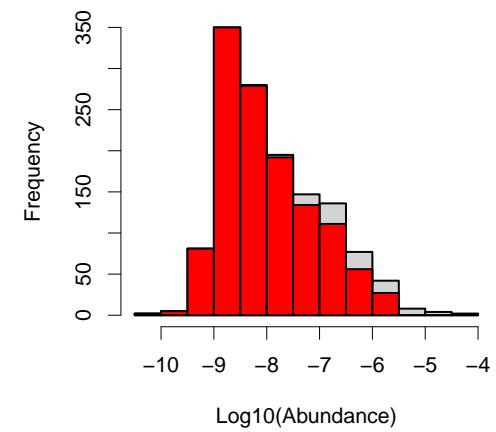
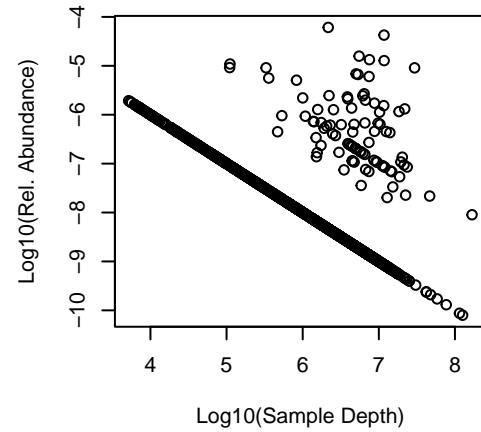
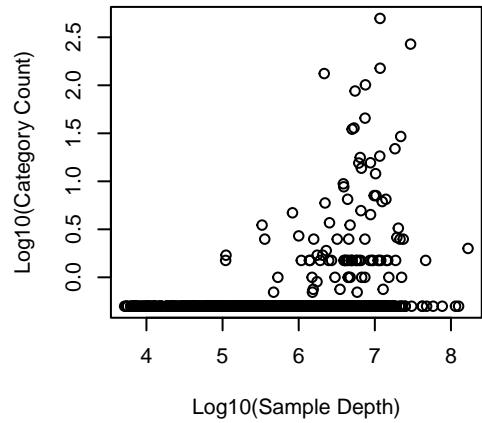
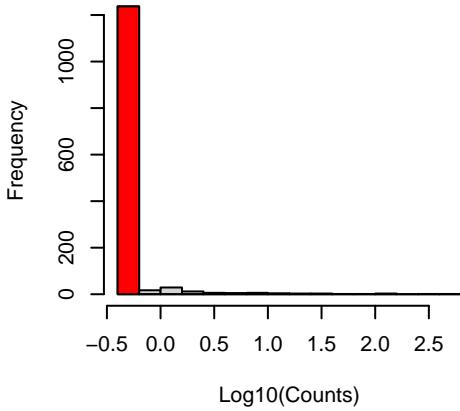


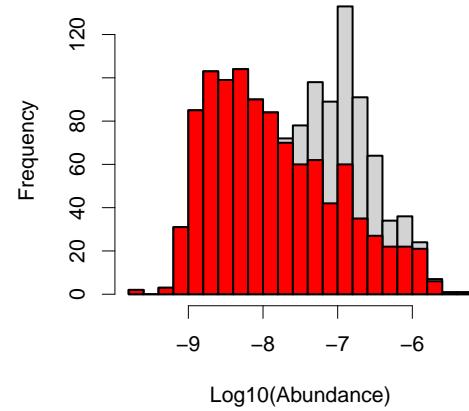
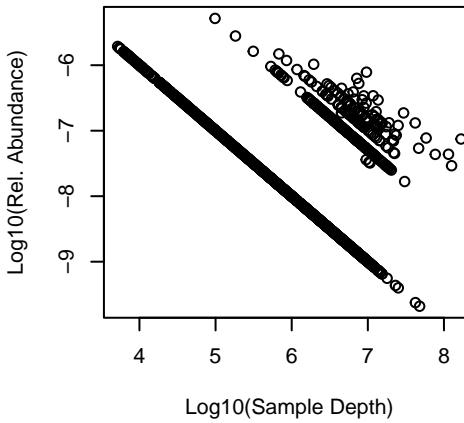
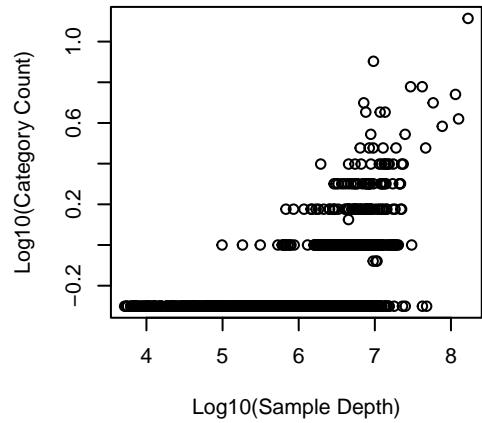
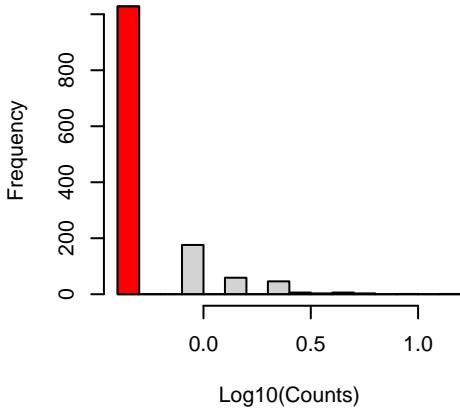


phenylalanine_tRNA_ligase_complex

Zero Count/Abundance Samples: 1173 (88.26%)

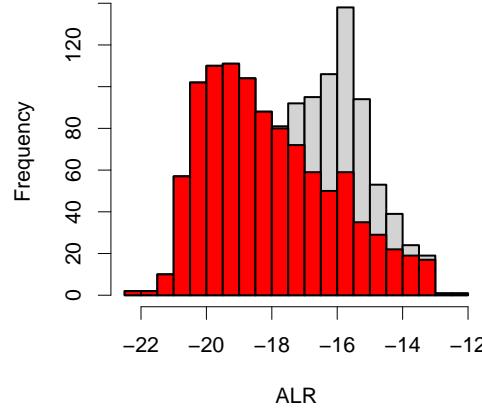
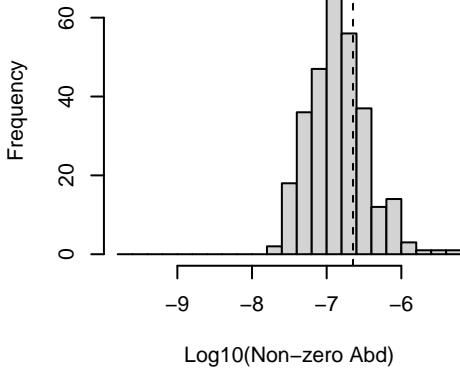


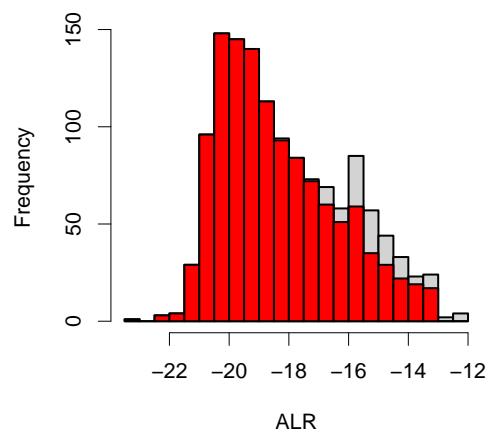
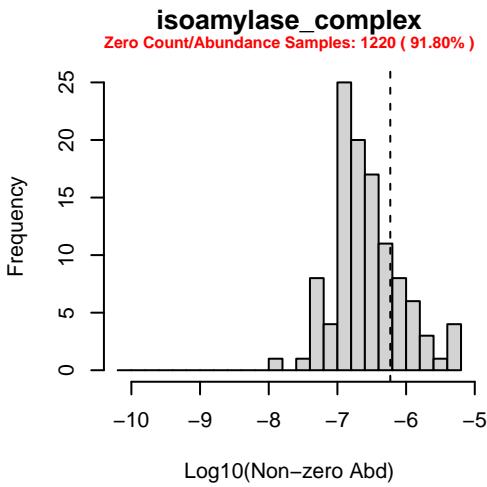
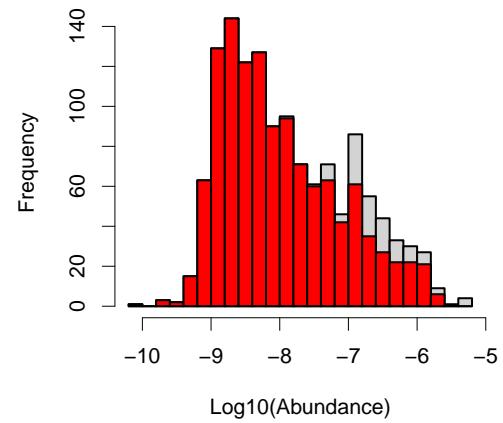
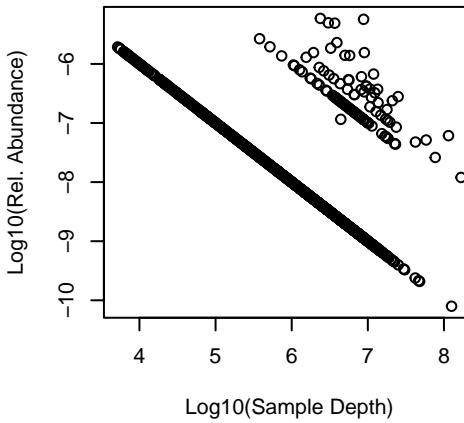
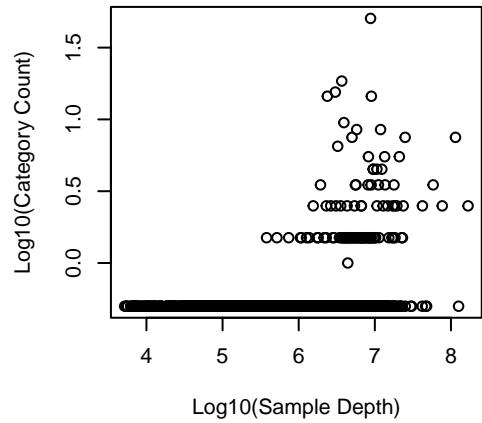
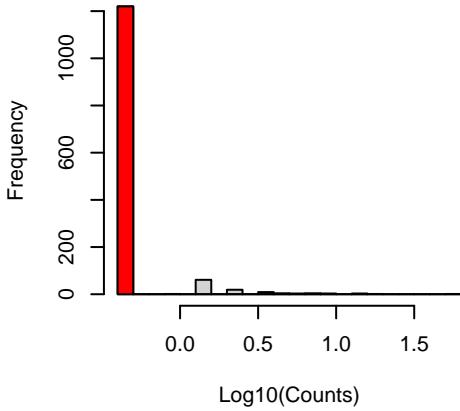


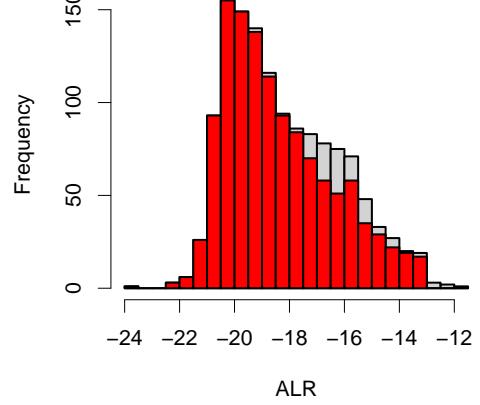
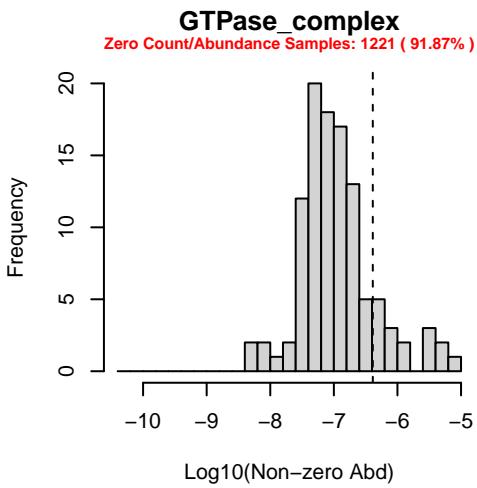
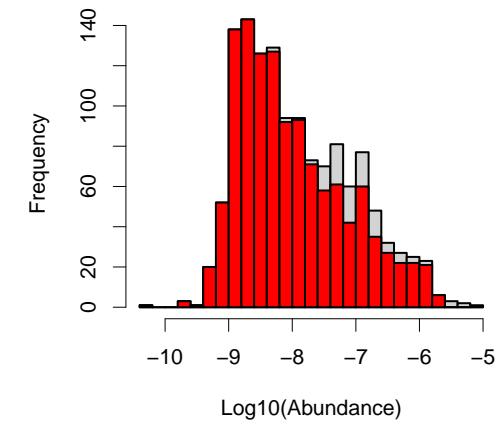
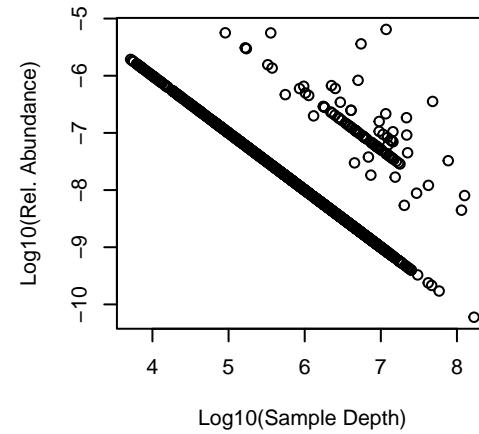
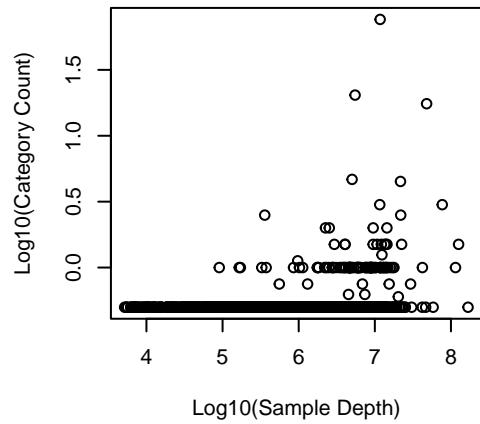
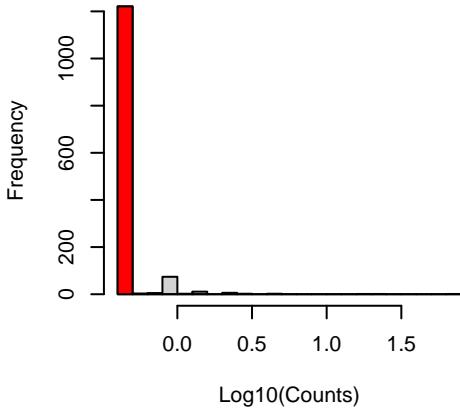


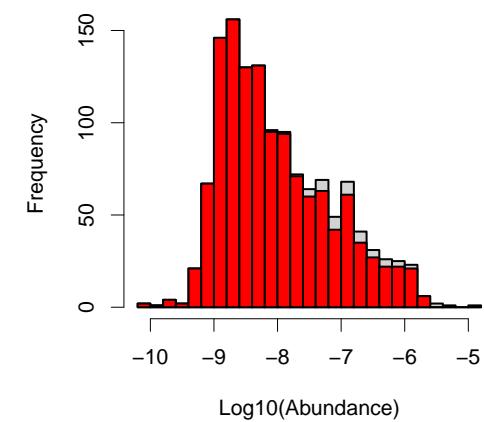
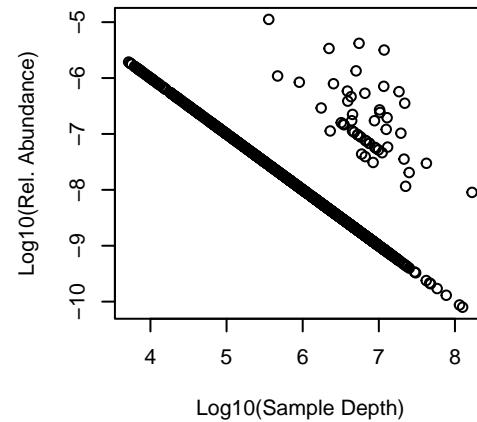
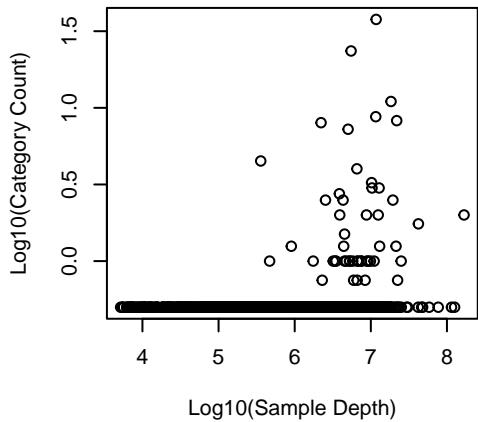
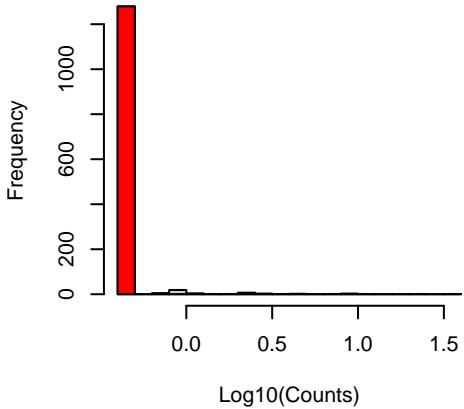
catalytic_complex

Zero Count/Abundance Samples: 1028 (77.35%)



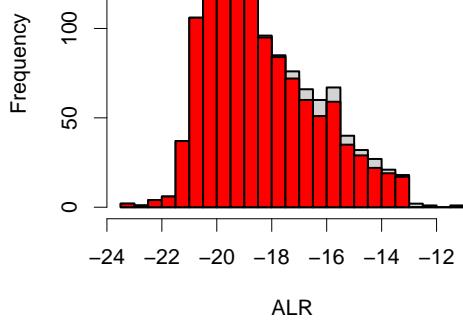
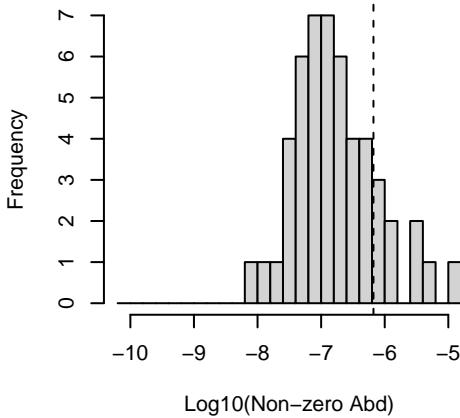


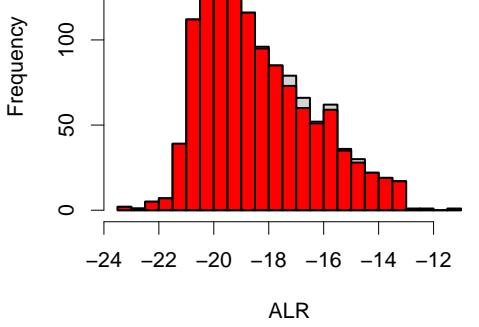
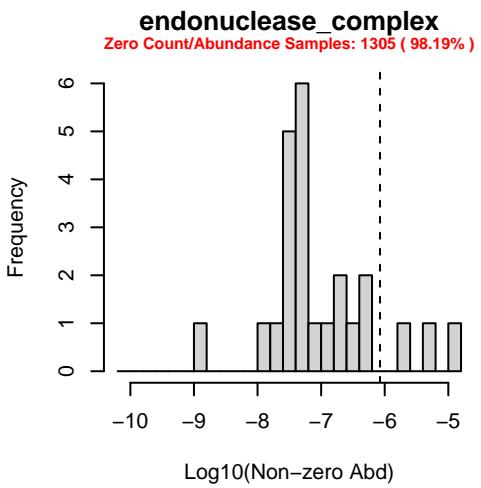
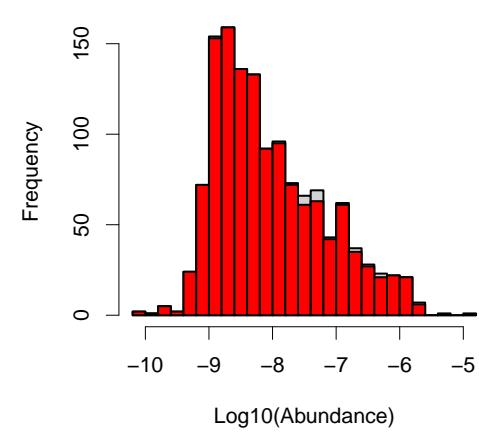
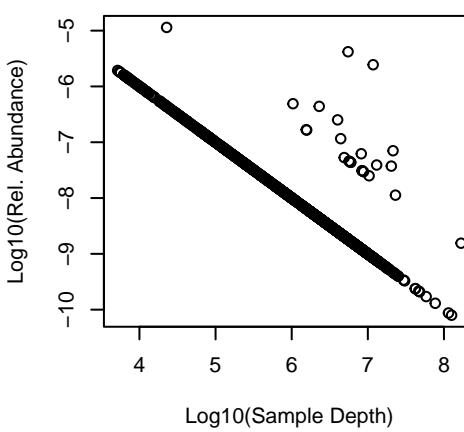
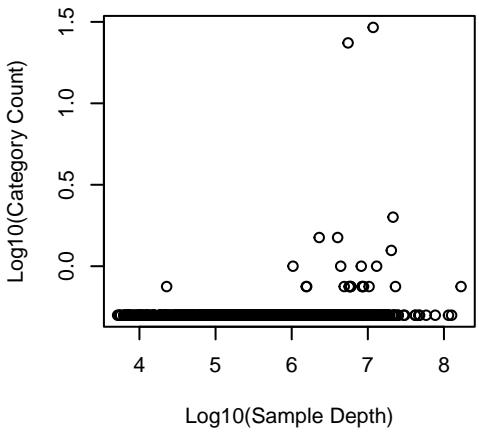
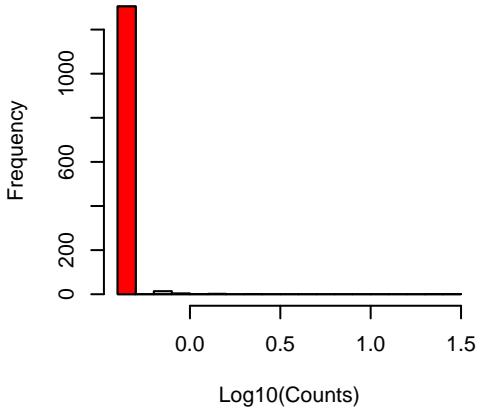


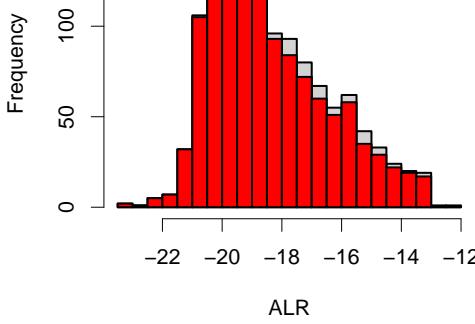
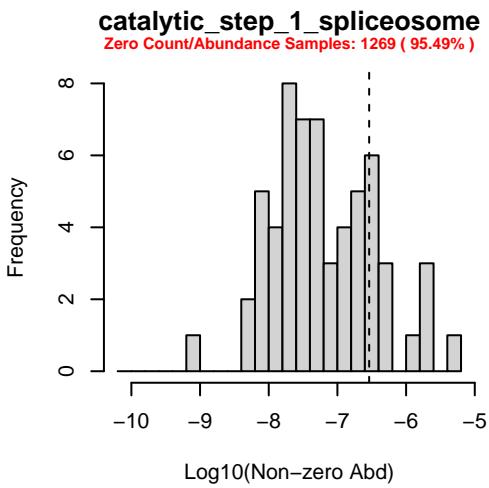
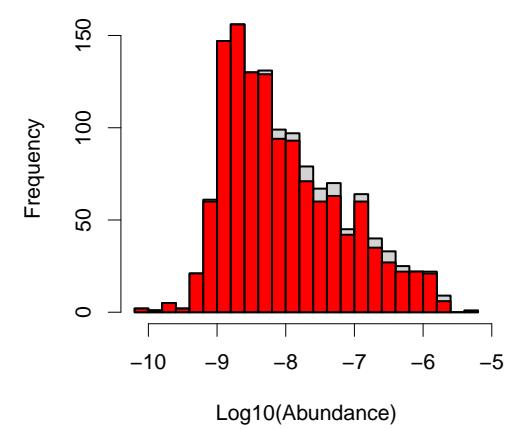
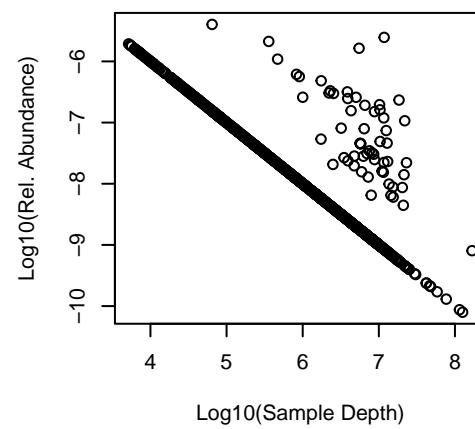
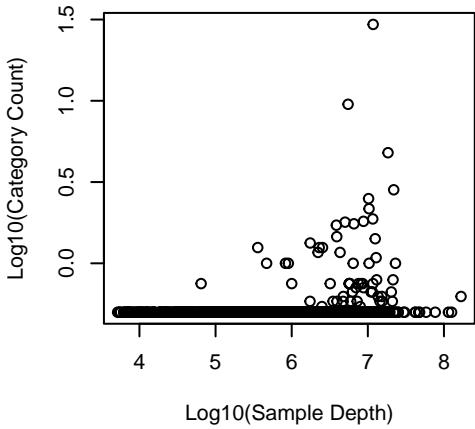
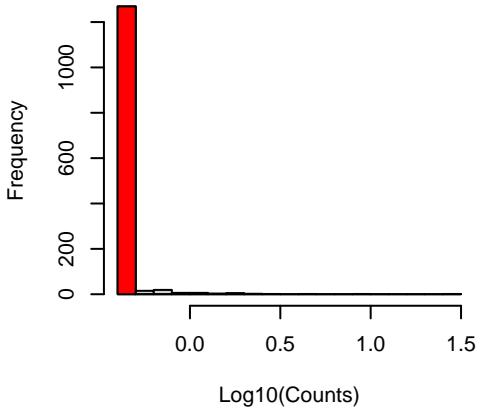


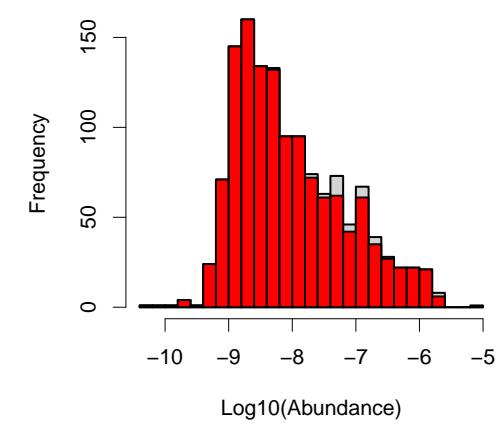
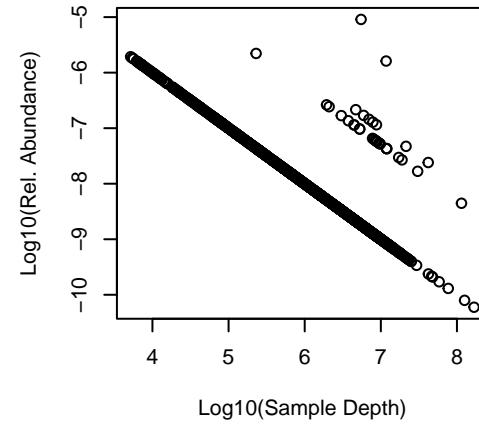
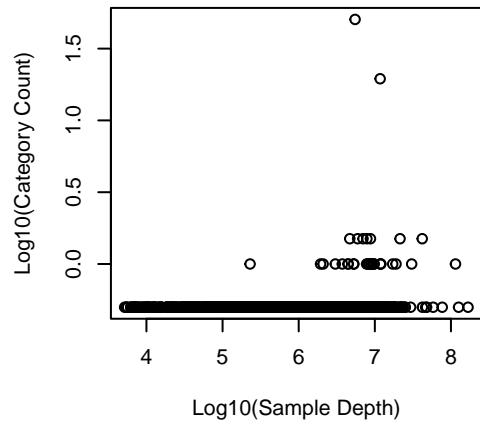
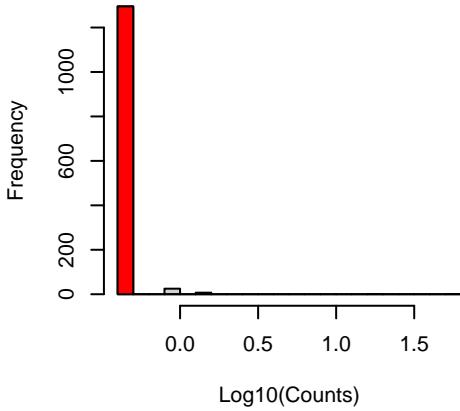
SUMO_activating_enzyme_complex

Zero Count/Abundance Samples: 1279 (96.24%)



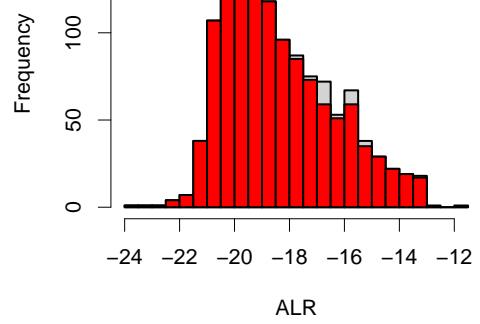
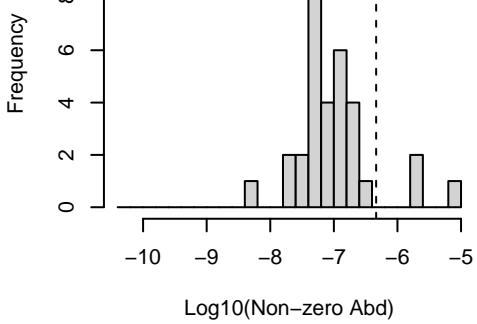


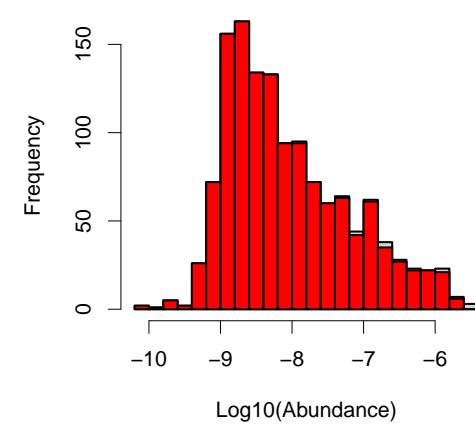
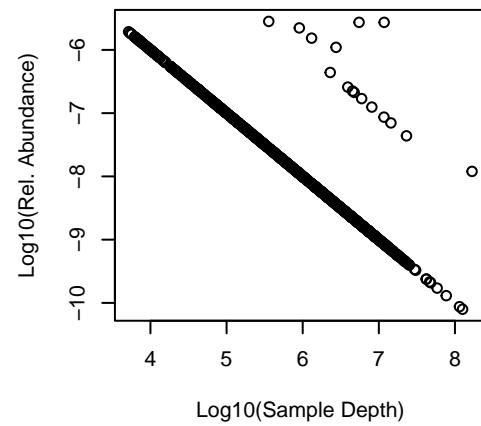
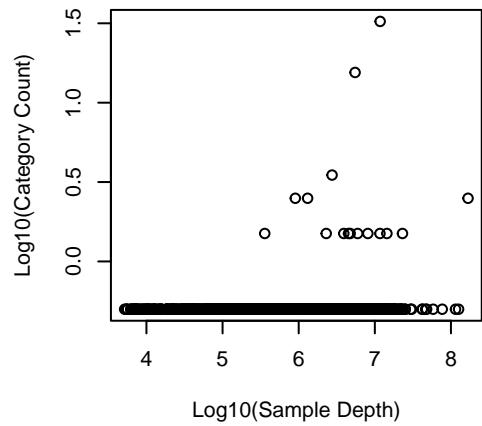
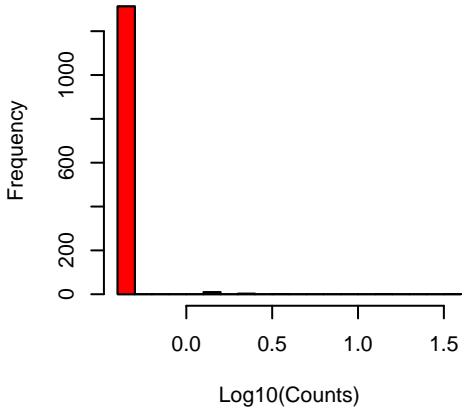




glutaminase_complex

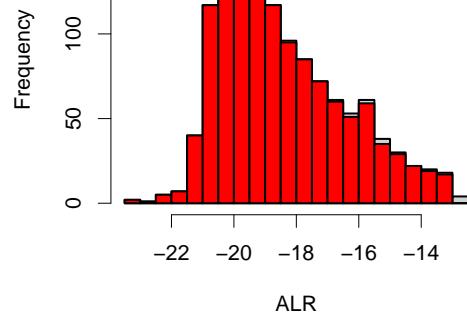
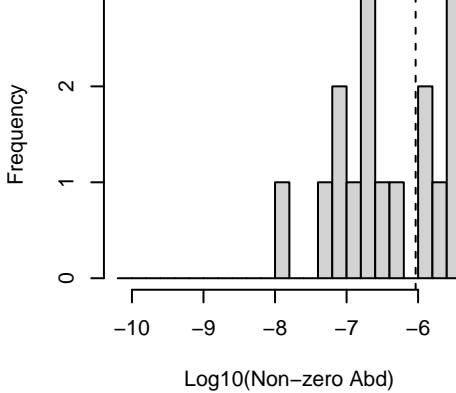
Zero Count/Abundance Samples: 1295 (97.44%)

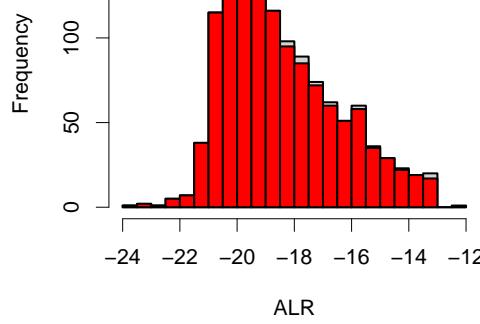
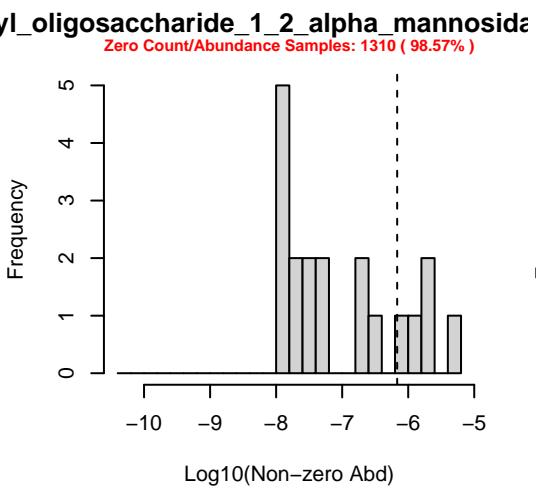
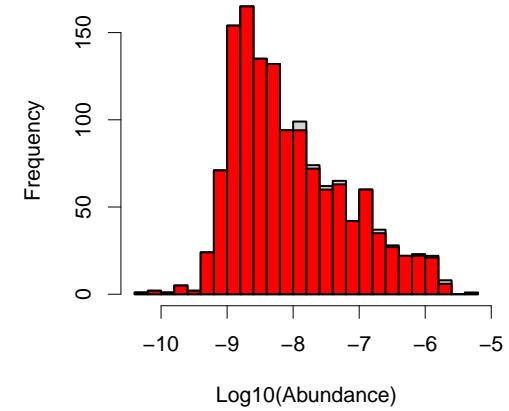
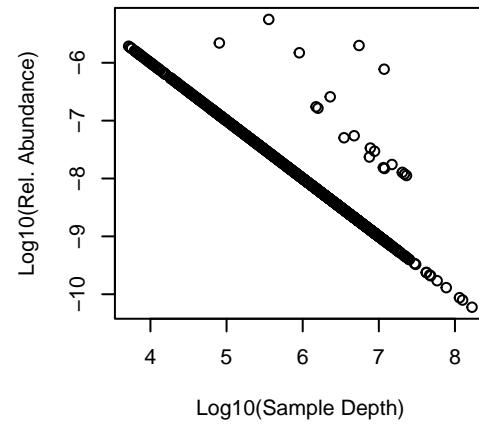
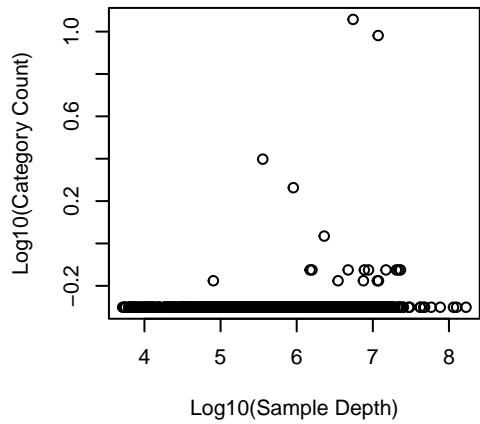
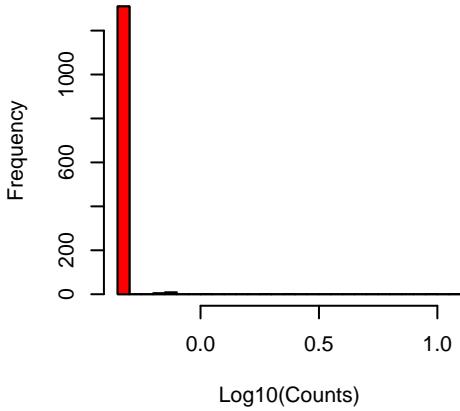


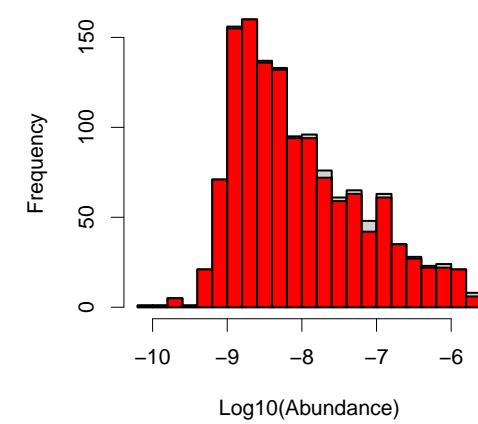
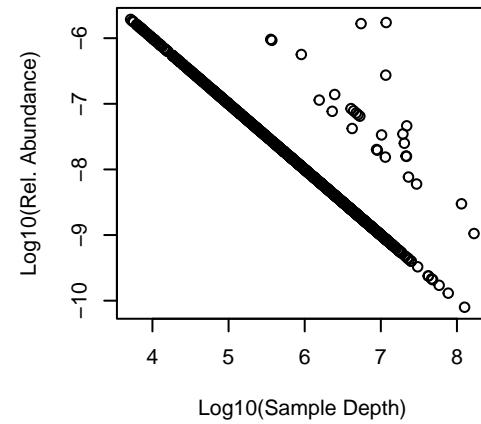
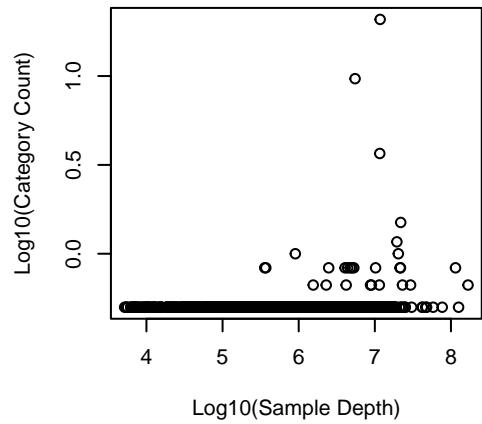
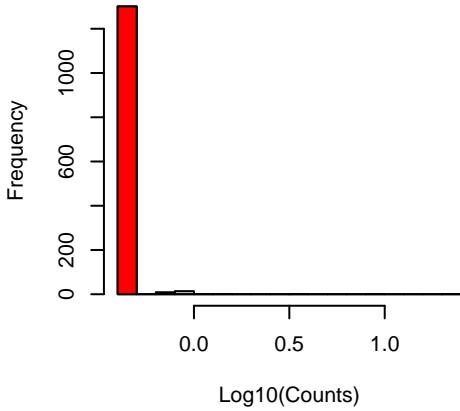


glutathione_synthase_complex

Zero Count/Abundance Samples: 1313 (98.80%)

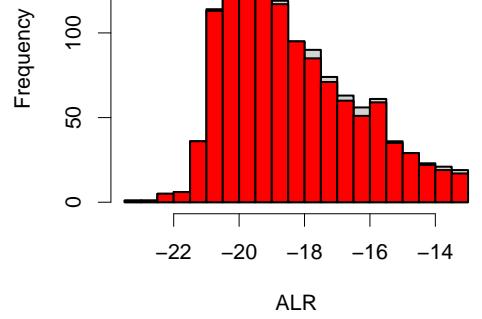
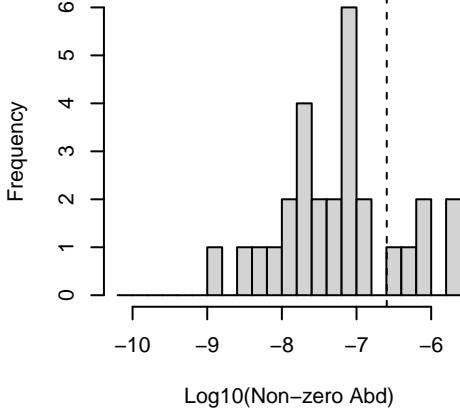


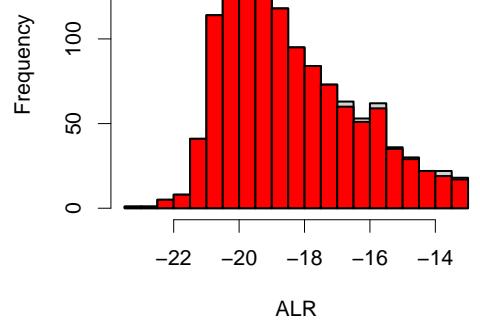
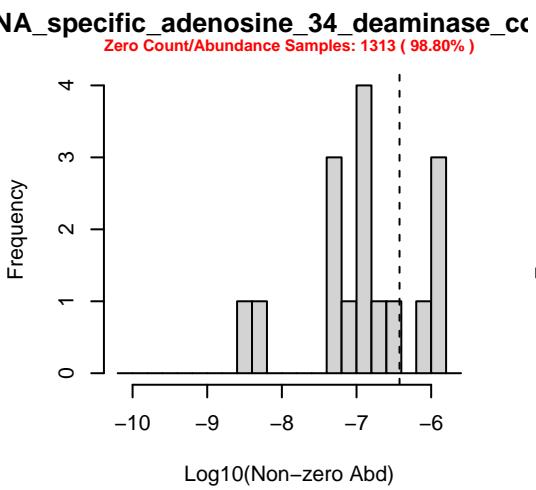
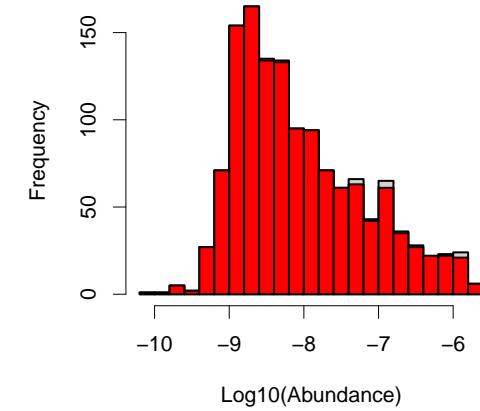
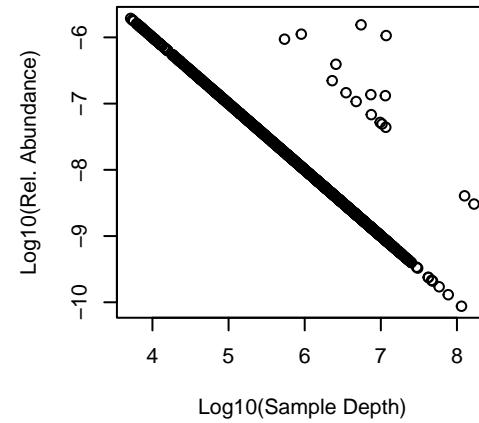
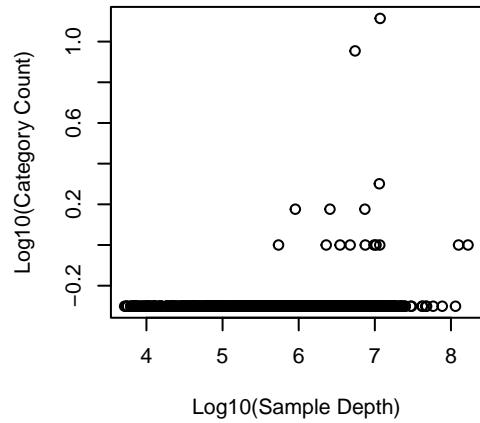
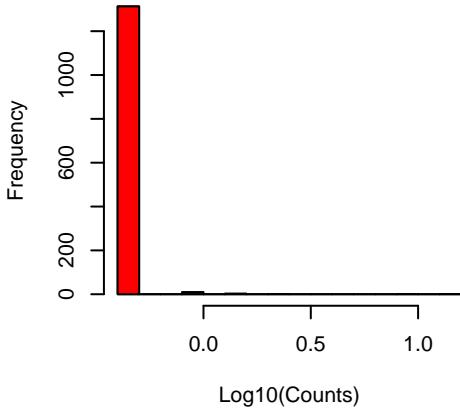


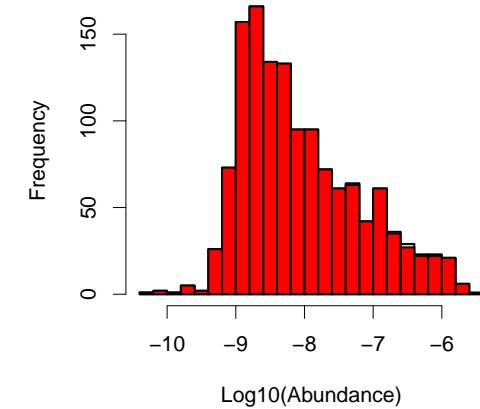
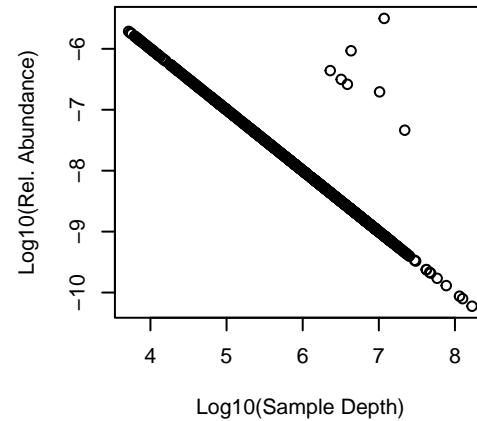
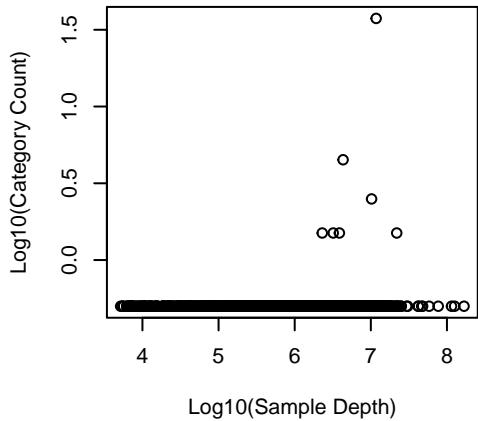
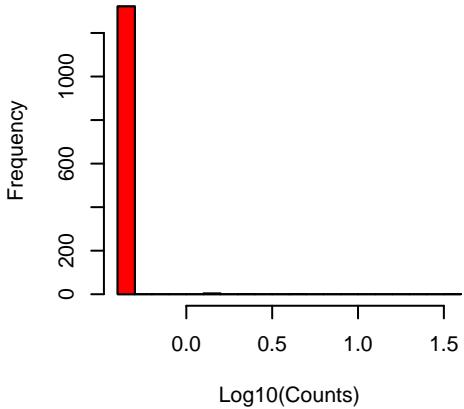


box_H ACA snoRNP complex

Zero Count/Abundance Samples: 1301 (97.89%)

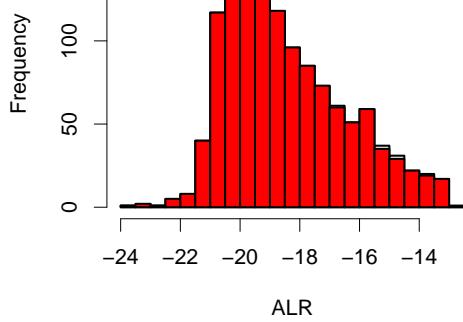
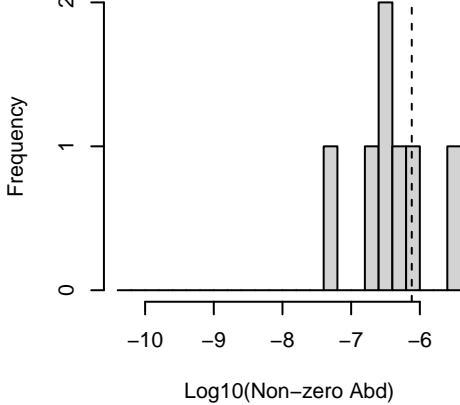


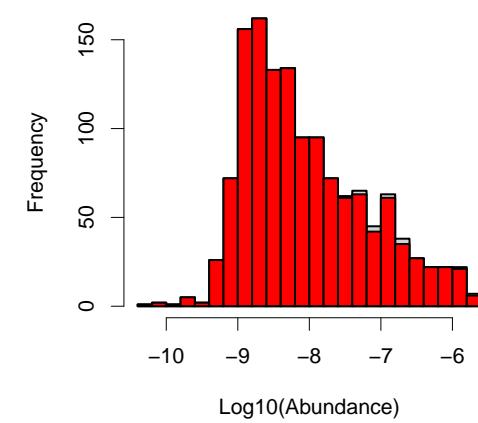
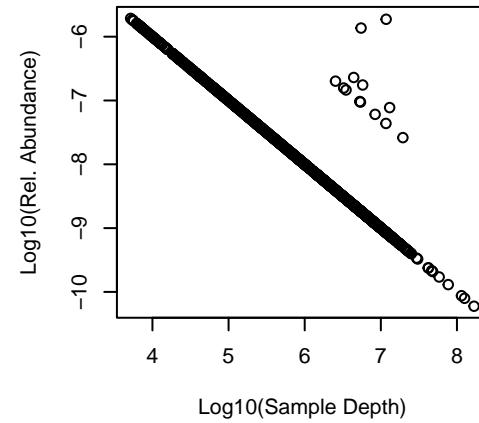
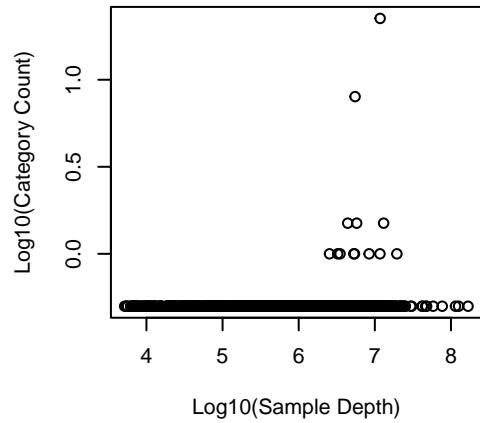
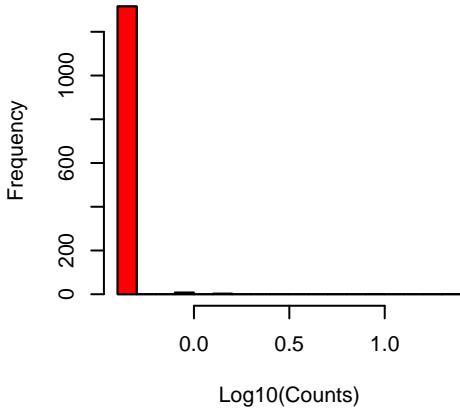




glutamate_cysteine_ligase_complex

Zero Count/Abundance Samples: 1322 (99.47%)





ubiquitin_activating_enzyme_complex

Zero Count/Abundance Samples: 1316 (99.02%)

