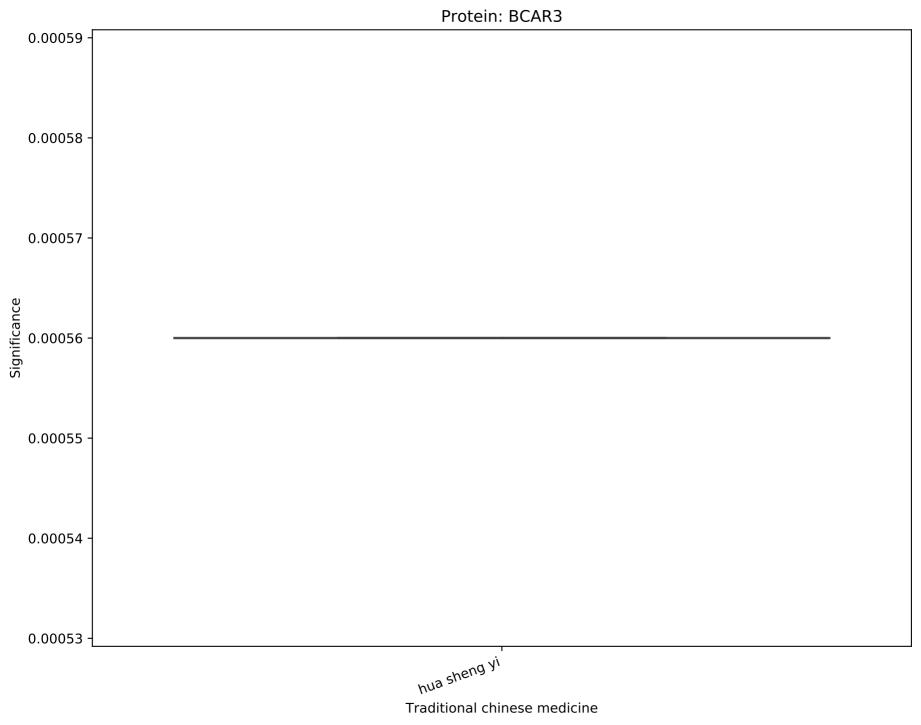
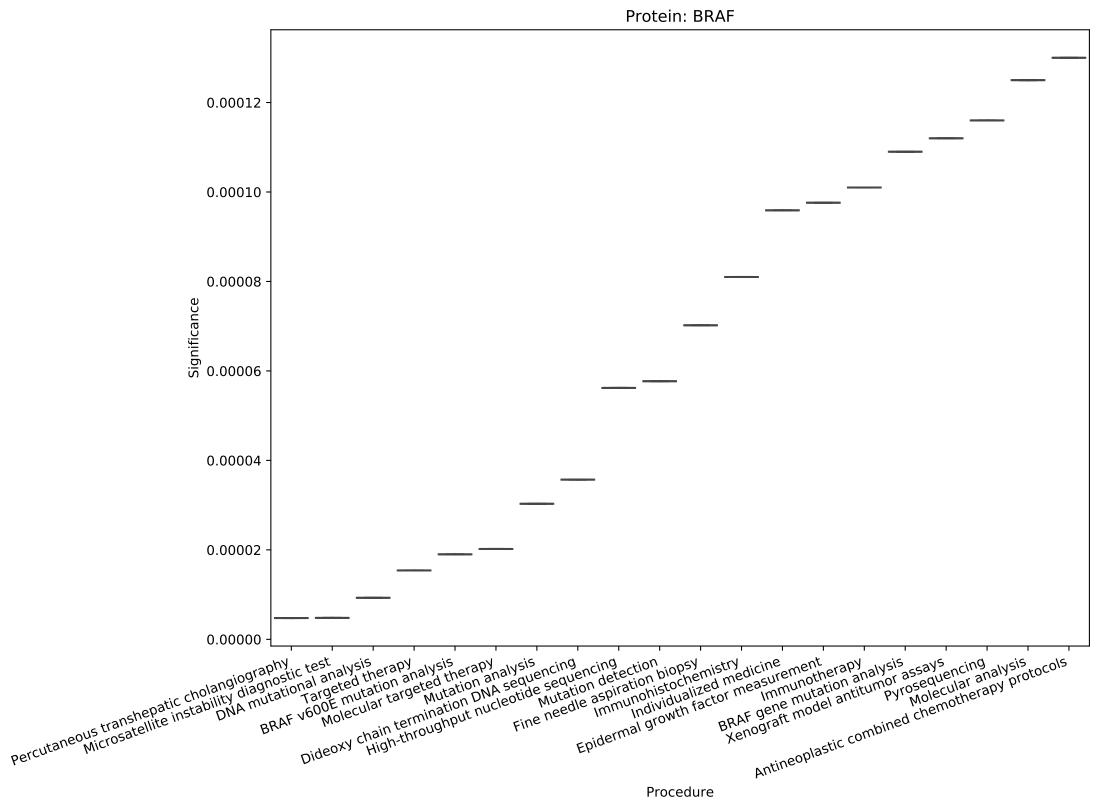
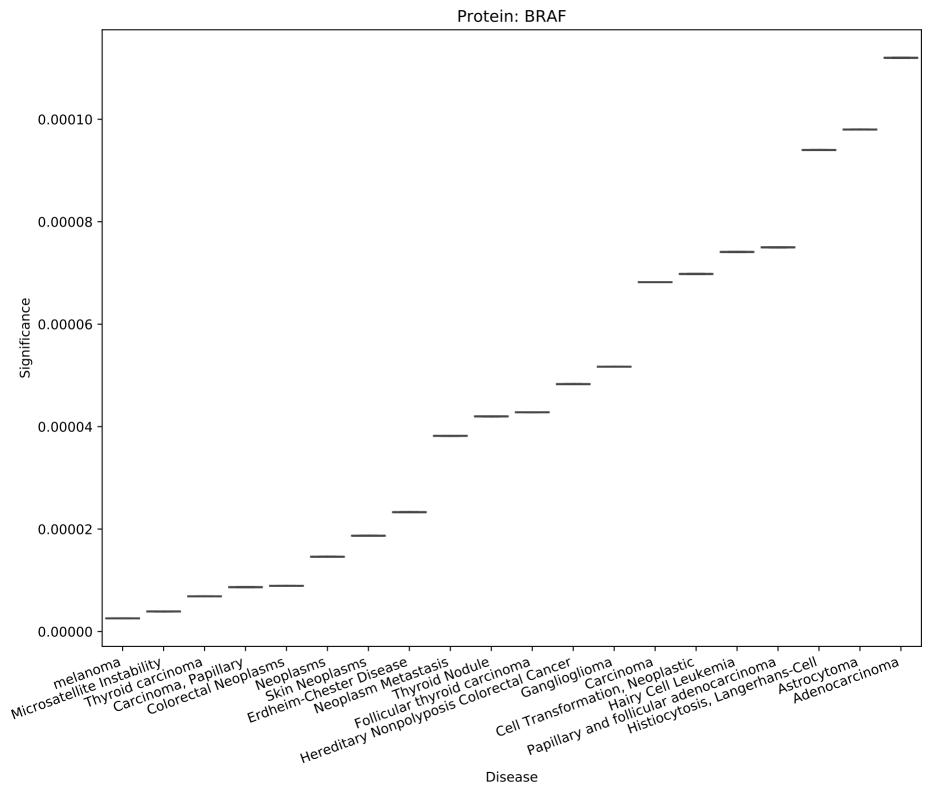
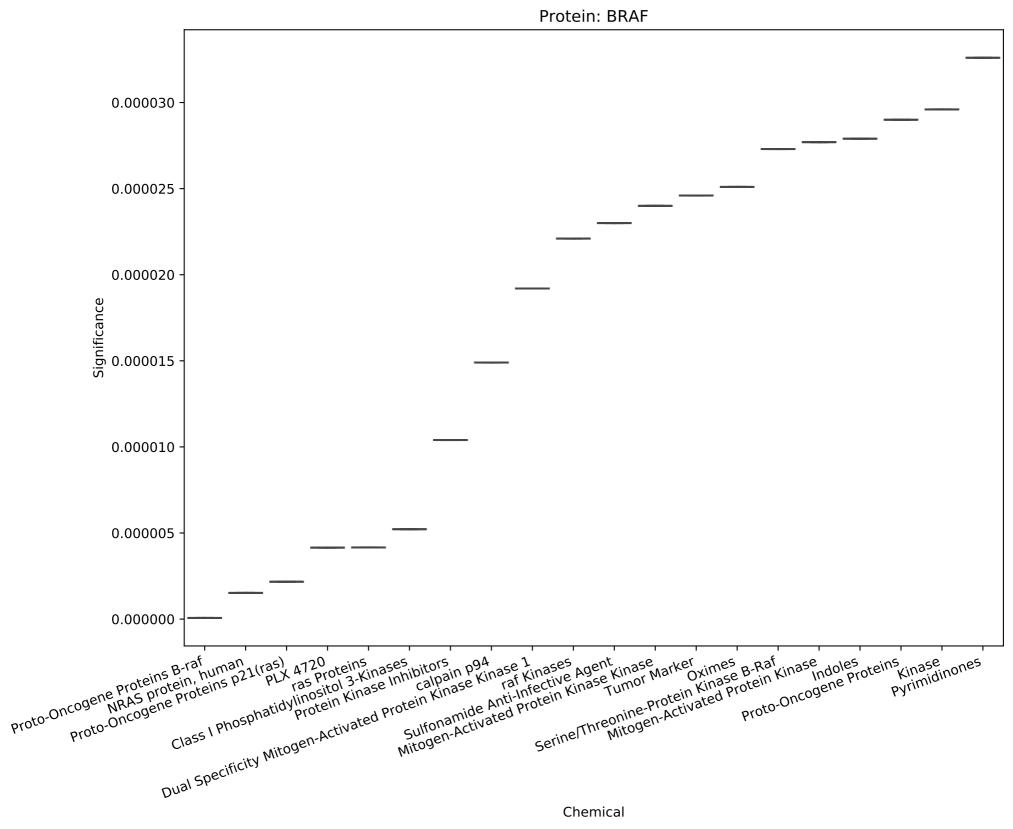


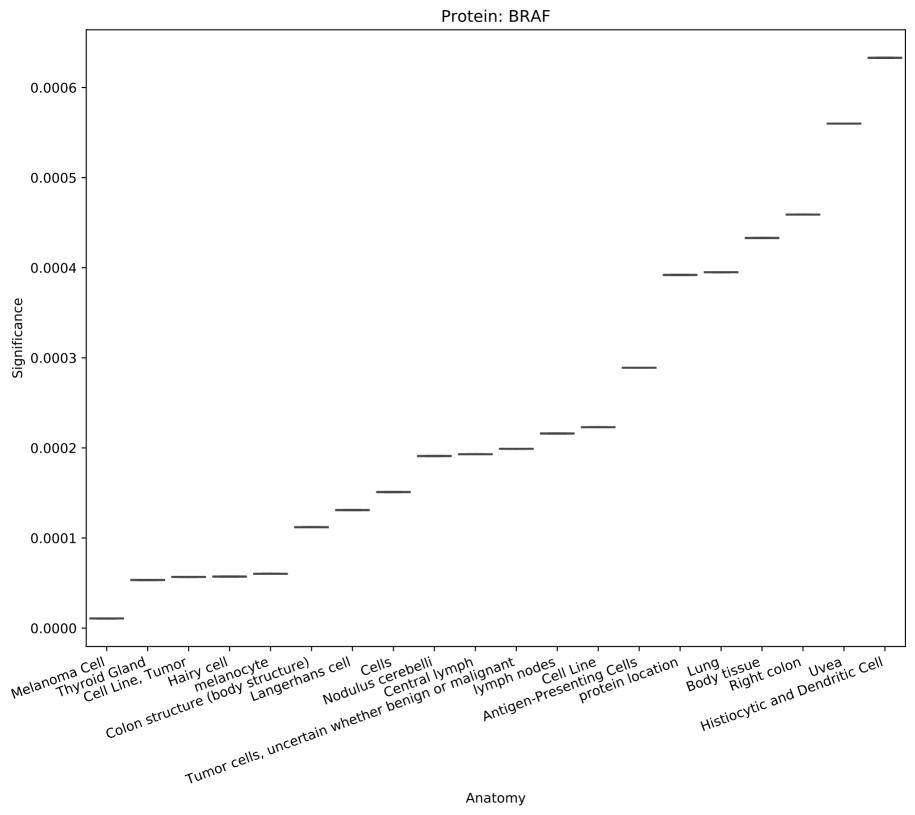
Protein: BCAR3 0.00014 0.00012 0.00010 0.00008 Significance 0.00006 0.00004 0.00002 Porcine respiratory and reproductive syndrome virus sylrus (virus virus virus

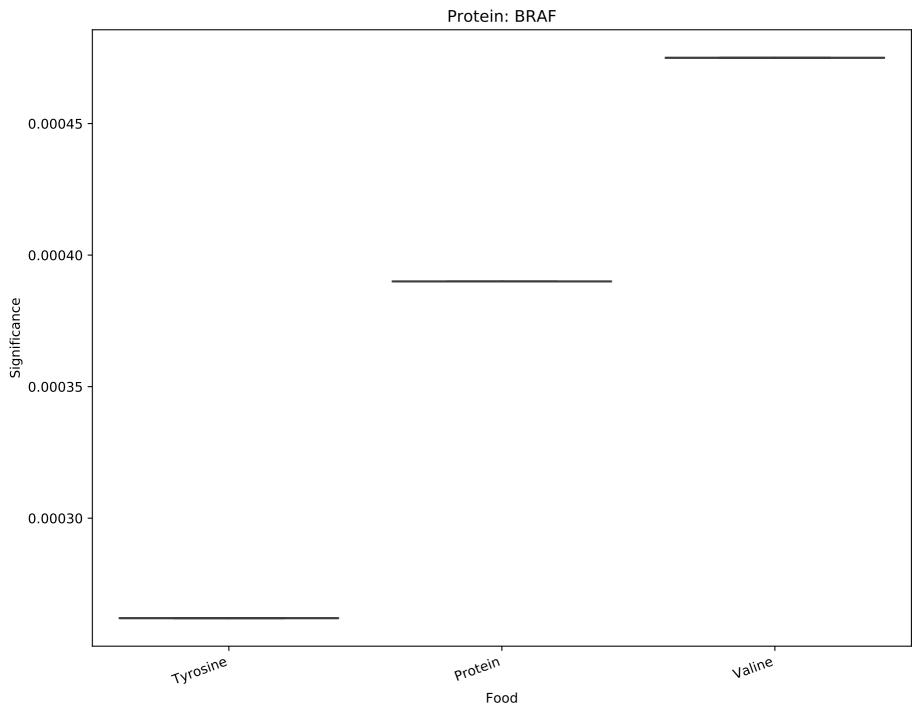


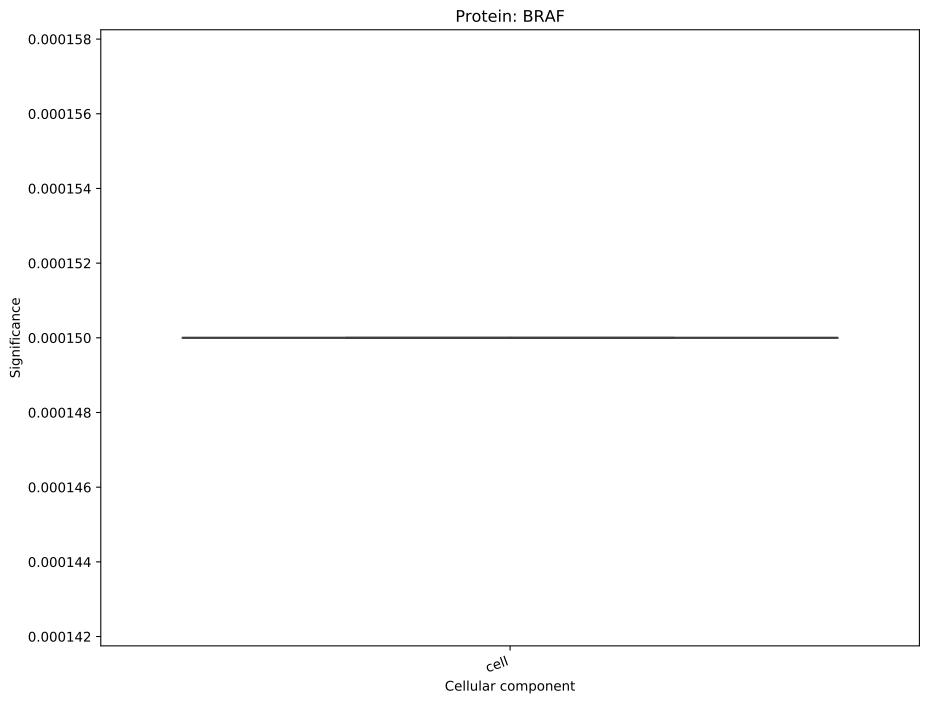






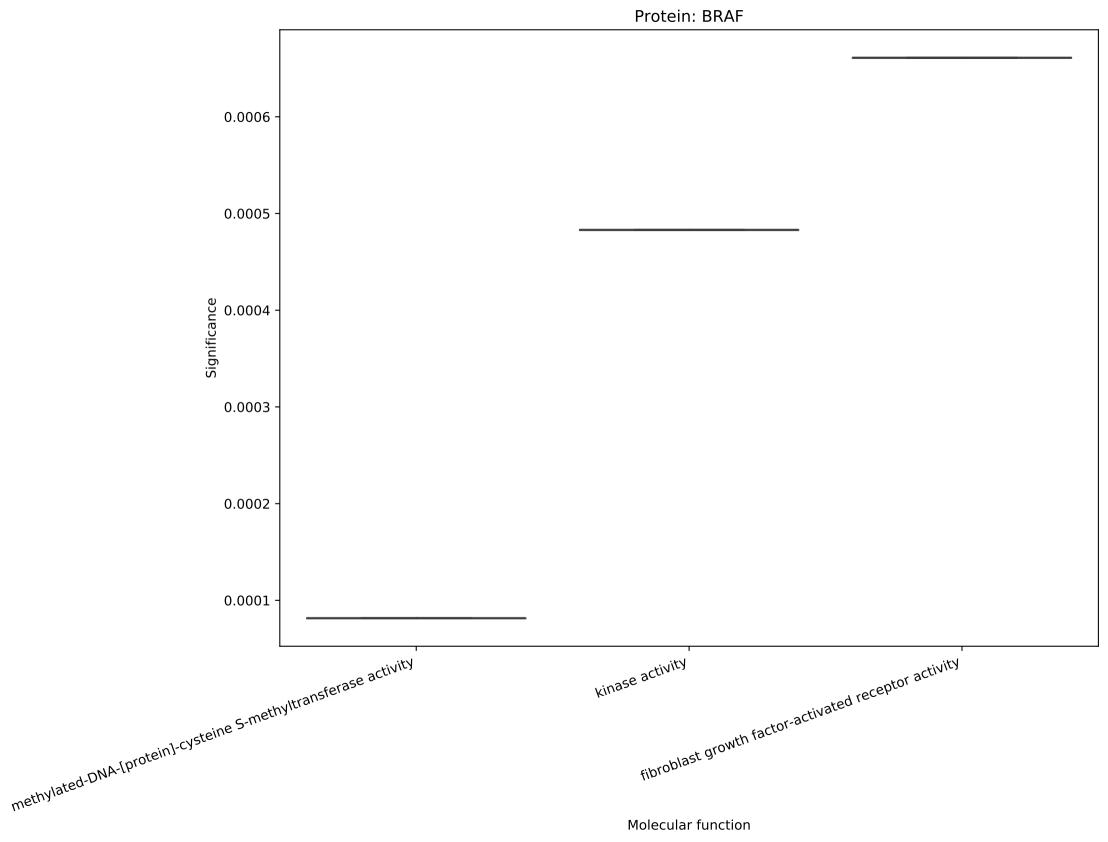


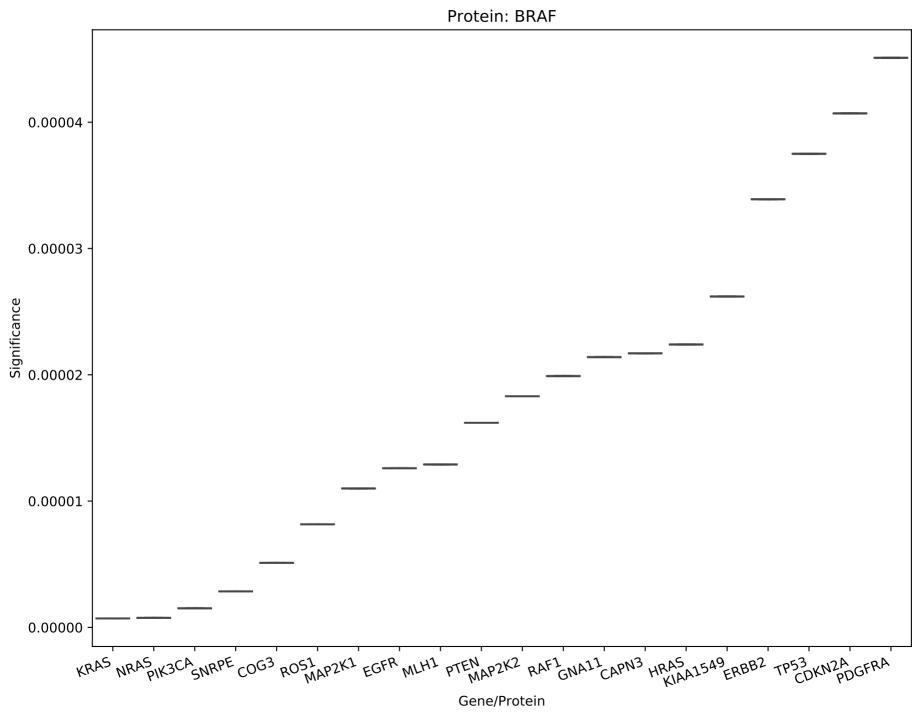


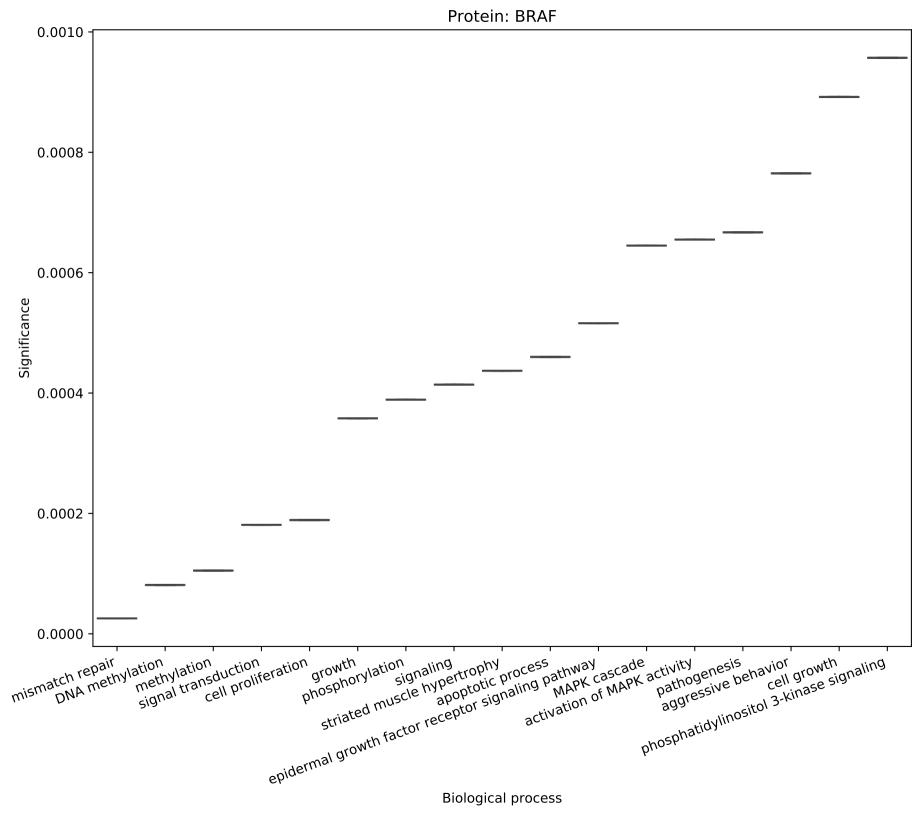


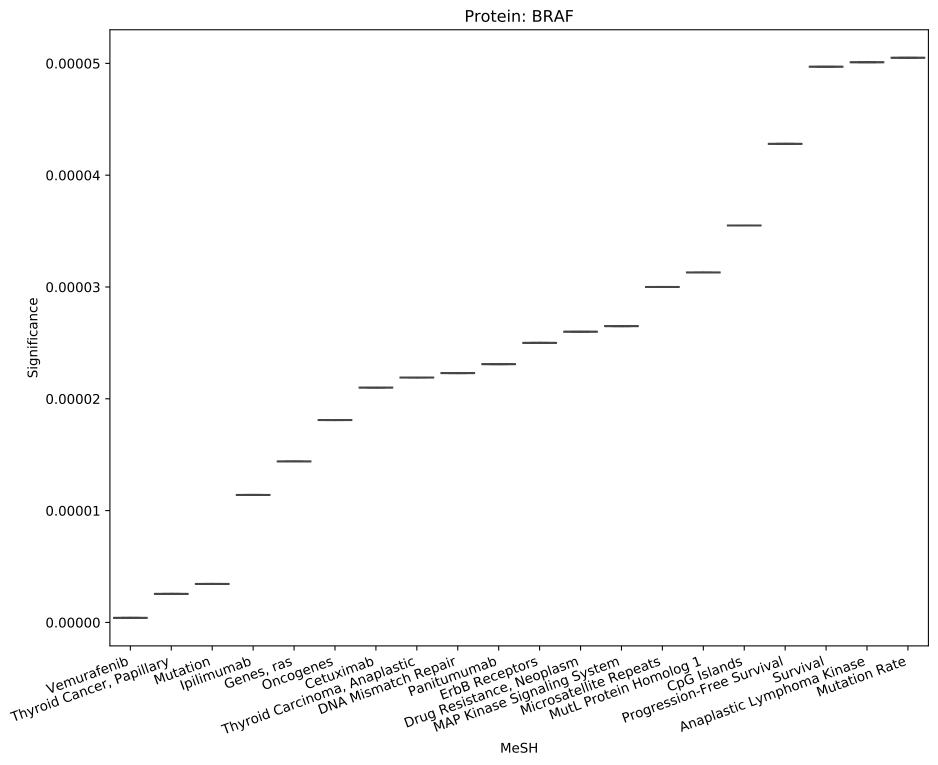
Protein: BRAF 0.000175 -0.000150 -0.000125 Significance 0.000100 -0.000075 0.000050 -0.000025 -0.000000 VEMURAFENIB TINIB TINIB

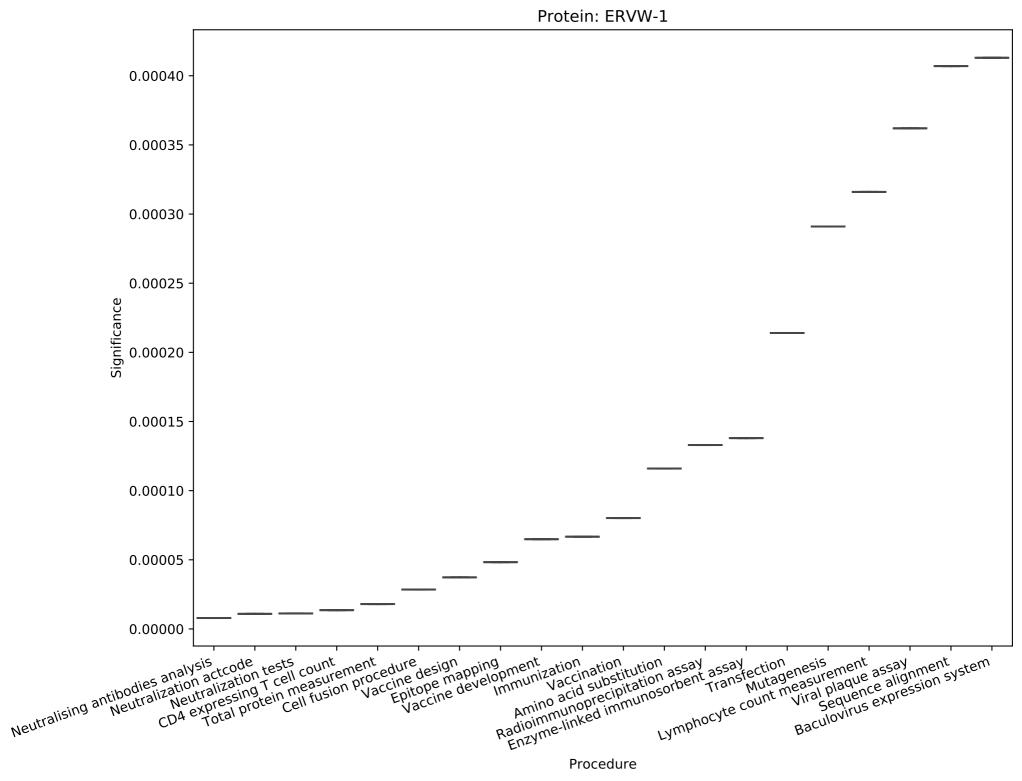
Drug

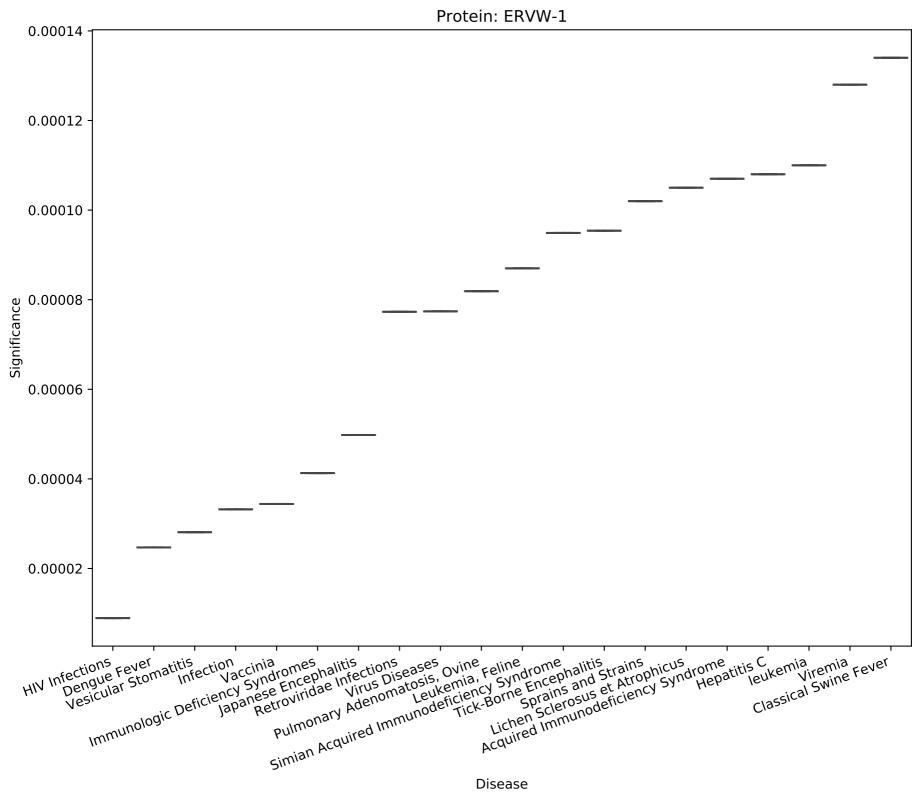




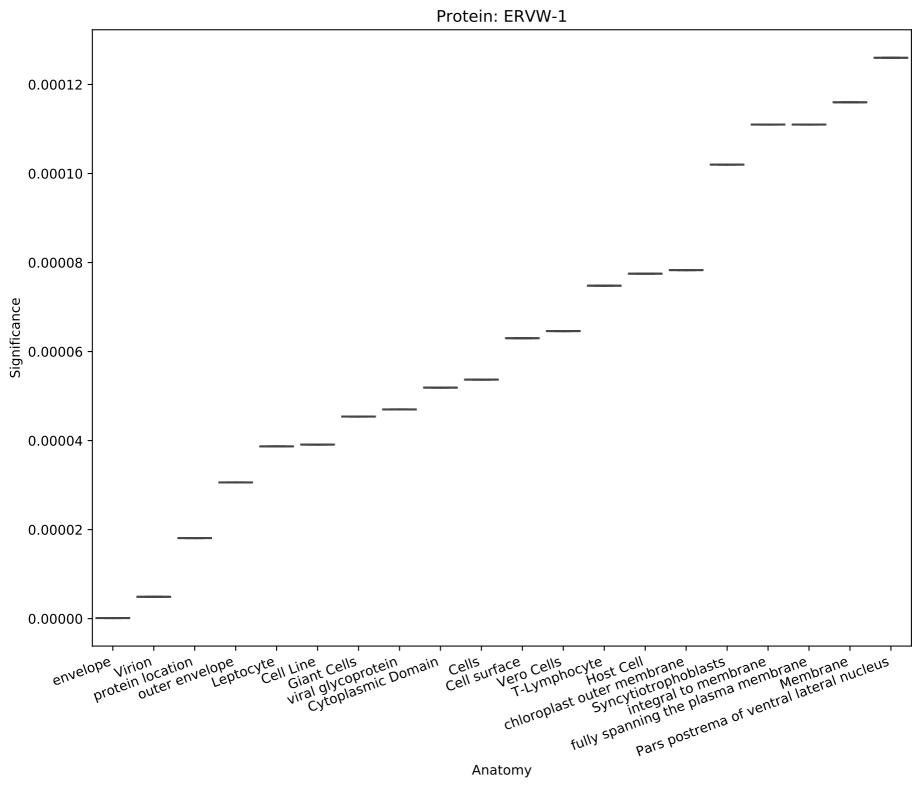


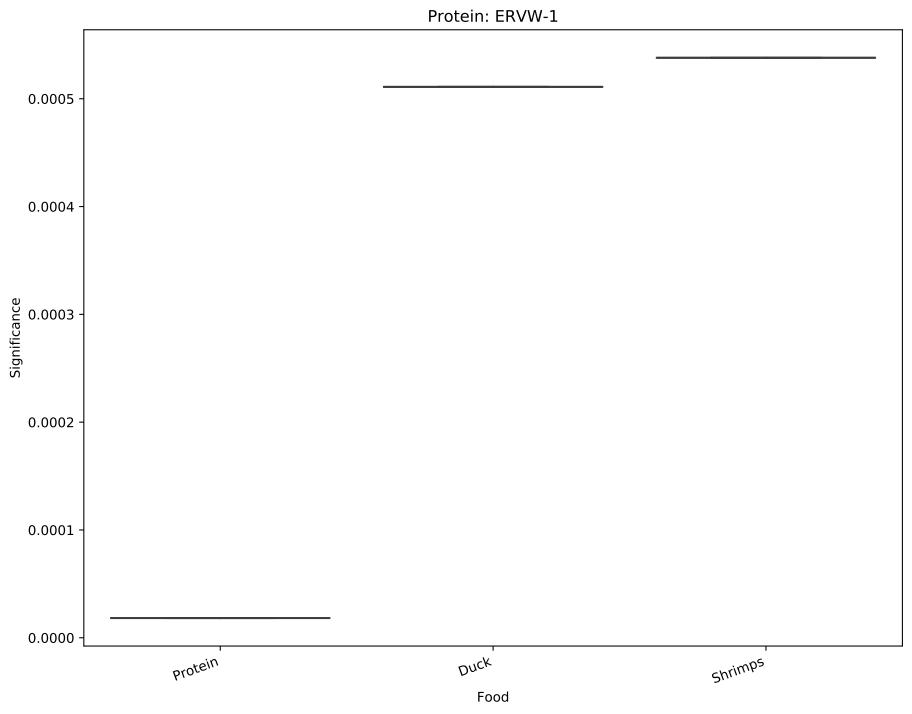


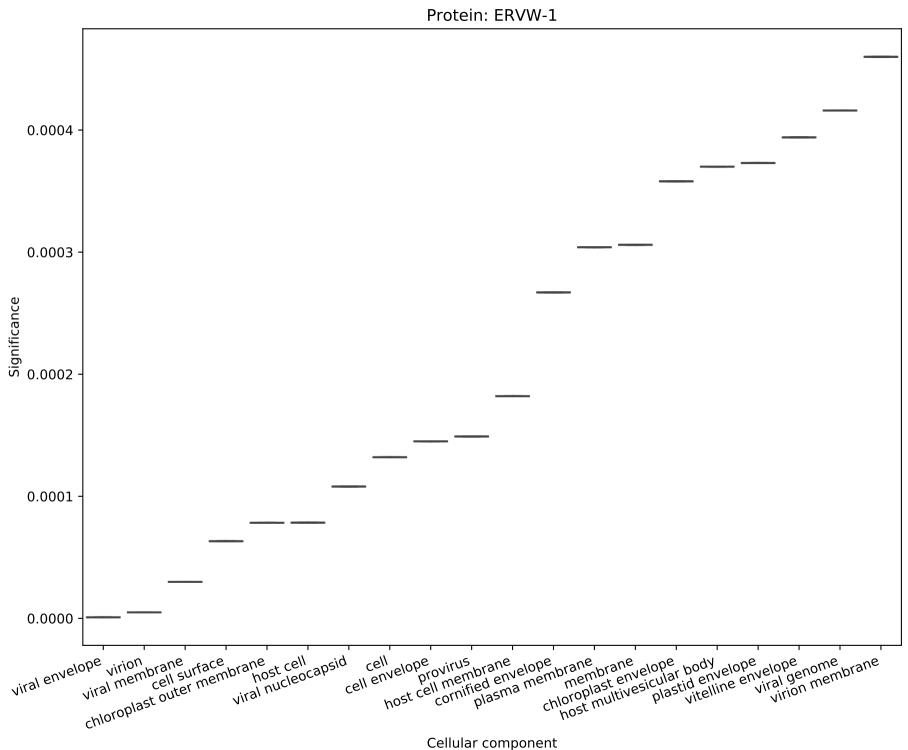




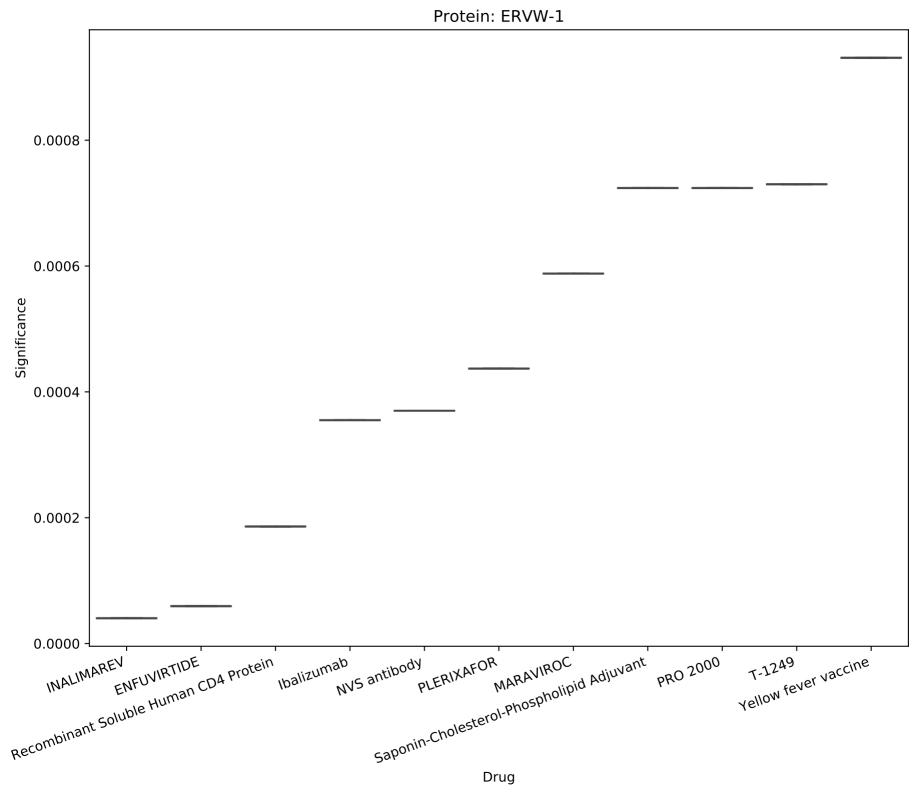
Protein: ERVW-1 0.000014 0.000012 0.000010 Significance 8000008 0.000006 0.000004 0.000002 HIV Vaccine
HIV Vaccine
FIV Vaccine
THIV Vaccine
THIV Vaccine
This type 2
This Viral Envelope Proteins gp120 Givcobri HIV Envelope Protein ENV Givcobri 0.000000 proteins gp120 protein of the syncytin gp41 gp160 virus lizing virus 1 glycoprotein gp720 protein gp720 protein gp720 protein gp720 protein gp720 protein gp720 protein gp720 protein, Human glycoprotein, Human immunodeficiency virus Er gp720 protein, Human gp720 protein, Human gp720 protein, Human gp740 envelope protein, Gramical gp740 envelope grotein, Human gp747 gp74 Antibodies virus 1 gp74 Antibodies v 'Glycoprotein Chemical

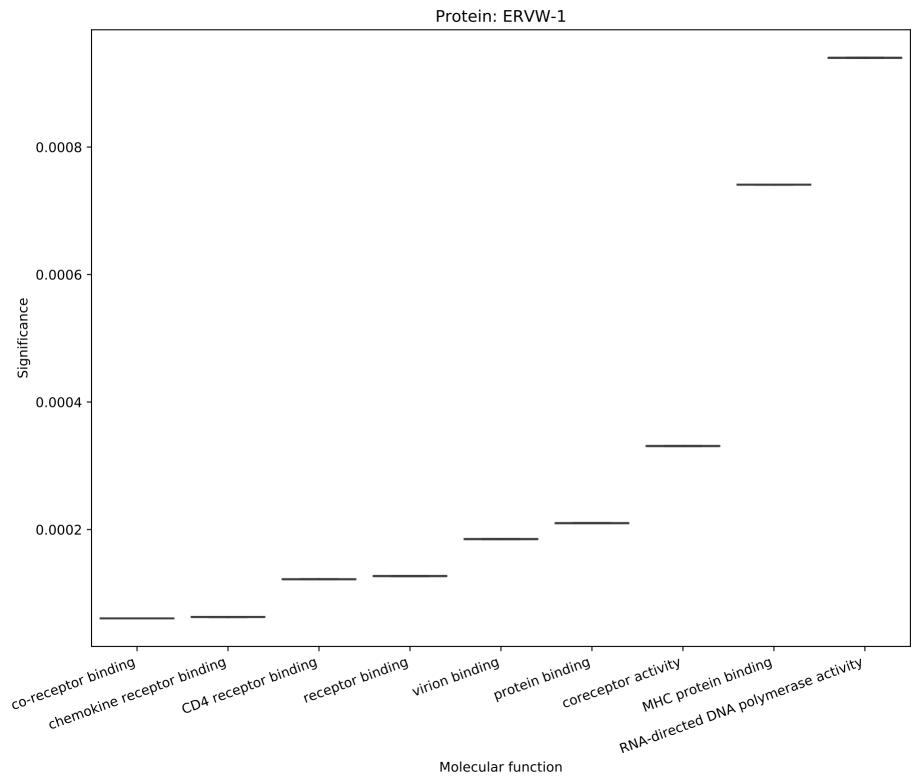


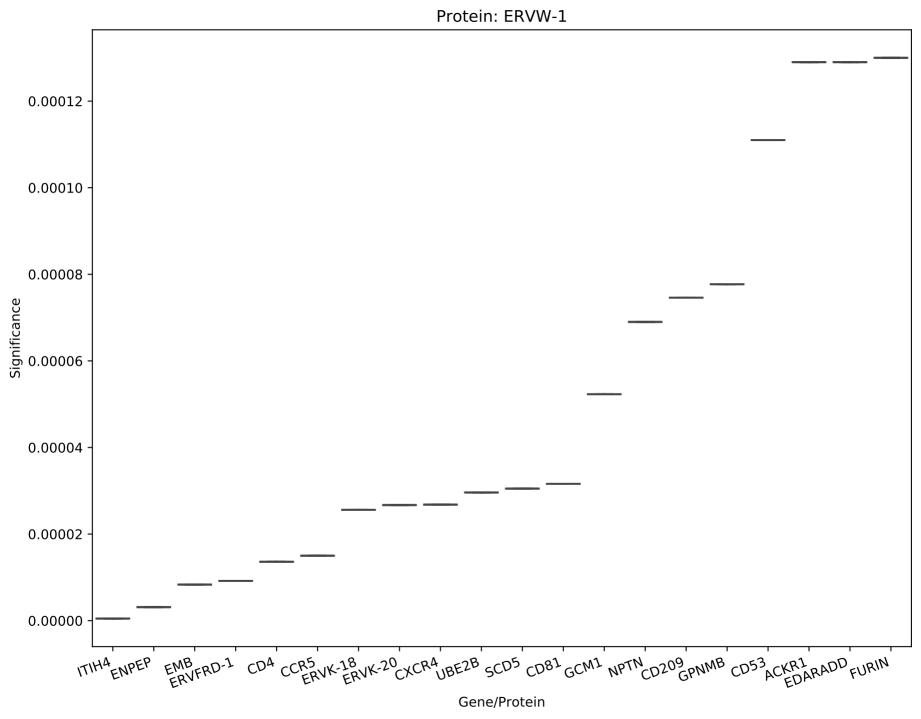


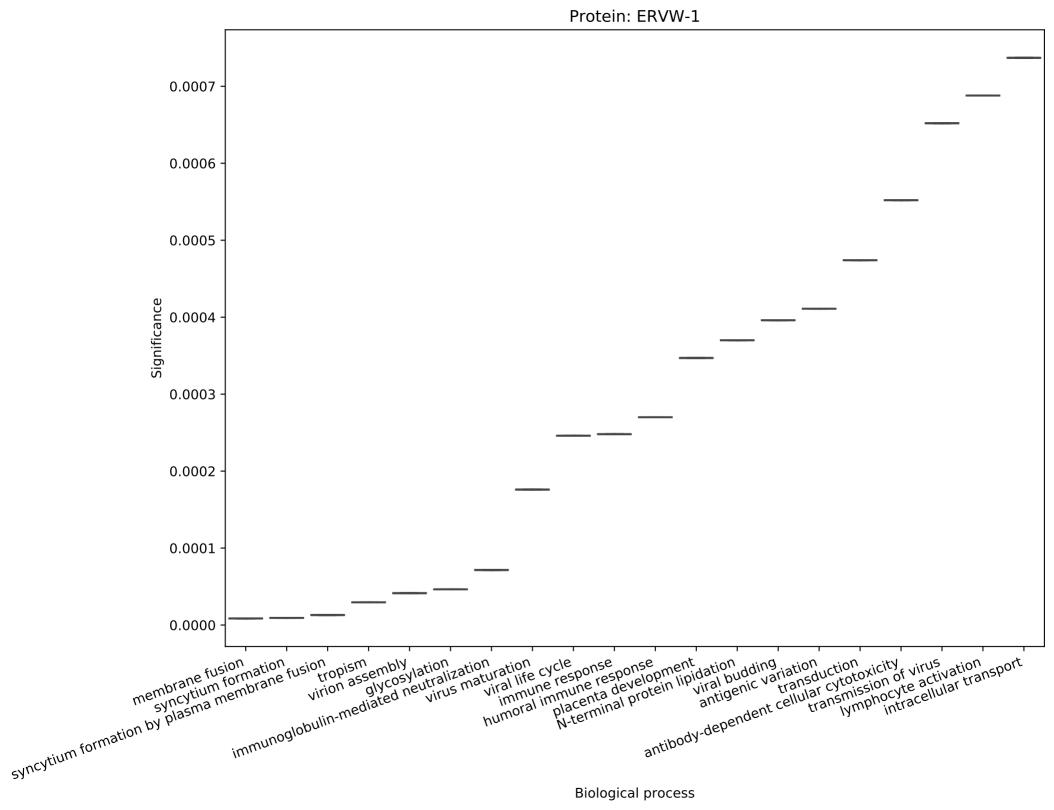


Cellular component



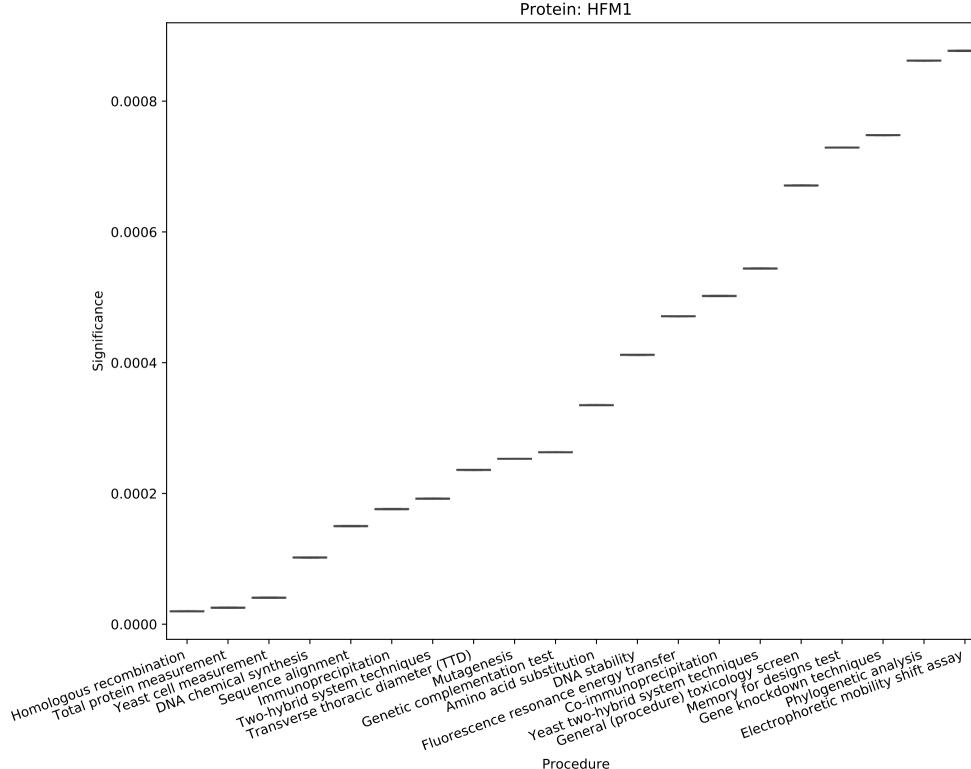




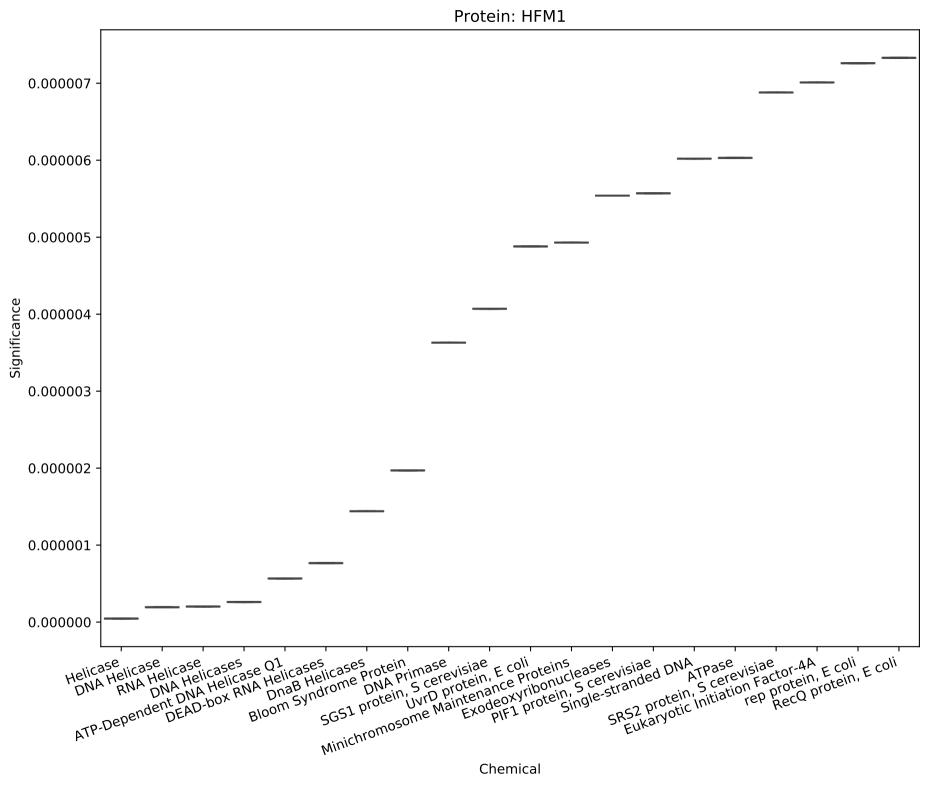


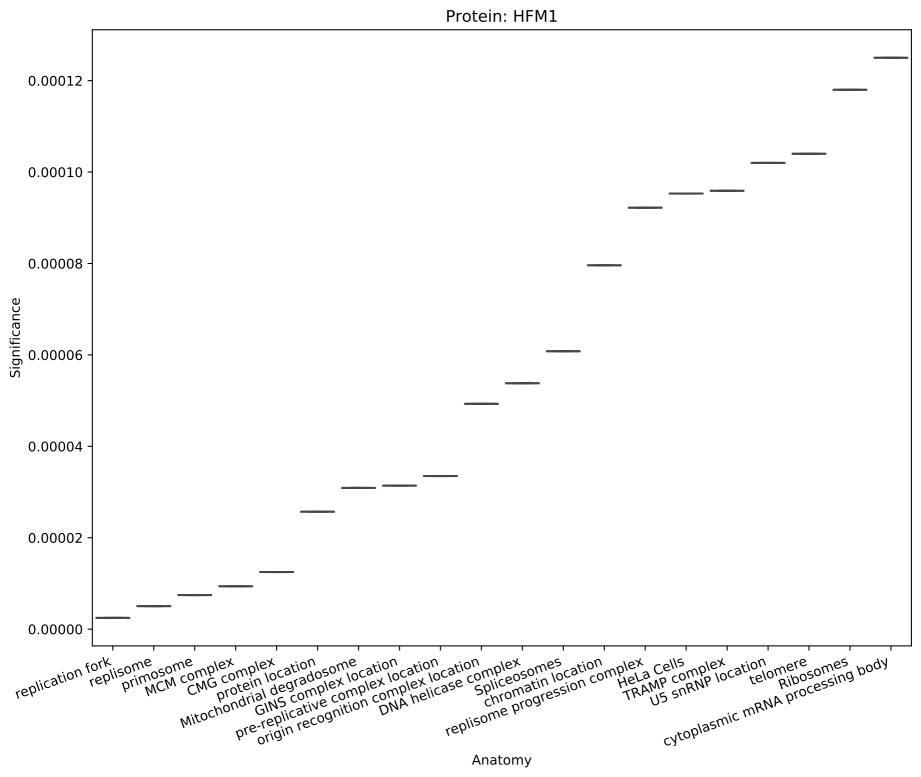
Protein: ERVW-1 0.000040 -0.000035 -0.000030 -0.000025 Significance 0.0000000 0.000015 -0.000010 -0.000005 -HIV-1 Viruses, env Genes, env Gen

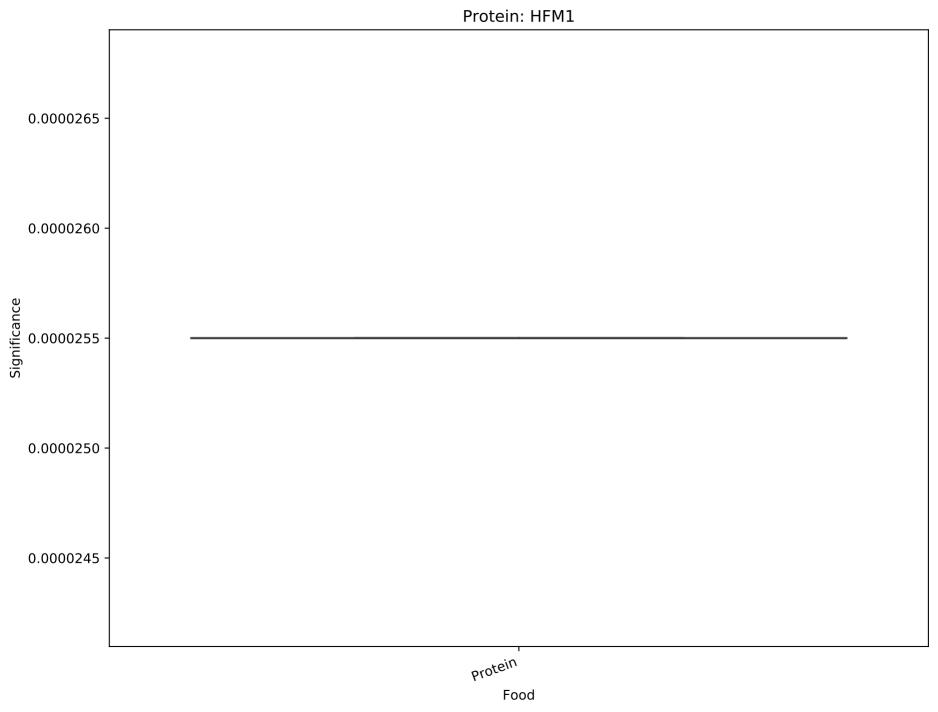
MeSH

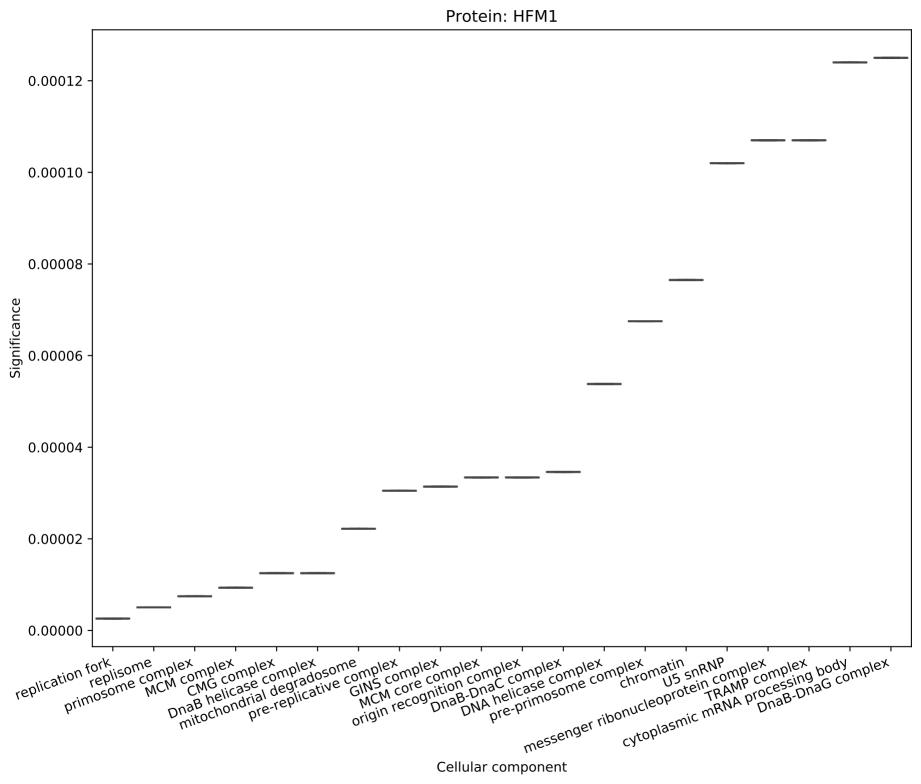


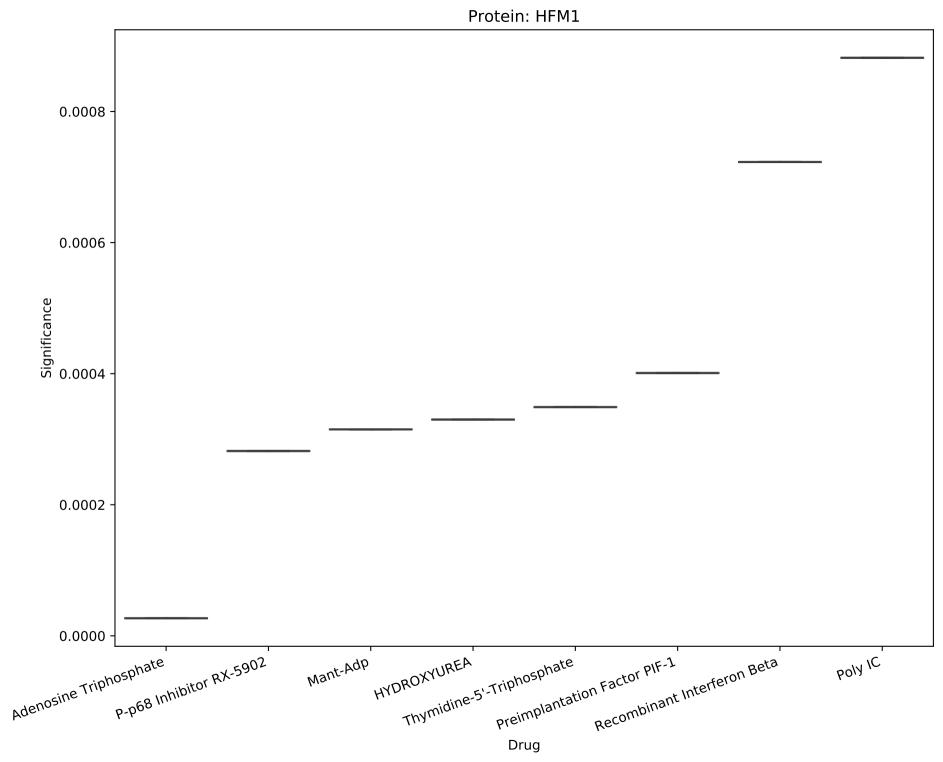
Protein: HFM1 0.0007 -0.0006 0.0005 Significance 0.0004 0.0003 0.0002 0.0001 Chronic progressive external Oculovestibuloauditory chronic progressiv 0.0000 r Syndrome r Syndrome Syndrom Chromosome Virus Diseases Merner Syndrome

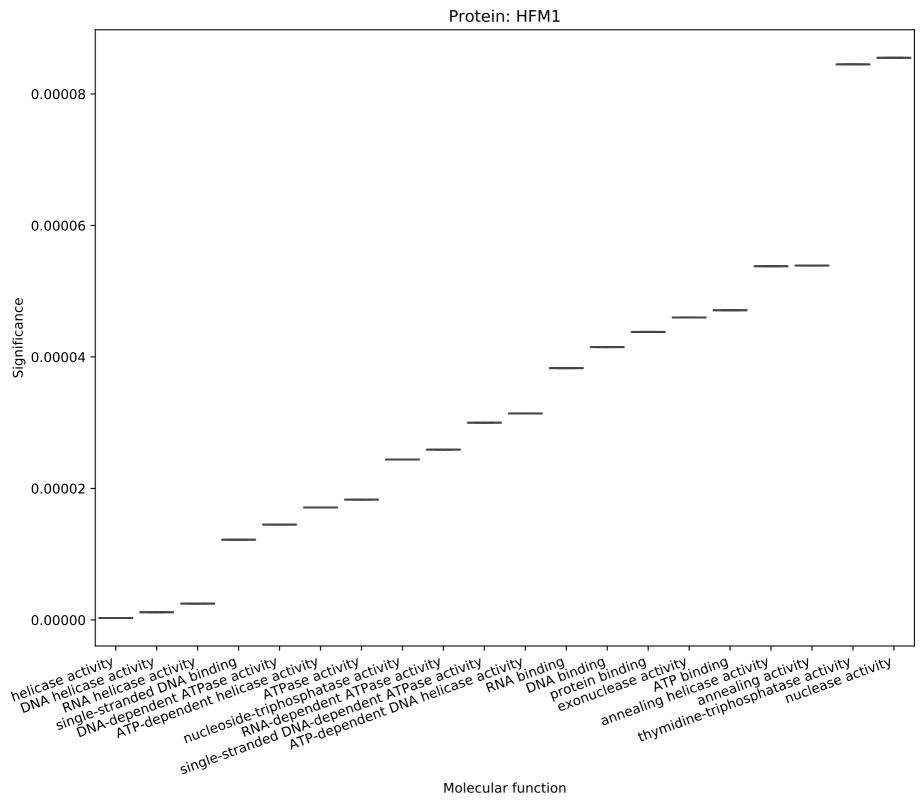


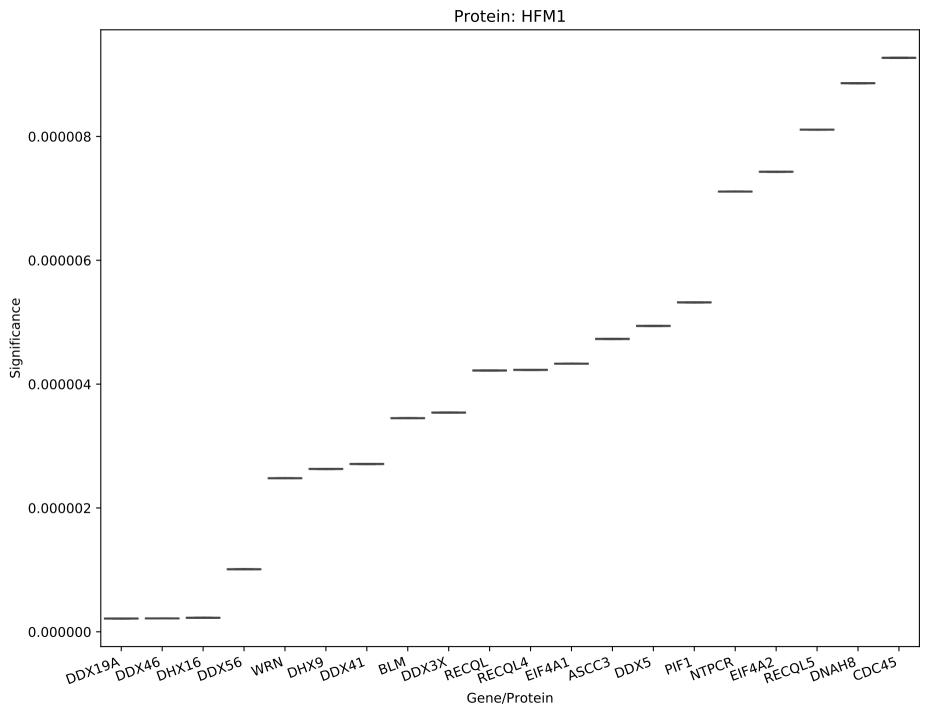








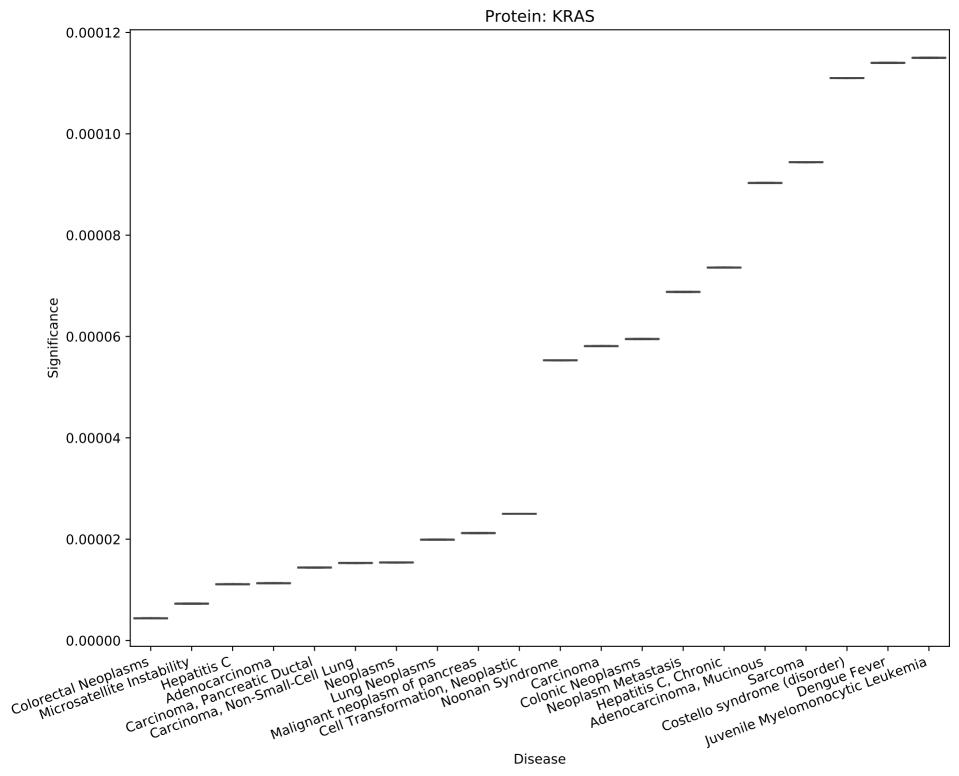


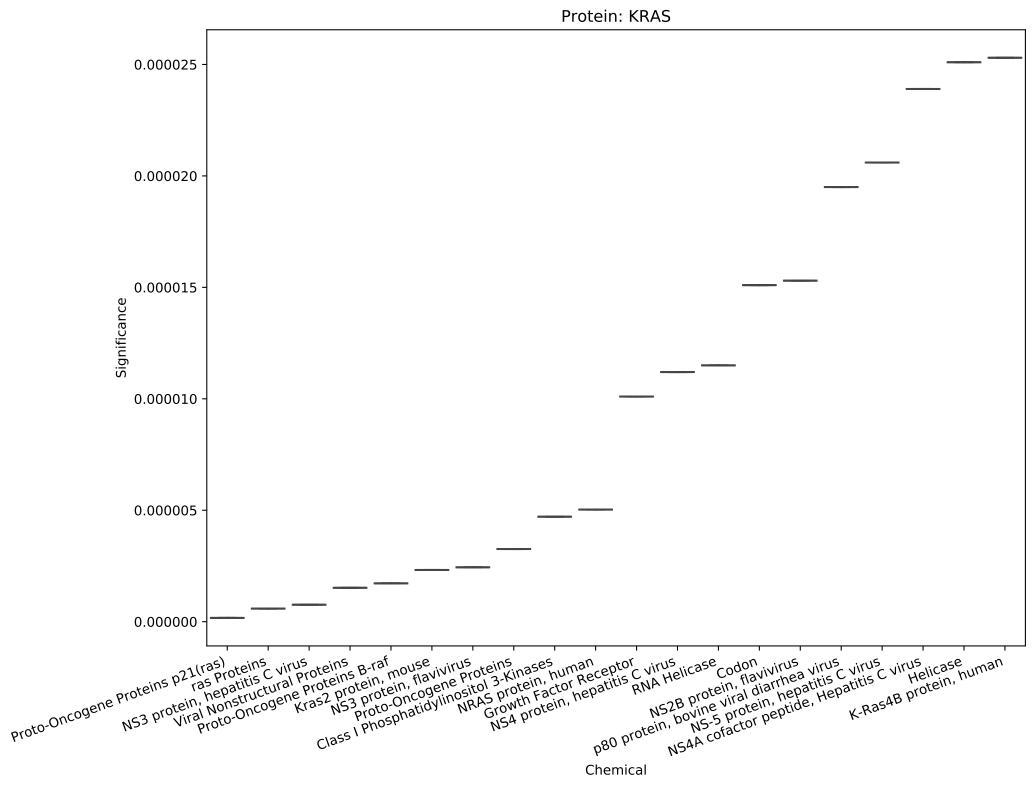


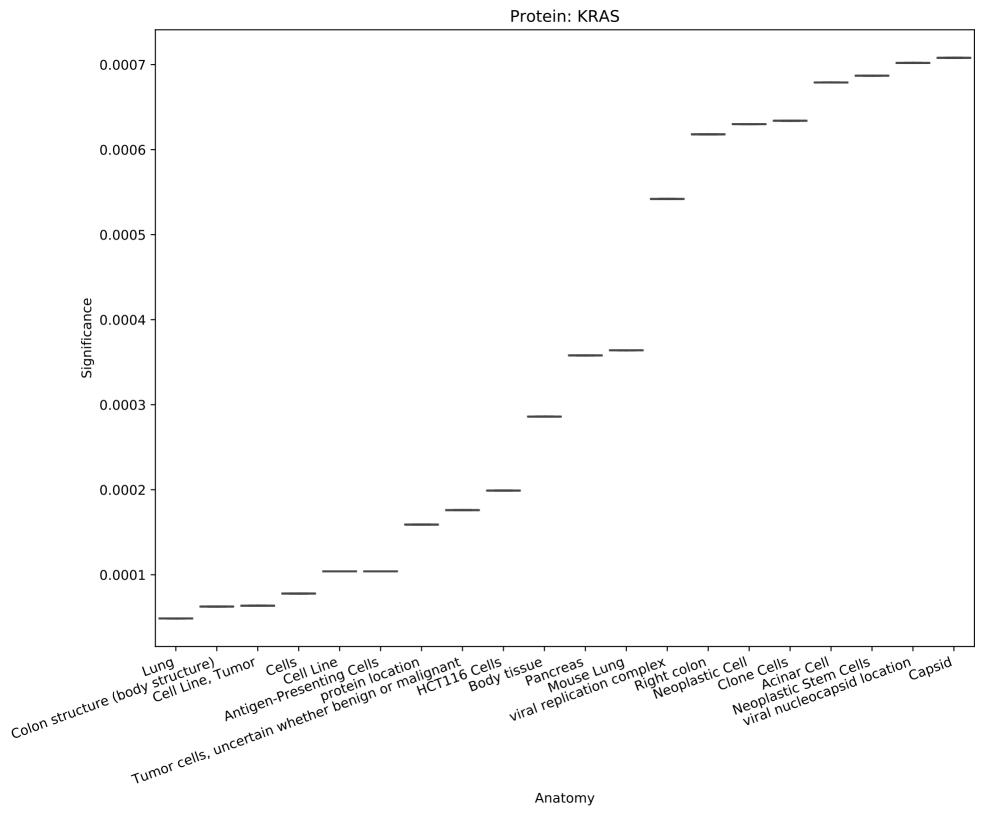
Protein: HFM1 0.00008 0.00006 -Significance 0.00004 0.00002 0.00000 DNA duplex unwinding cation repair DNA replic pro Ading ation repair ocess ocess arrest ation along ation repair ment ation repair ment ation notic process ocess arrest ation and initiation repair ment ation of the polic process ocess arrest ation and initiation repair ment ation nests and eacaly initiation of the polic process ocess. The process ocess are strained initiation of the polic process ocess. The policy of the policy ocess ocess are strained initiation repair ment ation nests and eacaly initiation repair ment ation nests and eacaly initiation repair ment ation nests and eacaly initiation and initiation repair ment ation nests and eacaly initiation repair ment ation at the pair at the pair ation at th **Biological process**

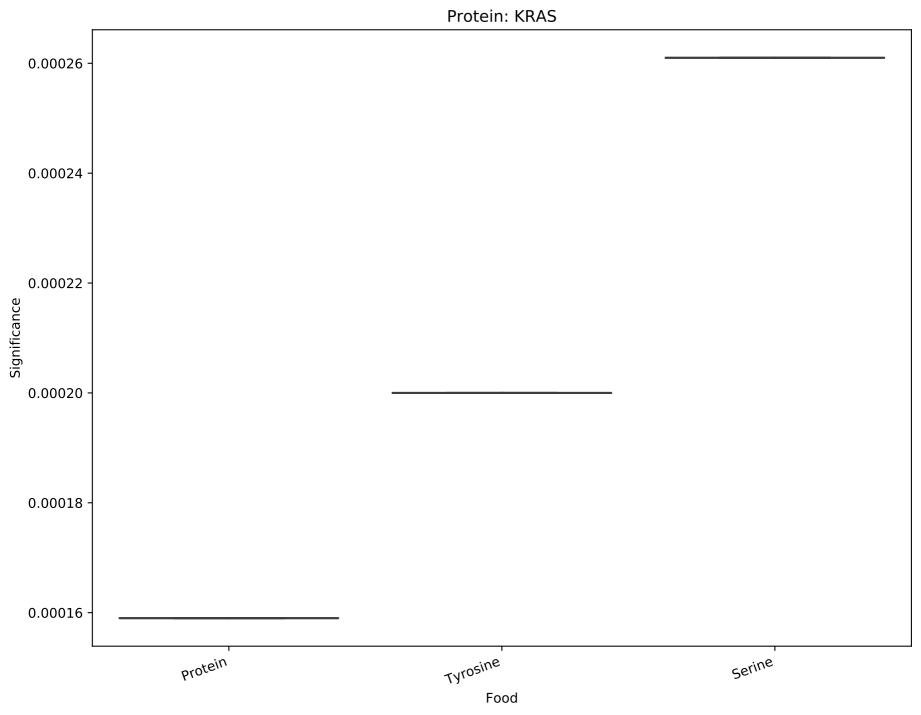
Protein: HFM1 0.00005 0.00004 Significance 0.00003 0.00002 0.00001 0.00000 Werner Syndrome Helicase Proteins Origin Ori nent 6 Genome Mutation DNA Dam Mutation Mutation and Data Strander Data Molecular Sequence Bacterional DNA RNA Viruses Bacterional DNA RNA Viruses Bacterional DNA RNA Viruses Recombinational DNA RNA Viruses MeSH

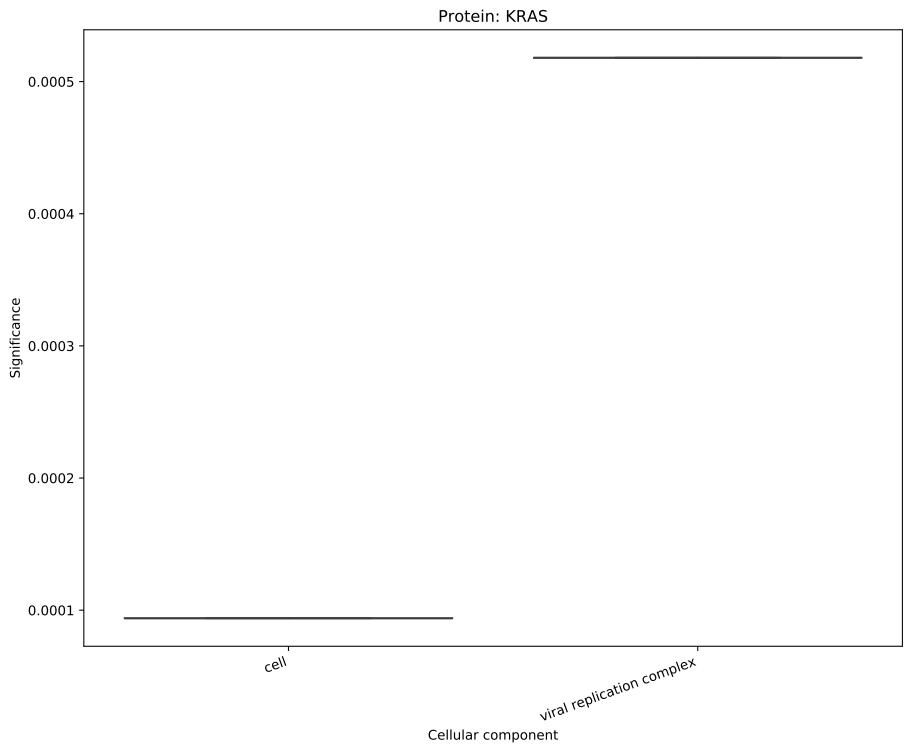
Protein: KRAS 0.00014 -0.00012 0.00010 Significance 0.00008 0.00006 0.00004 0.00002 Kras mutation analysis testing test analysis ment diagnostic analysis ment analysis measurement regimen regimen regimen measurement mutation determents mutation determents mutation determined in the management of the mutation determined in the mutation determined ng test lysis ment men clion cing cing rapy lysis men regimen regimen regimen regimen regimen detection as analysis men regimen regimen detection are sequenced there analysis men respondenced the rechnique chain reaction cols described the rechnique chain reaction cols described and respondenced the respondence to the respondenced the respondence to the respond Procedure

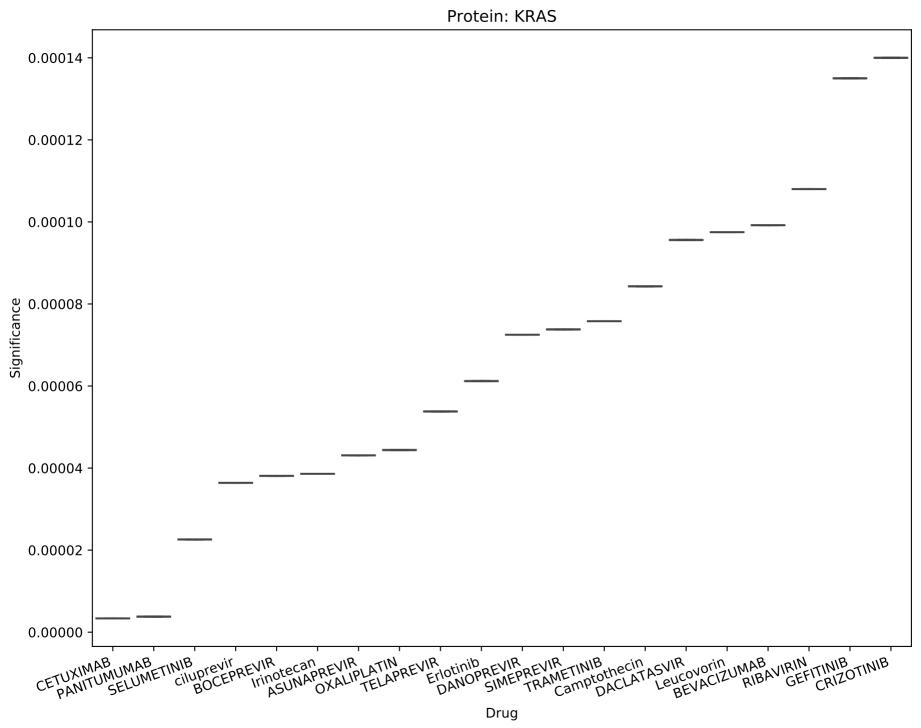


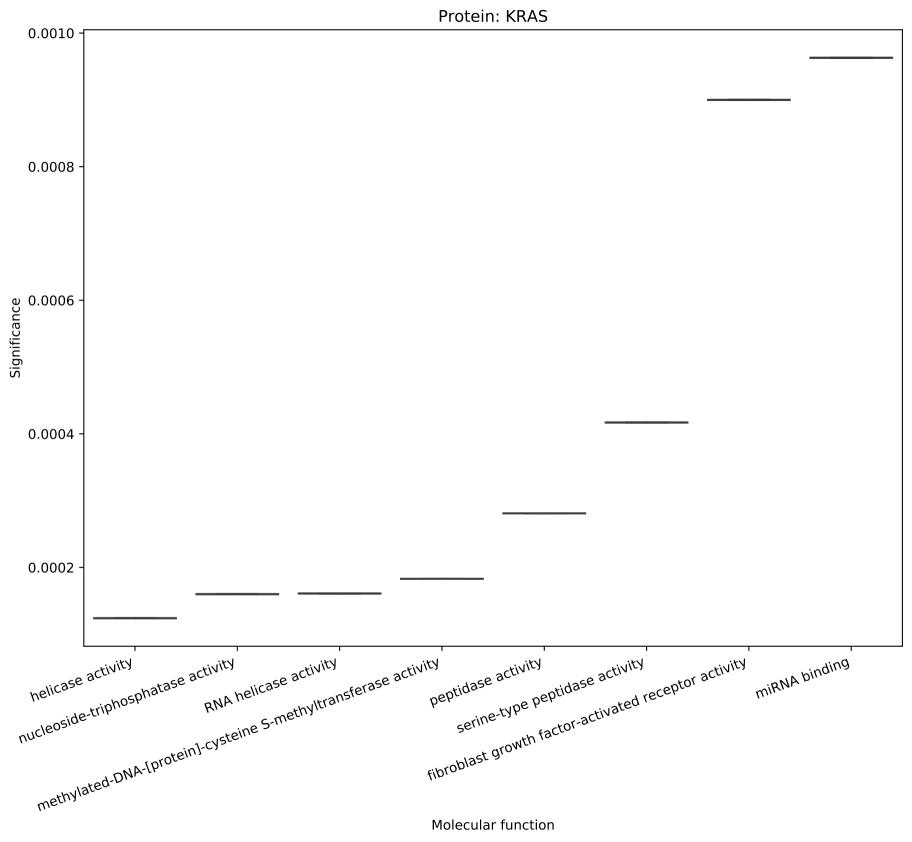


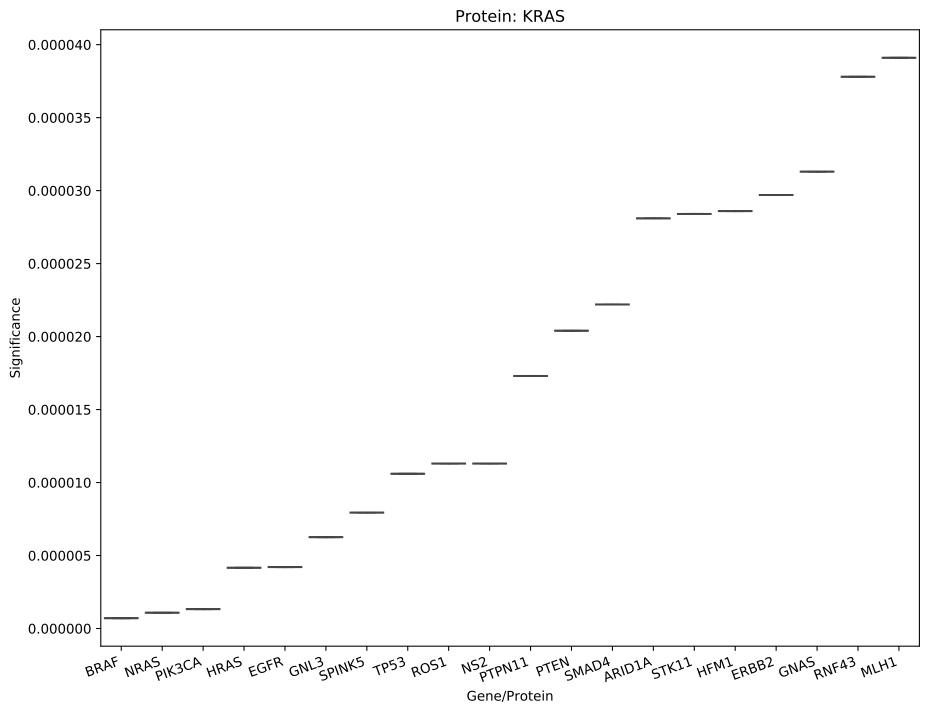


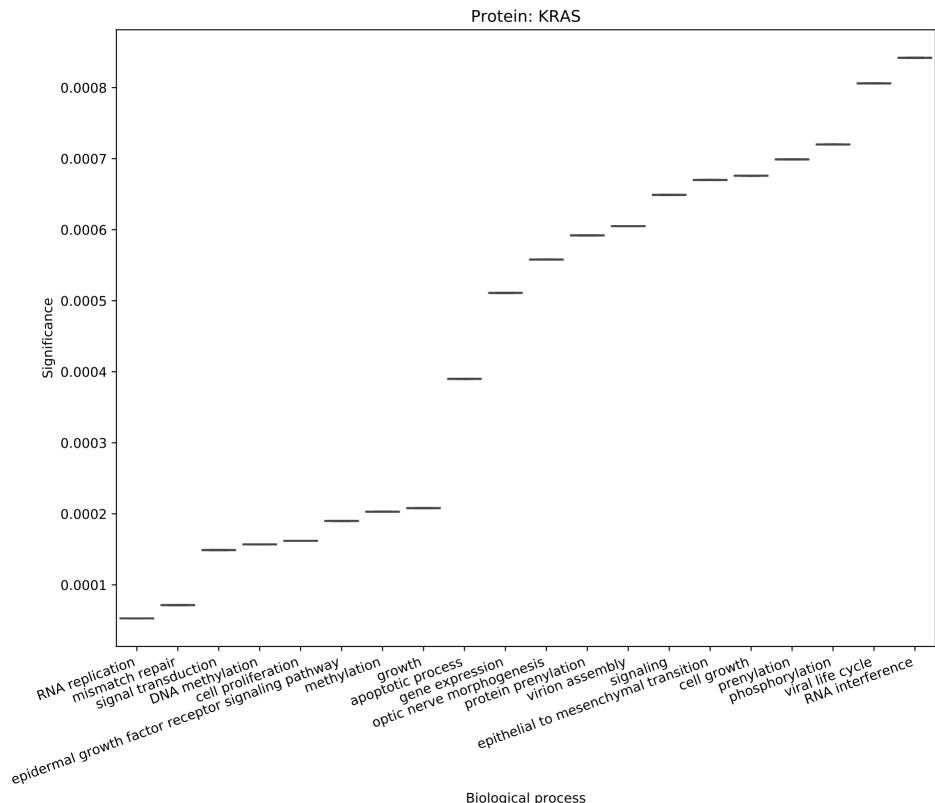




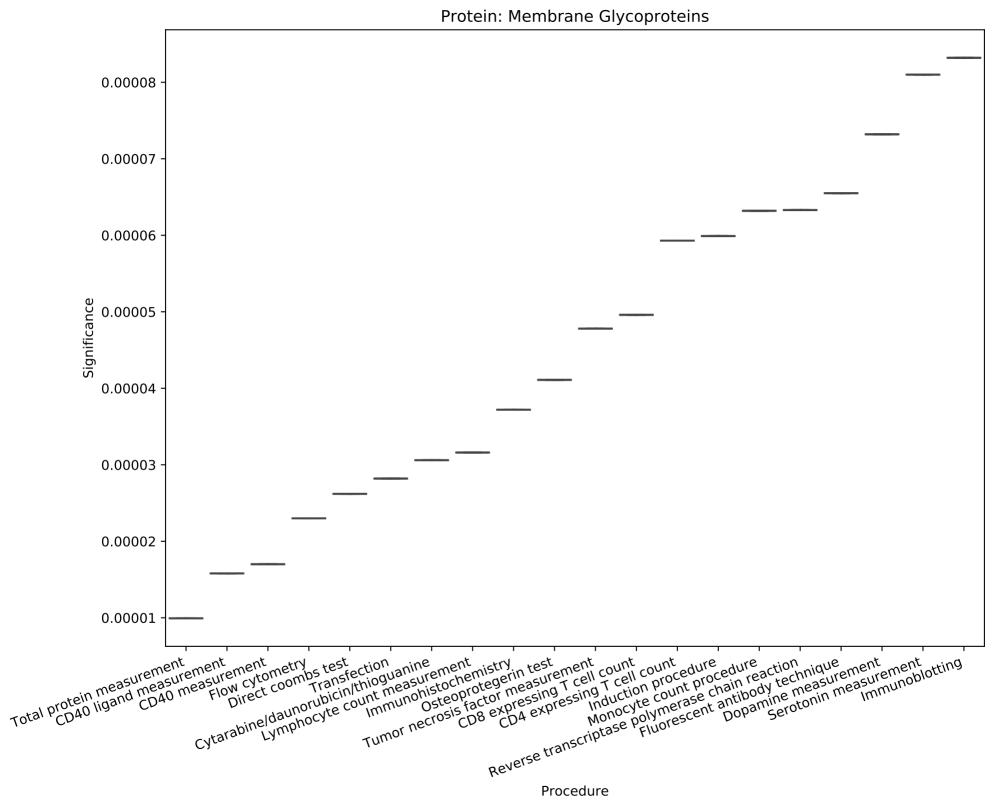


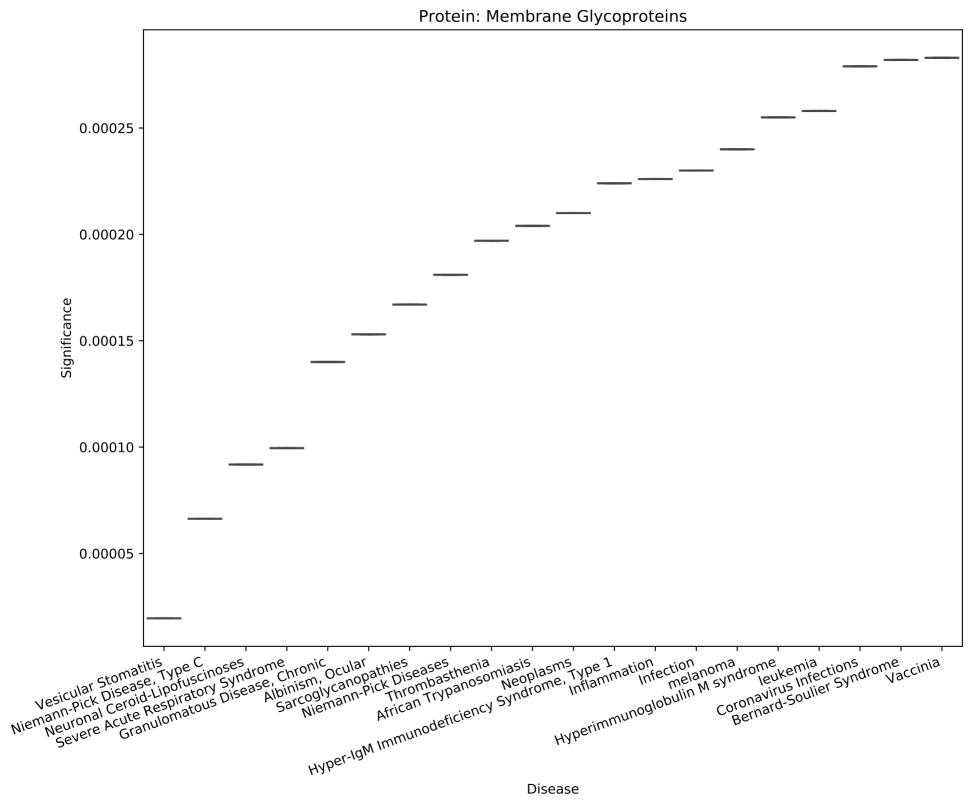


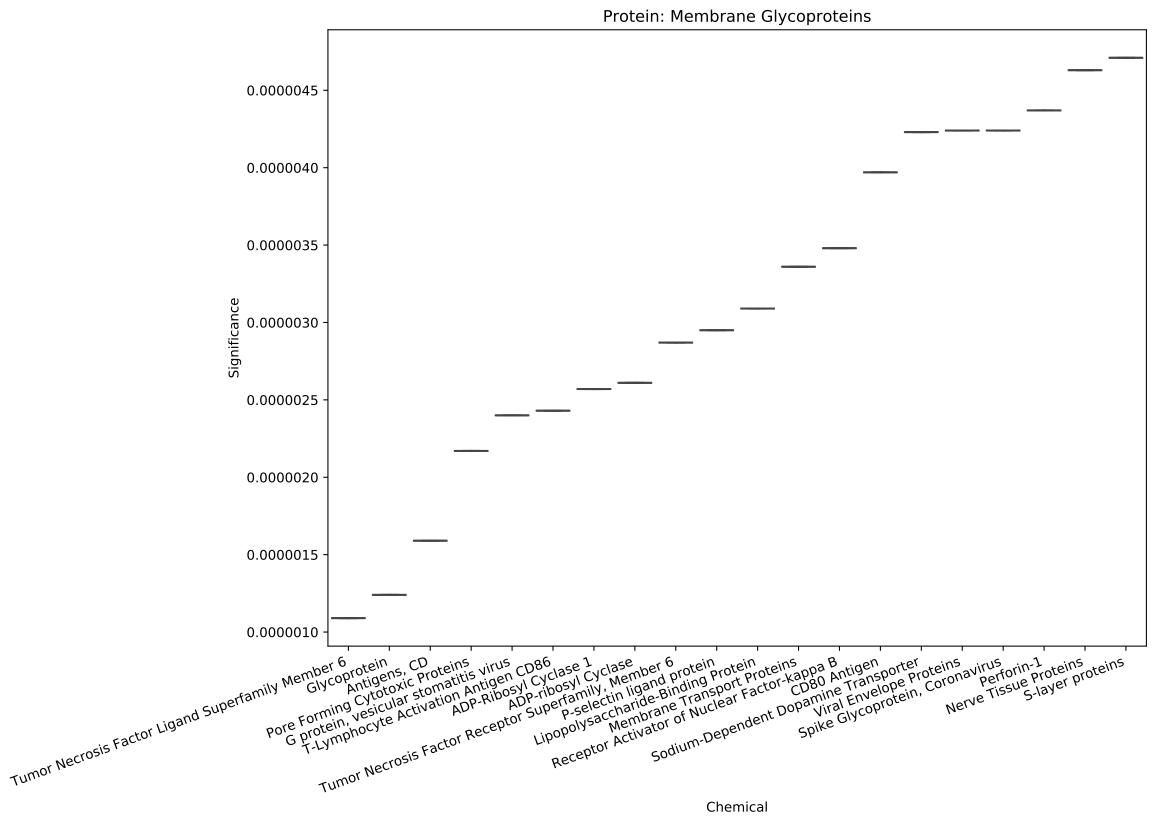


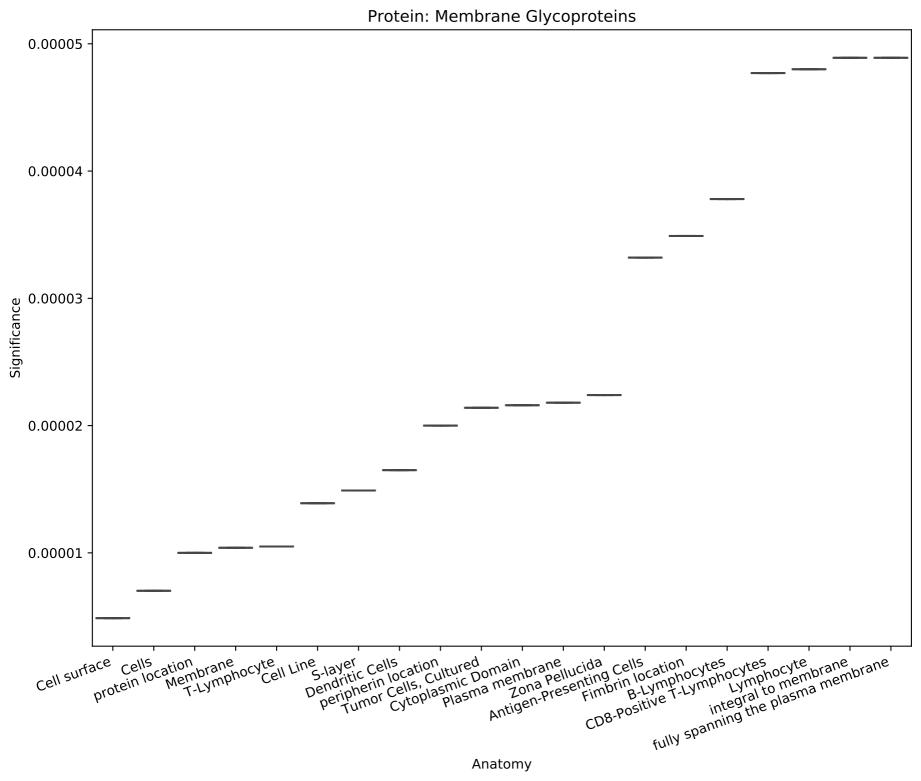


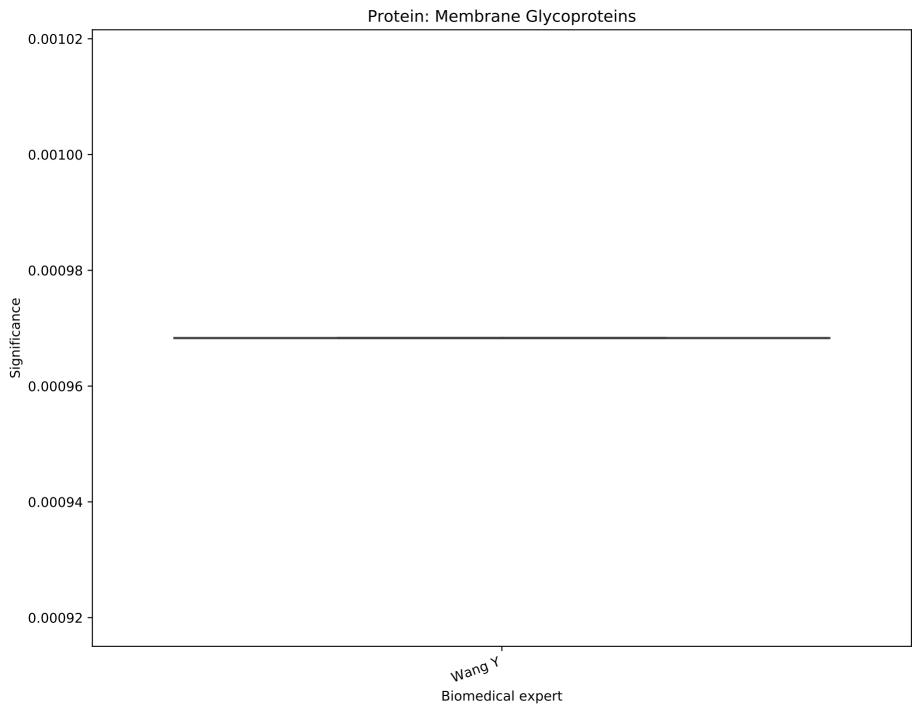
Protein: KRAS 0.00005 0.00004 0.00003 Significance 0.00002 0.00001 Irvival Genes (Oxaliplatin Replicon Replicon Neoplastic Survival Proto-Oncogenes Exons Flaviviridae Survival Replicon, Neoplastic Survival Regulation, Neoplastic Survival Survival Gene Expression Regulation 0.00000 Genes, ras imab Mutation ptors genes civirus Lung erbB-1 Kinase tecan Mutation Mutation Hepacivirus Genes, erbB Receon Hepacivirus Genes, erbB-1 Kinase Irinotecan Irinotecan Genes Genes Irinotecan Irinotecan Genes Genes Irinotecan Genes Genes Irinotecan Genes Genes Irinotecan Genes MeSH

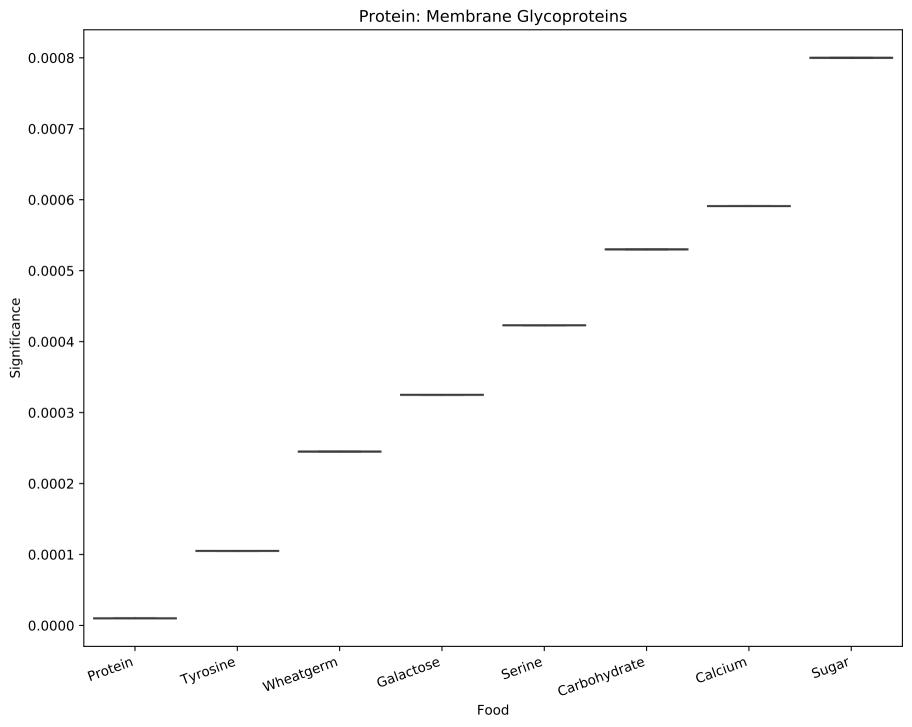


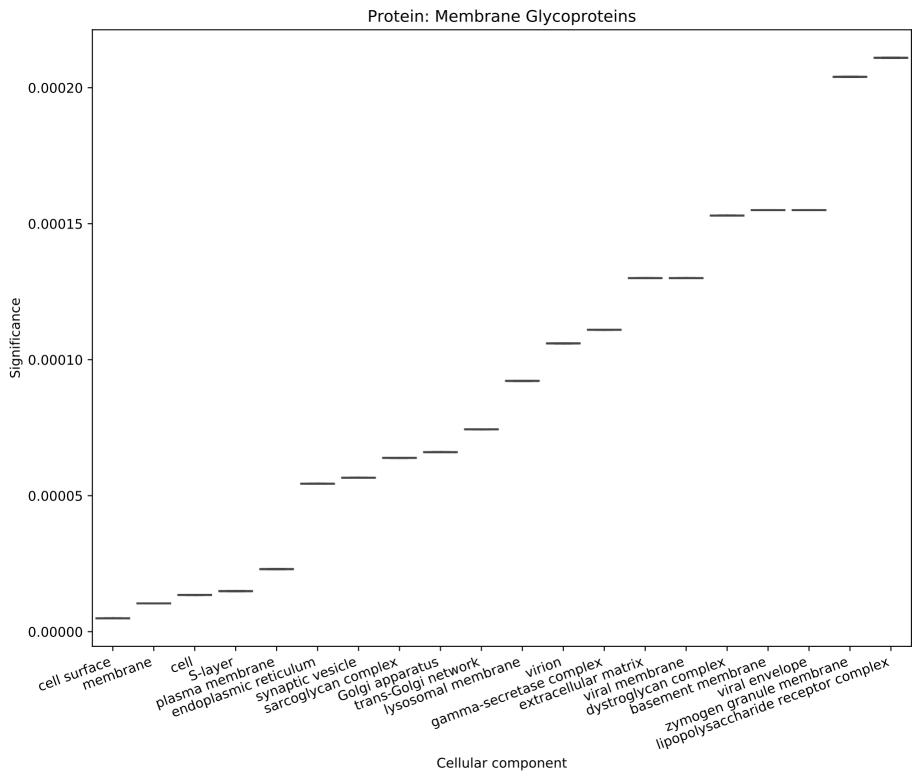


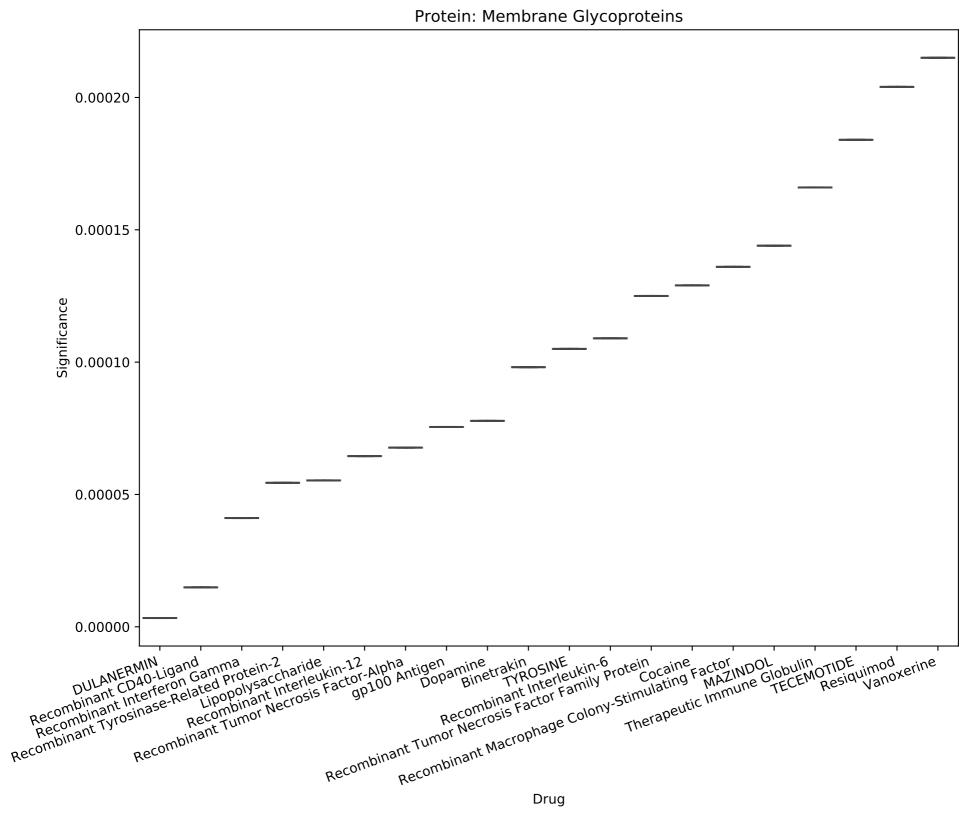


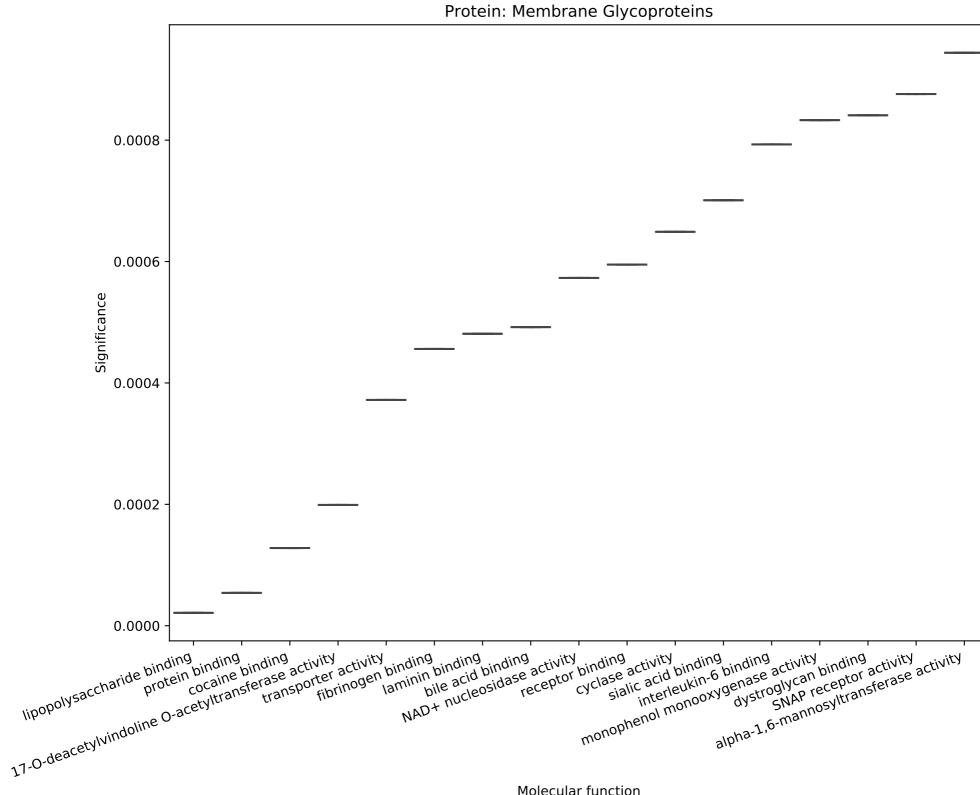


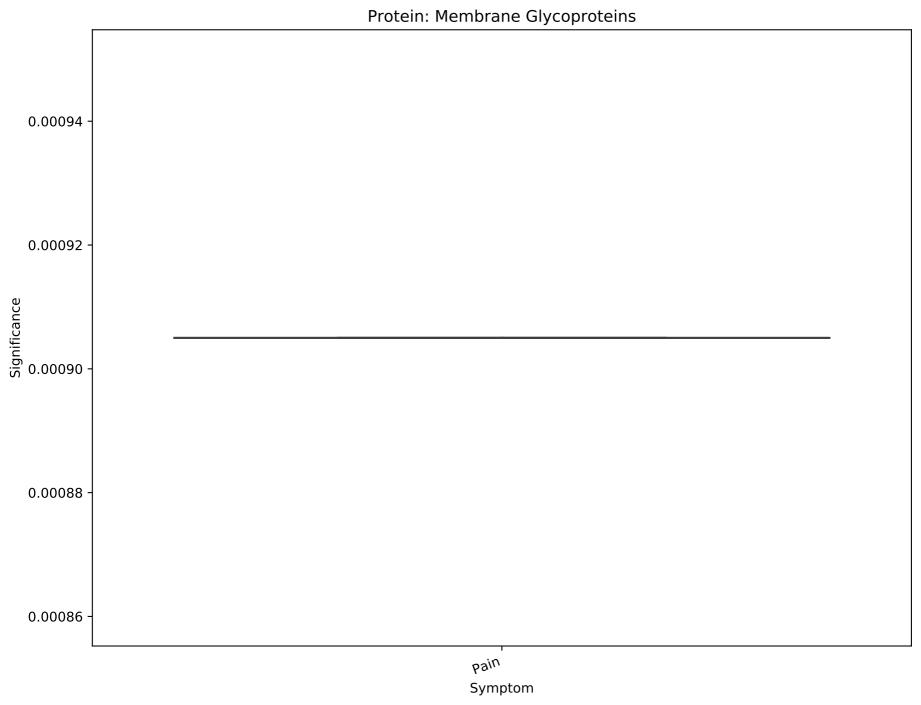


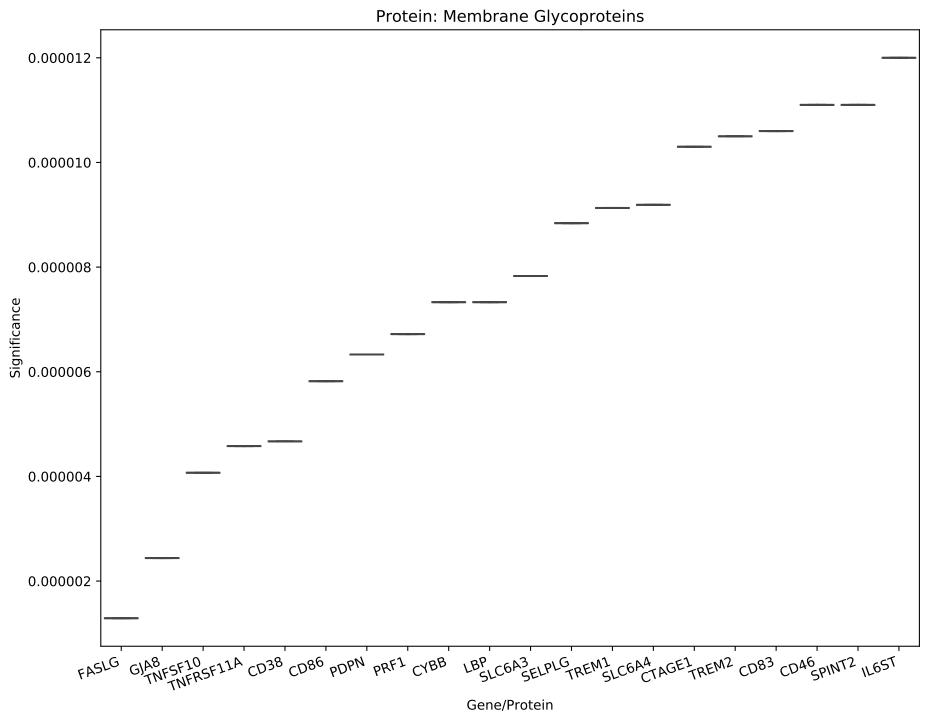


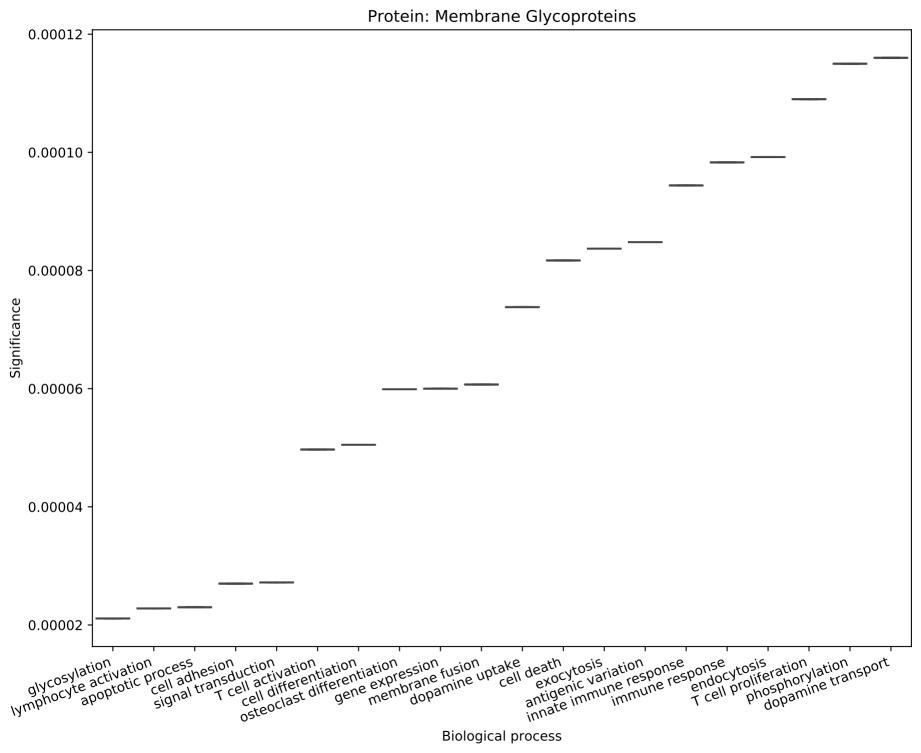


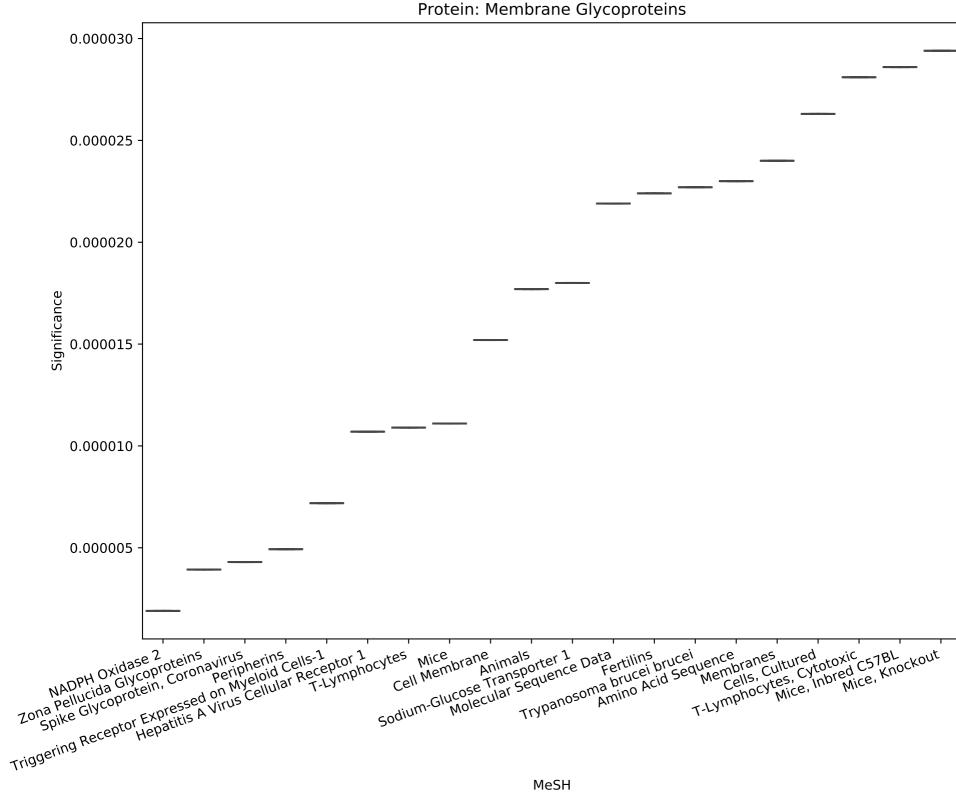


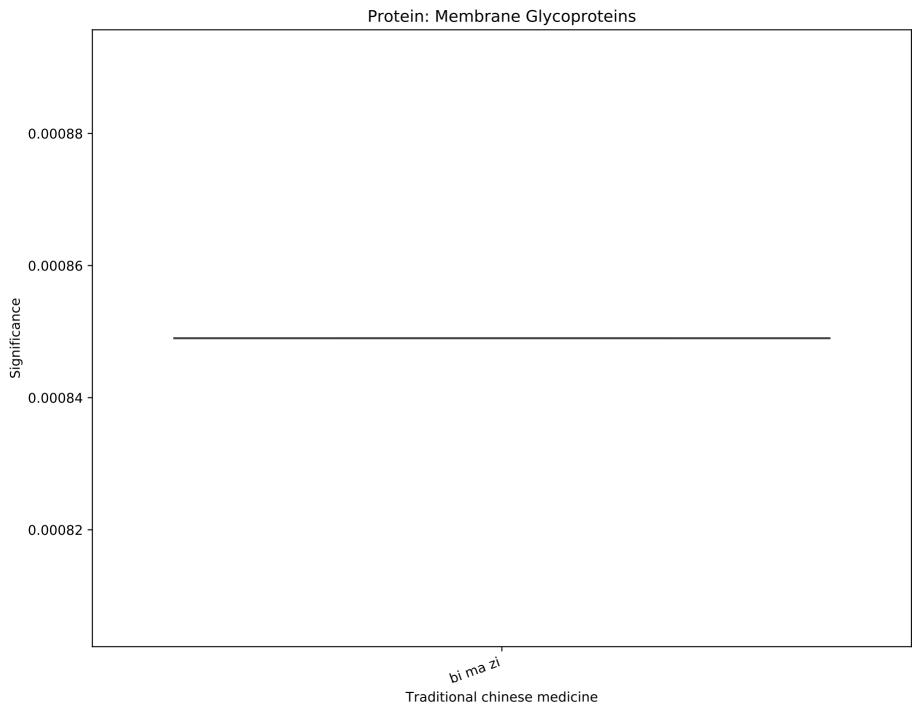


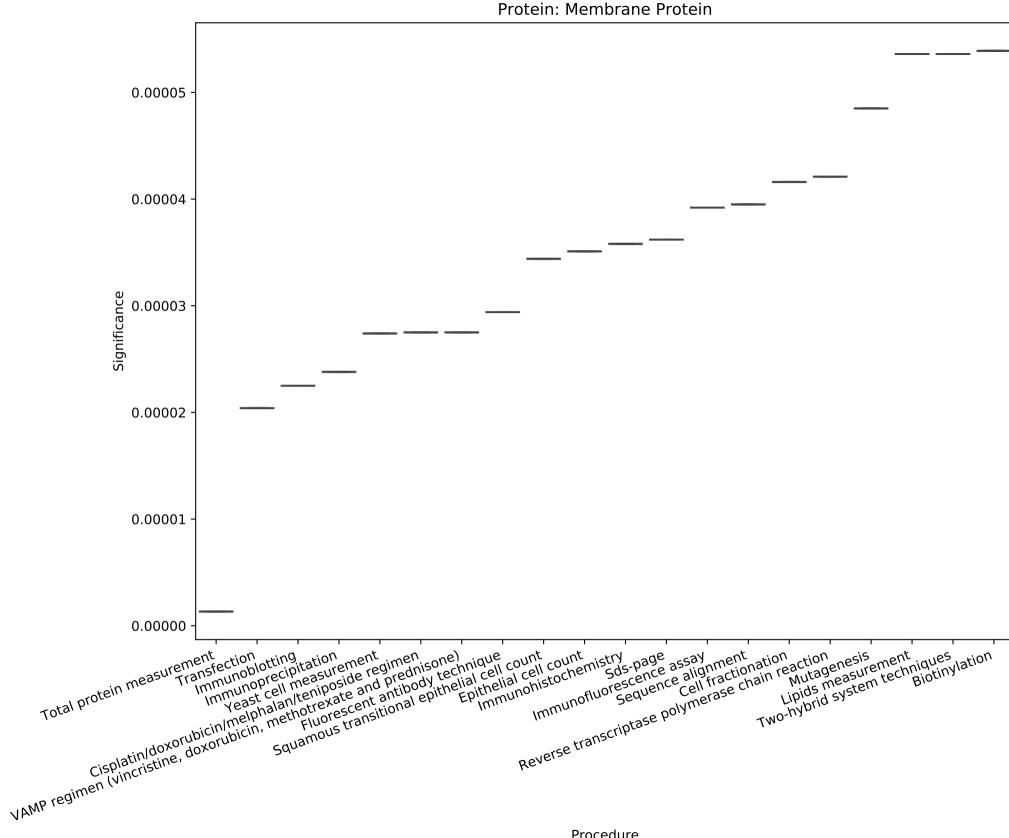


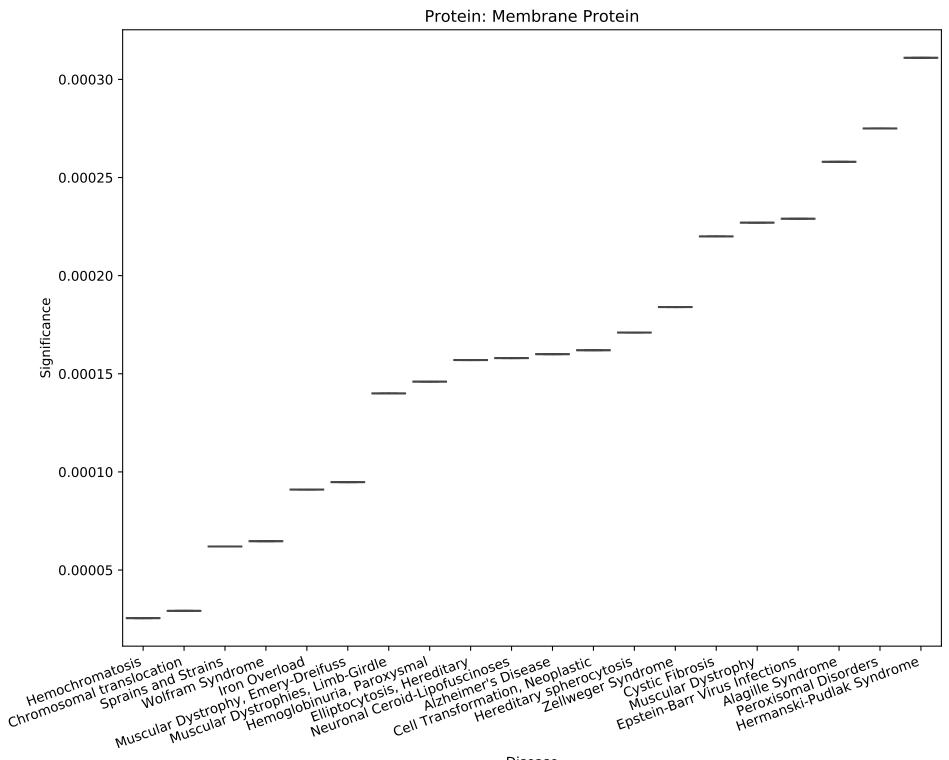




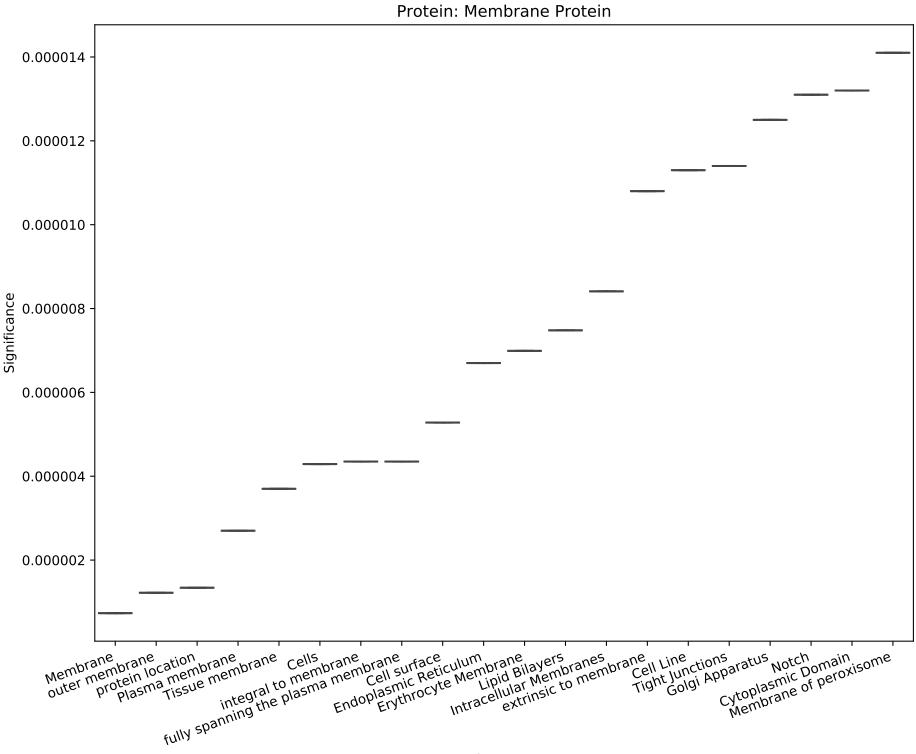




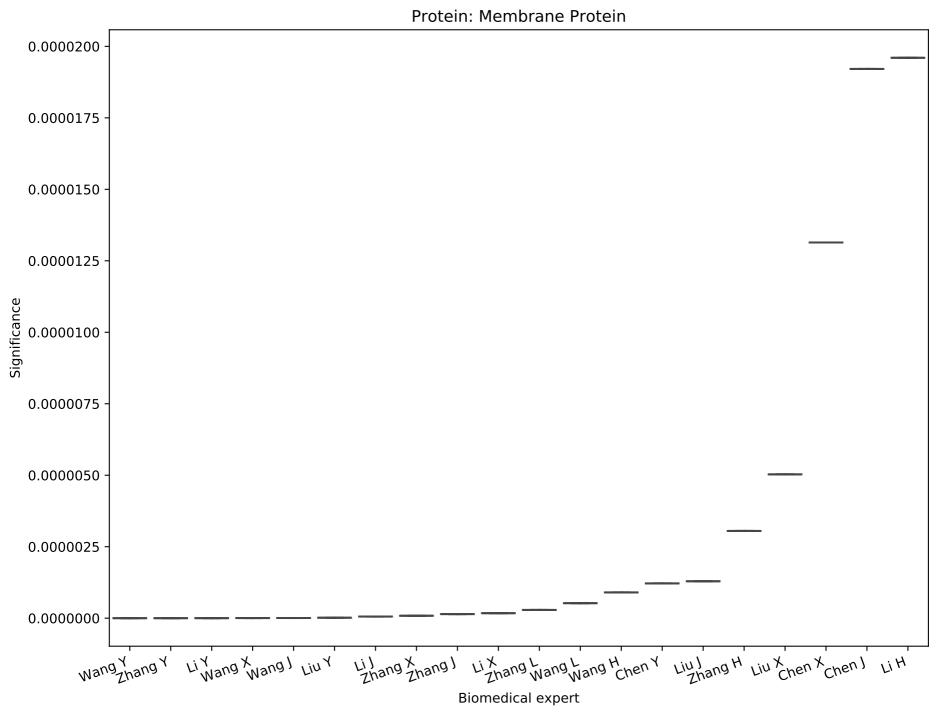


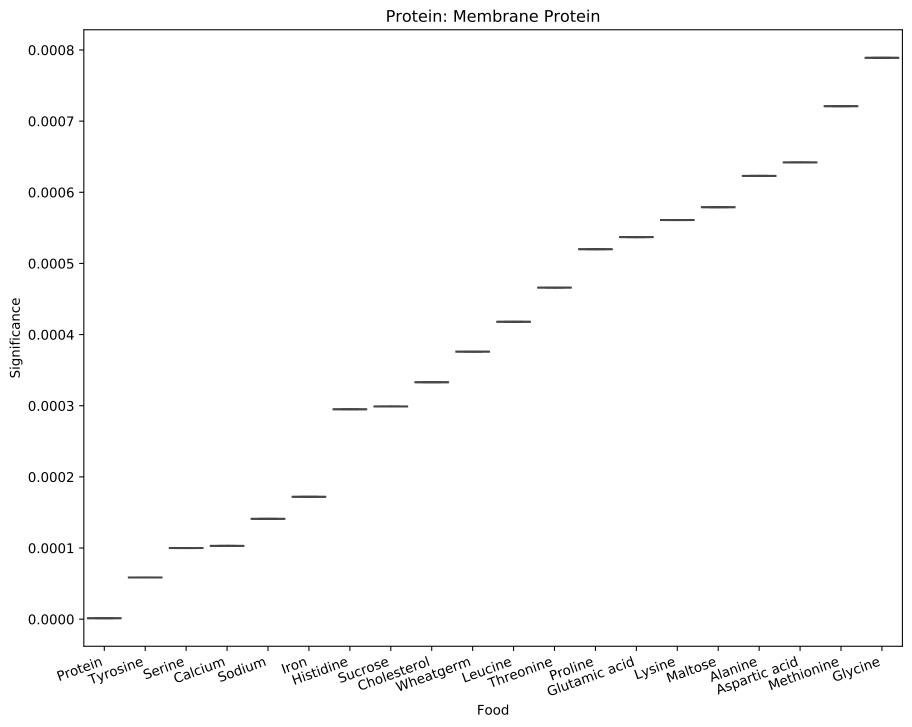


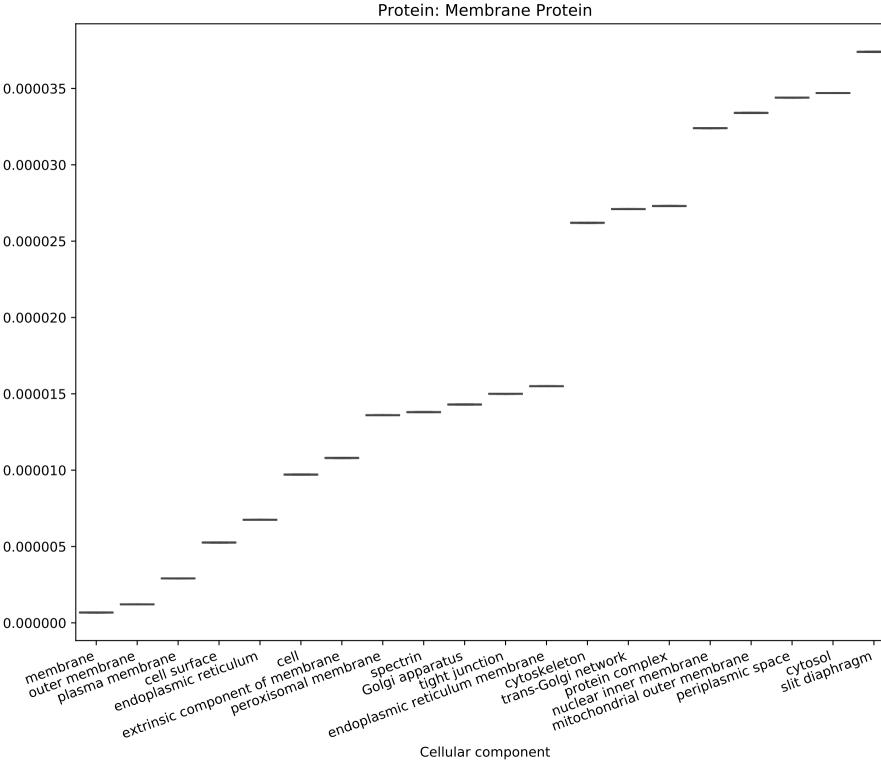
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Anatomy

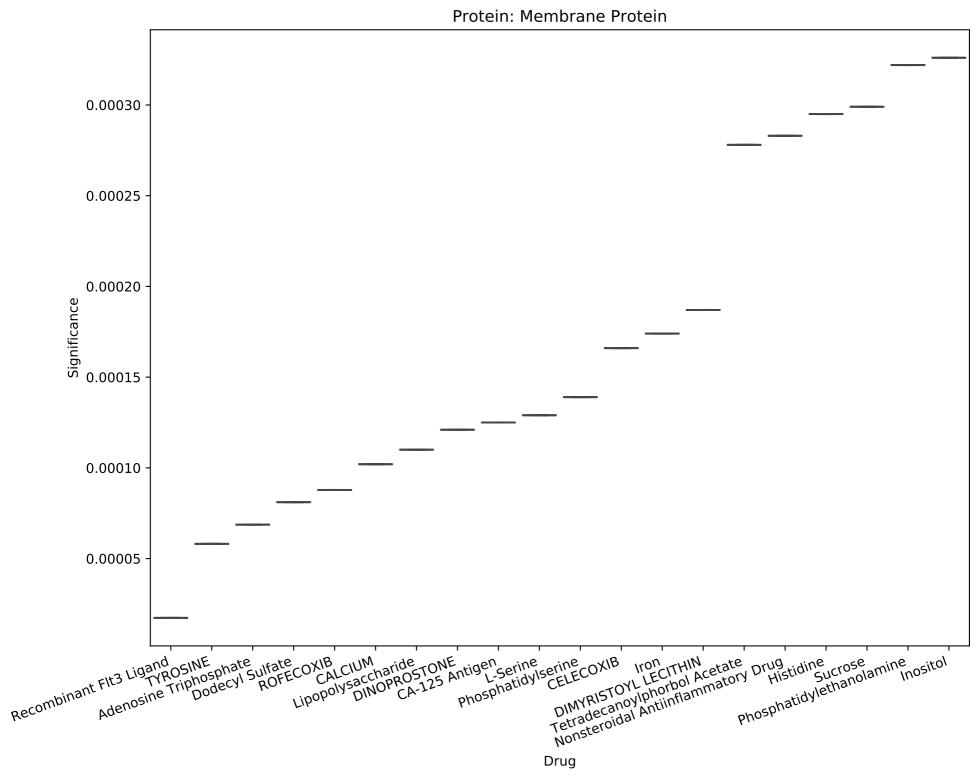


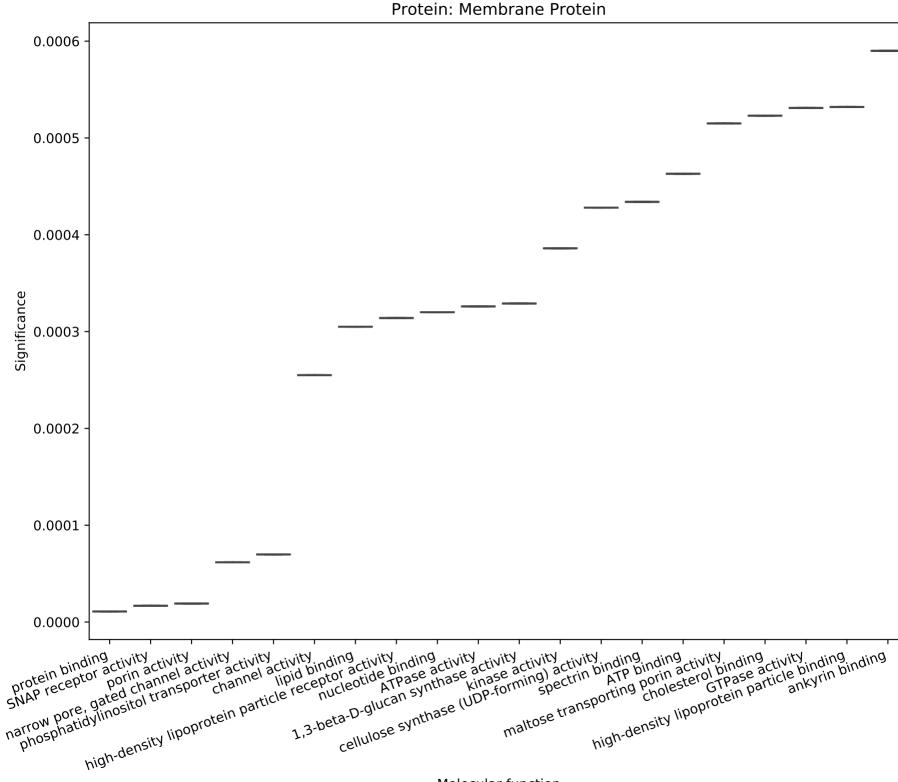


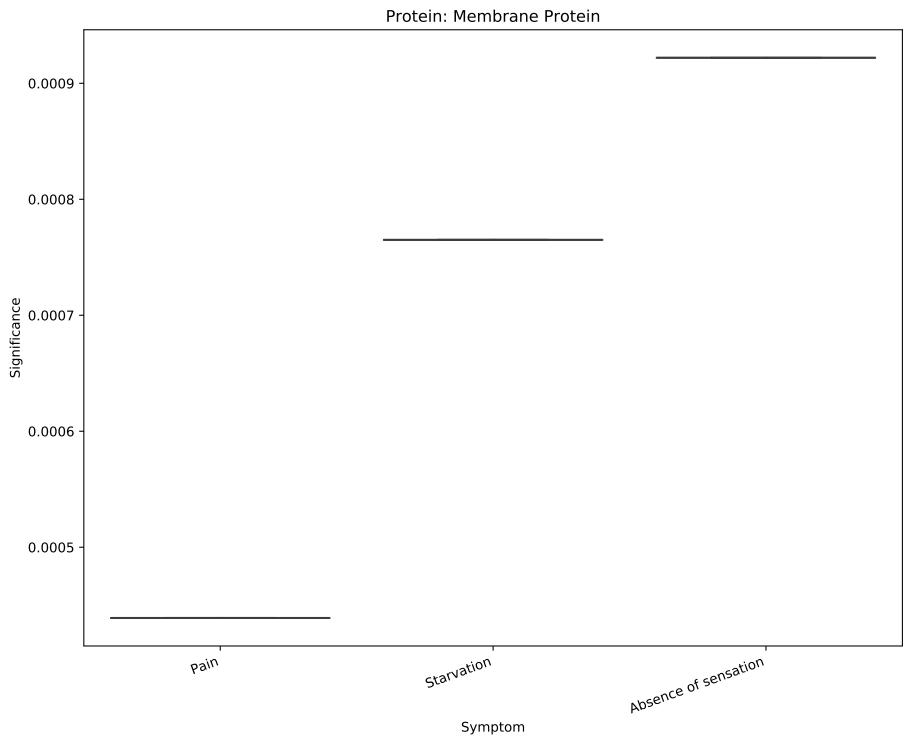


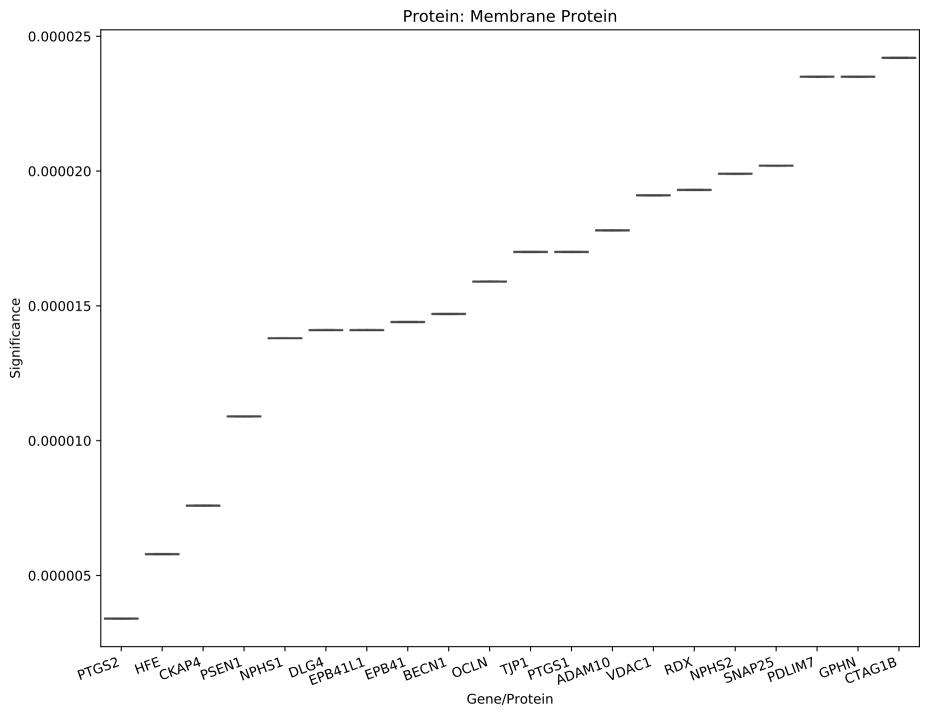
Significance

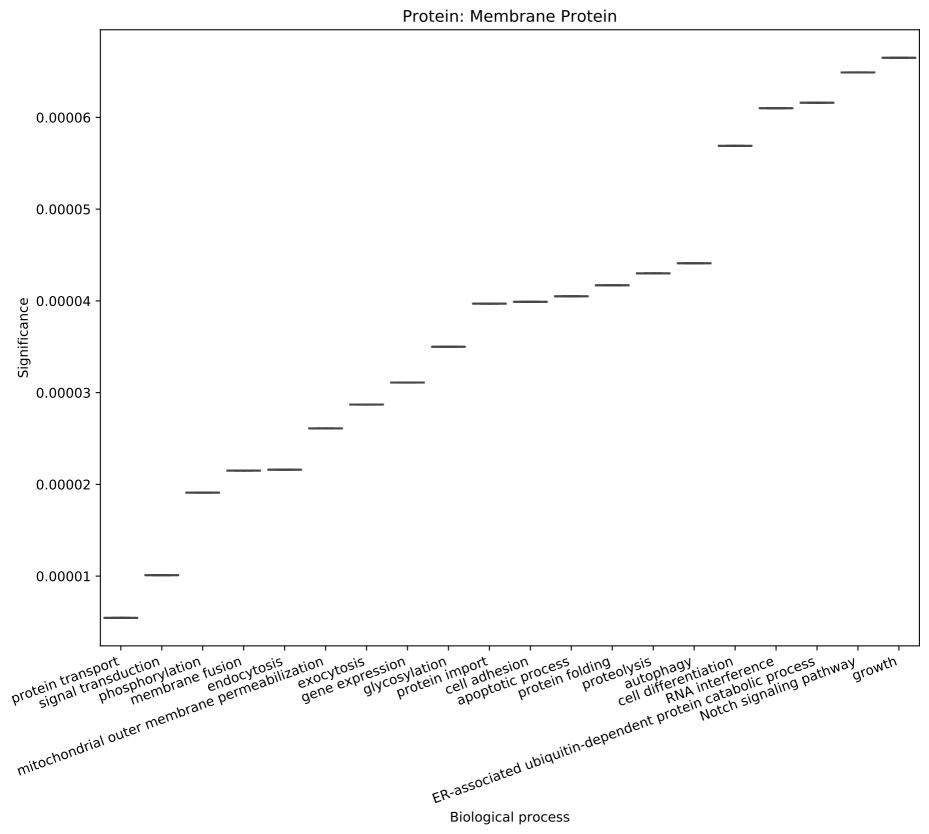
Cellular component

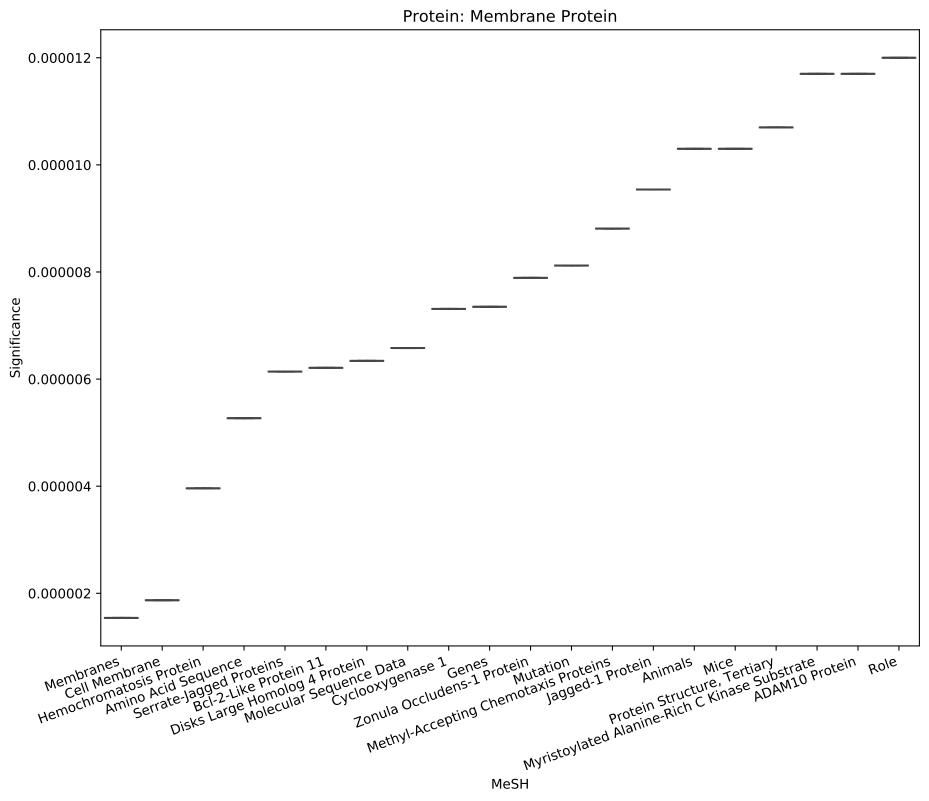


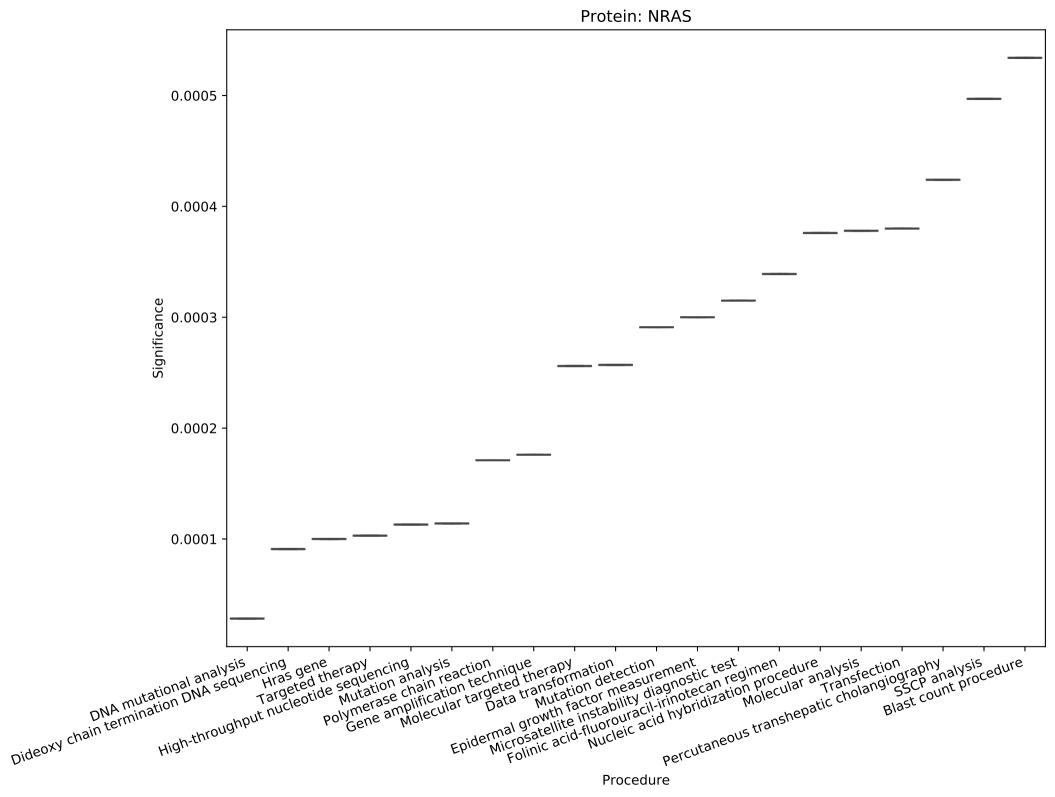


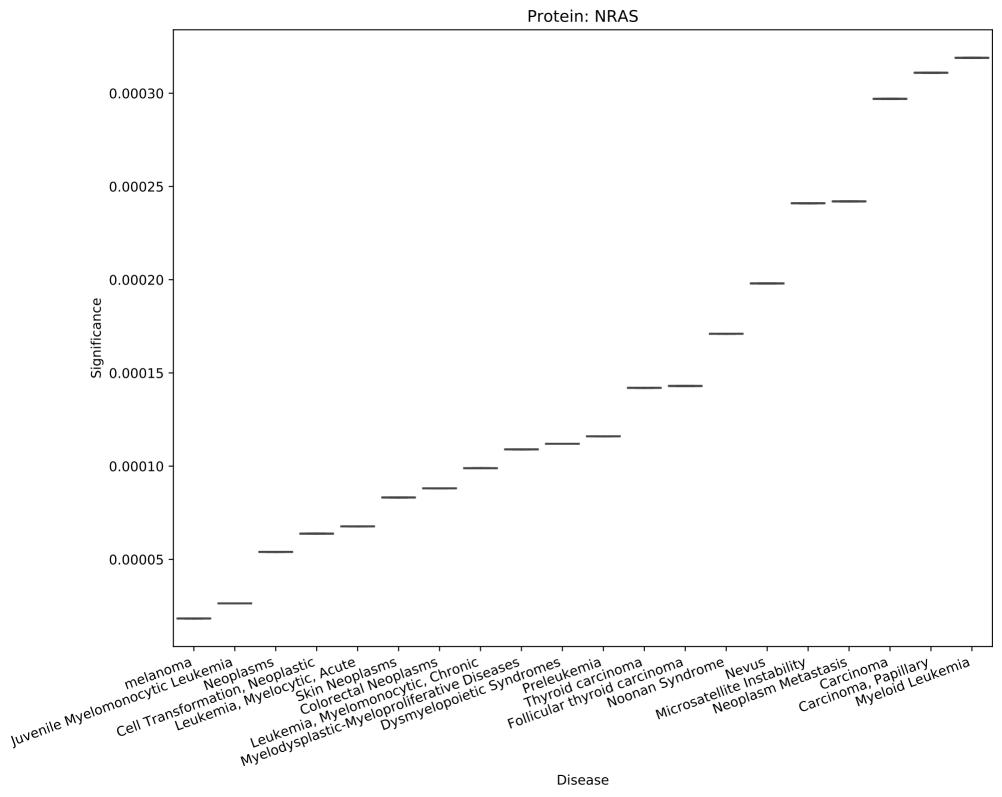


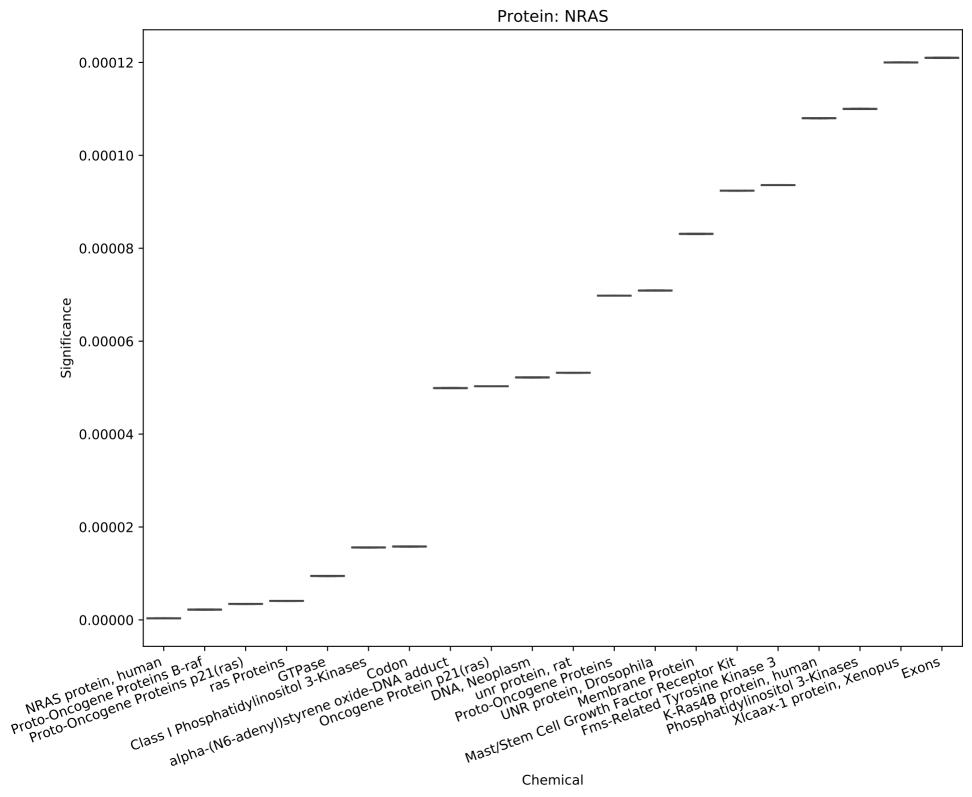


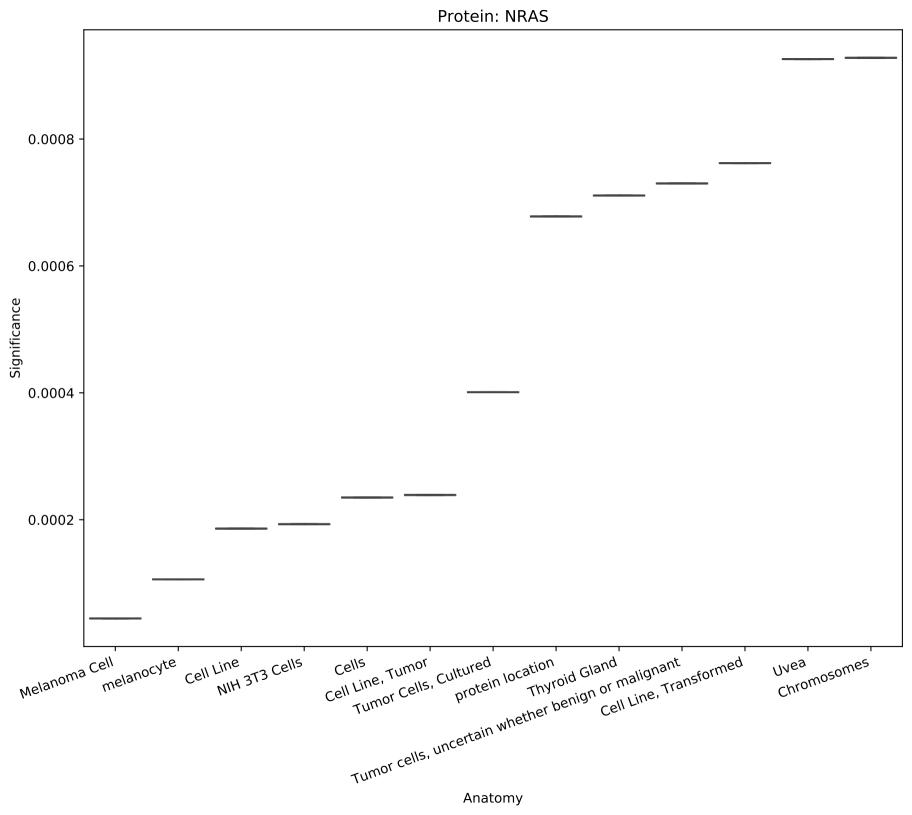


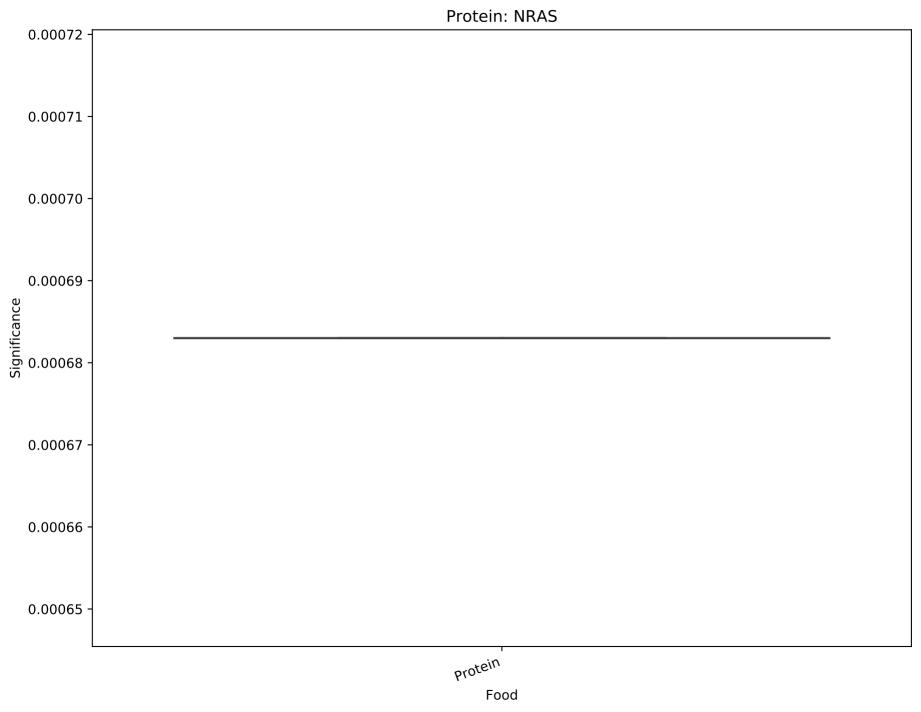


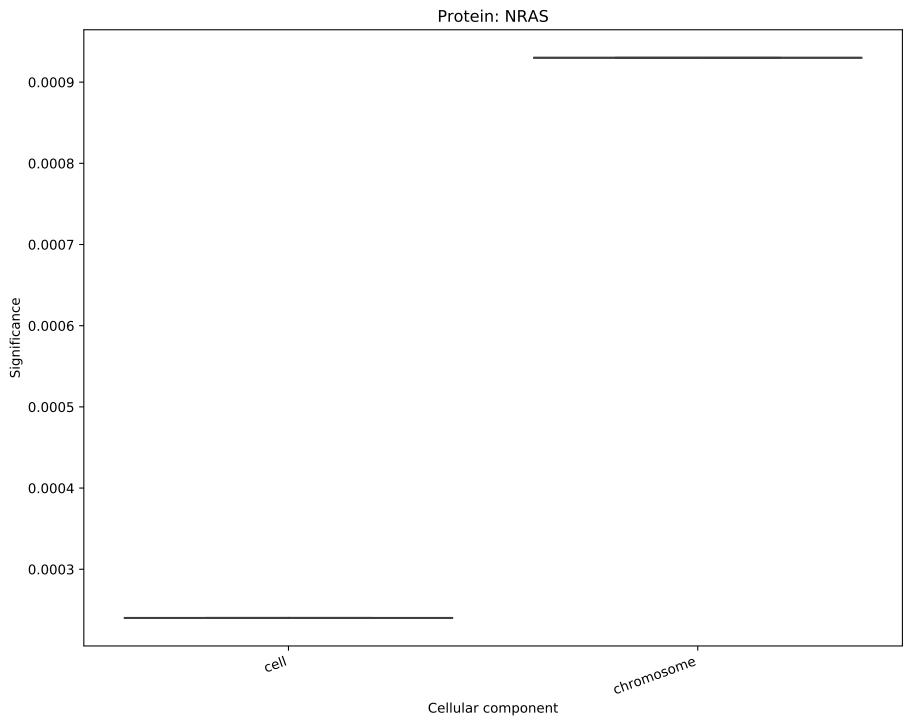


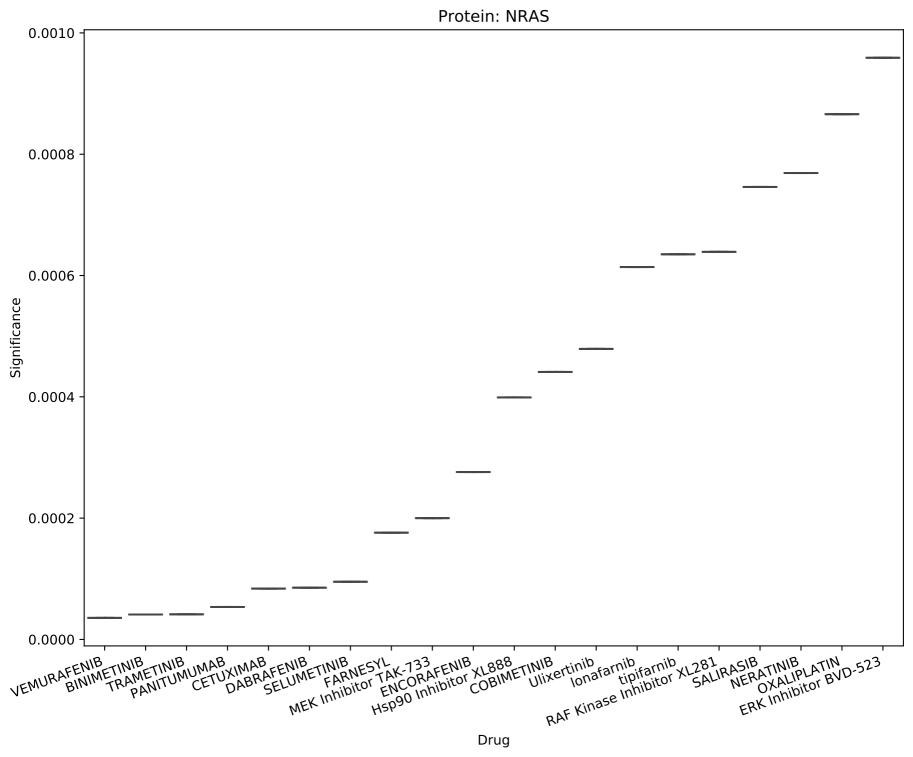


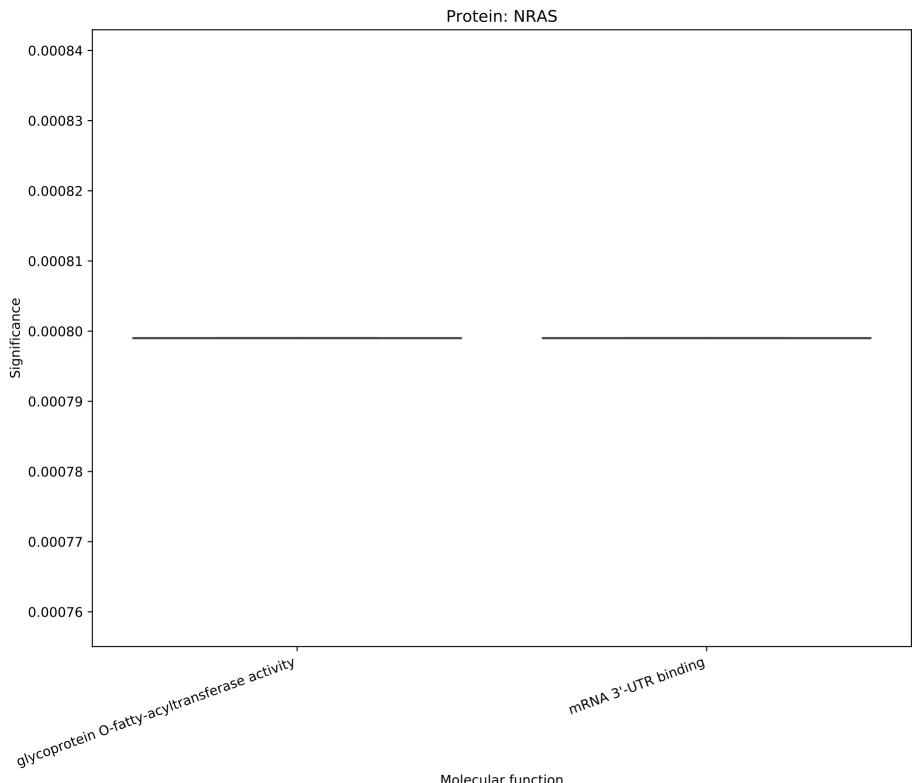


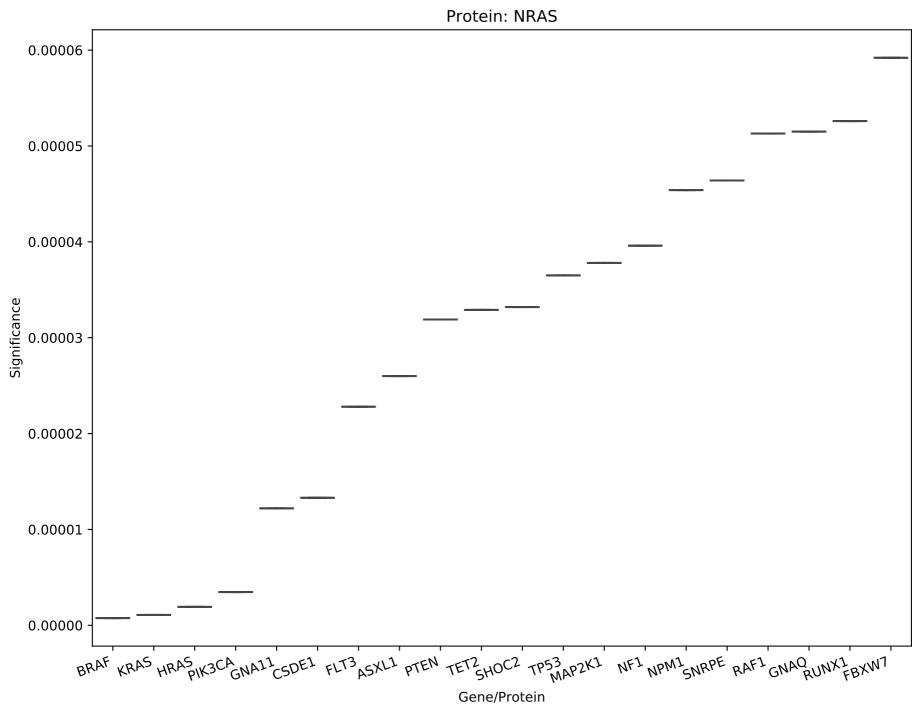


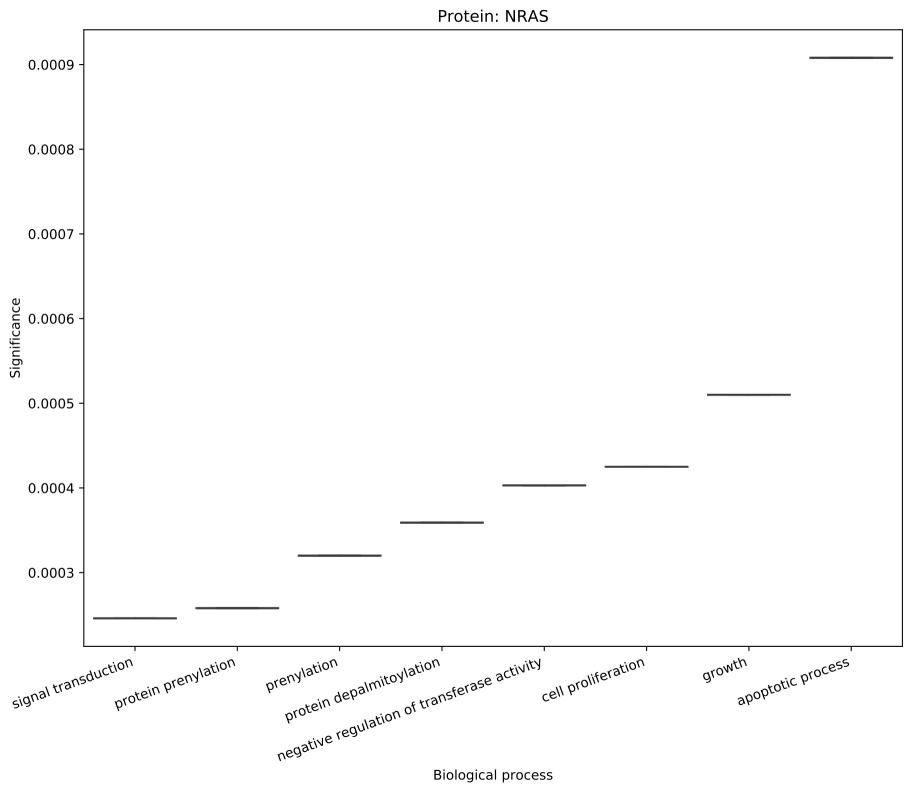




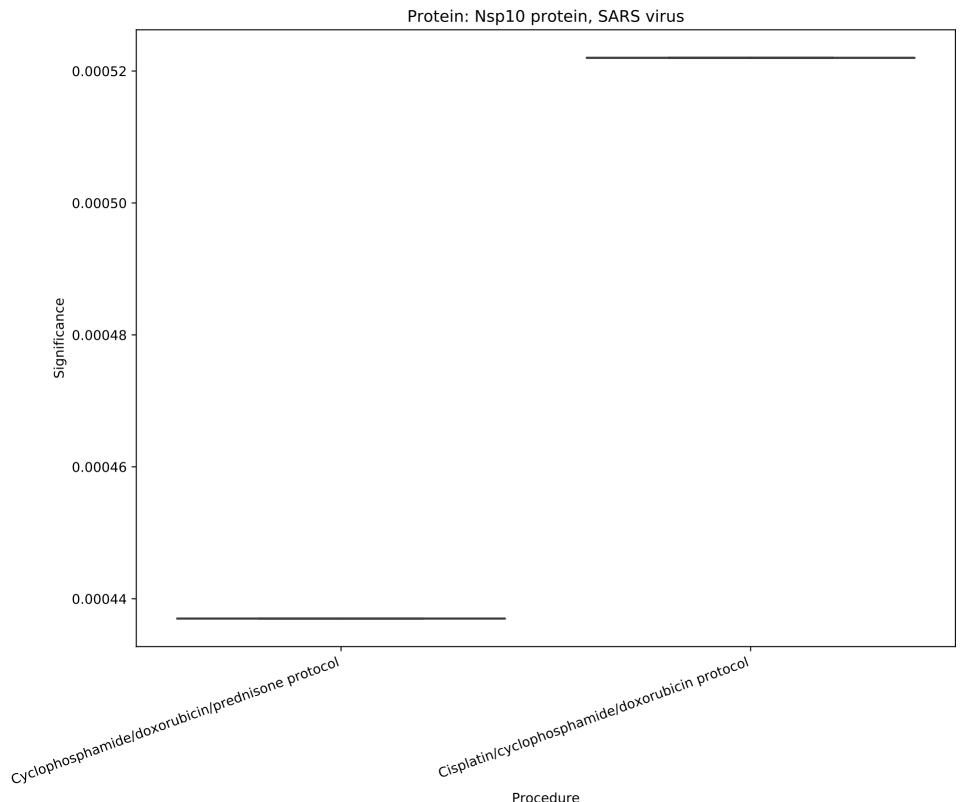


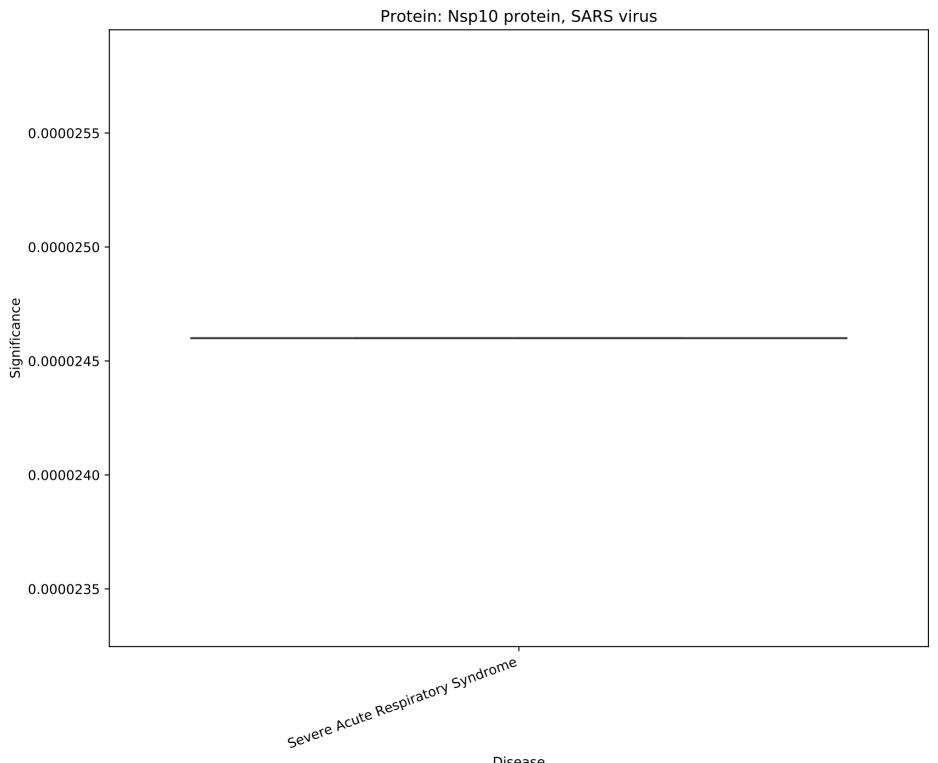


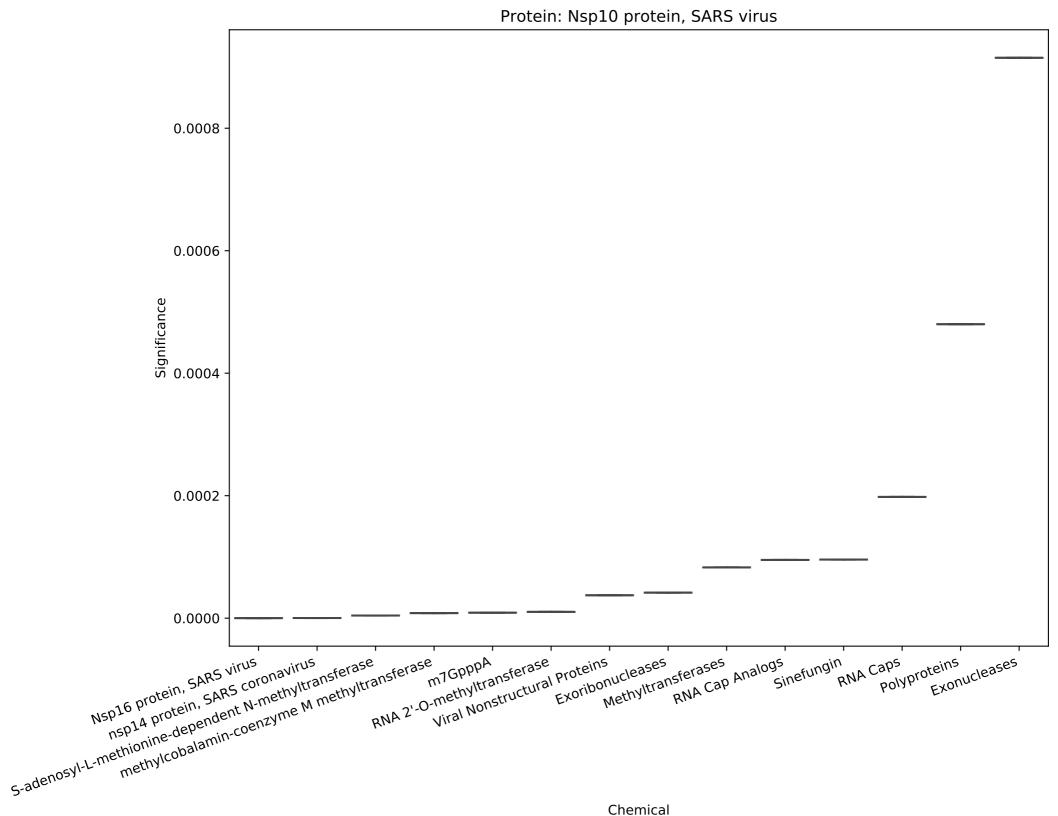


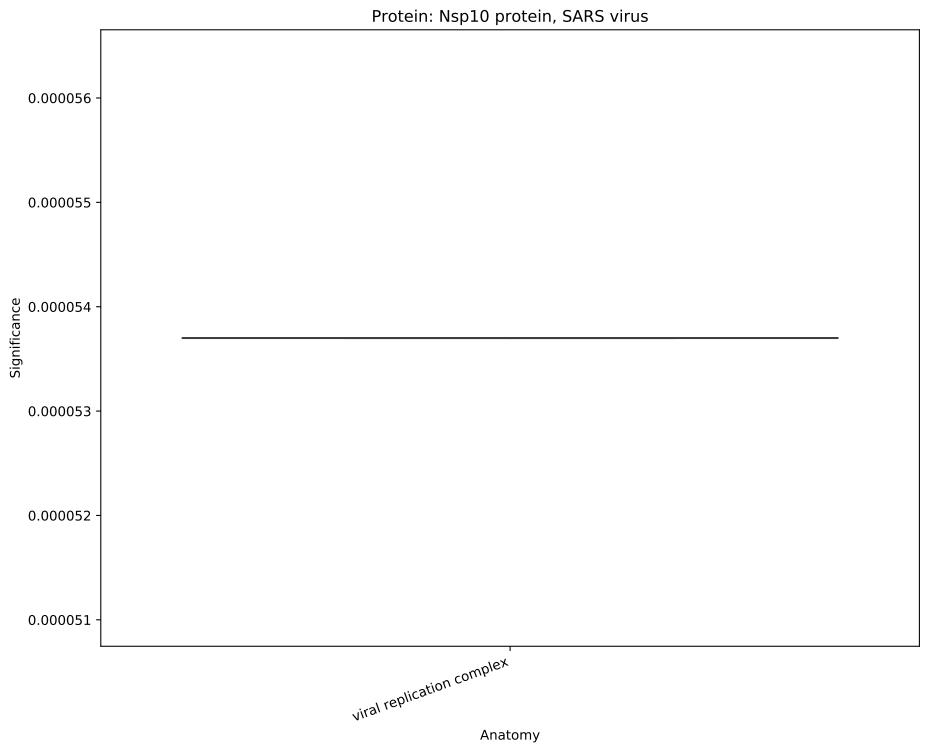


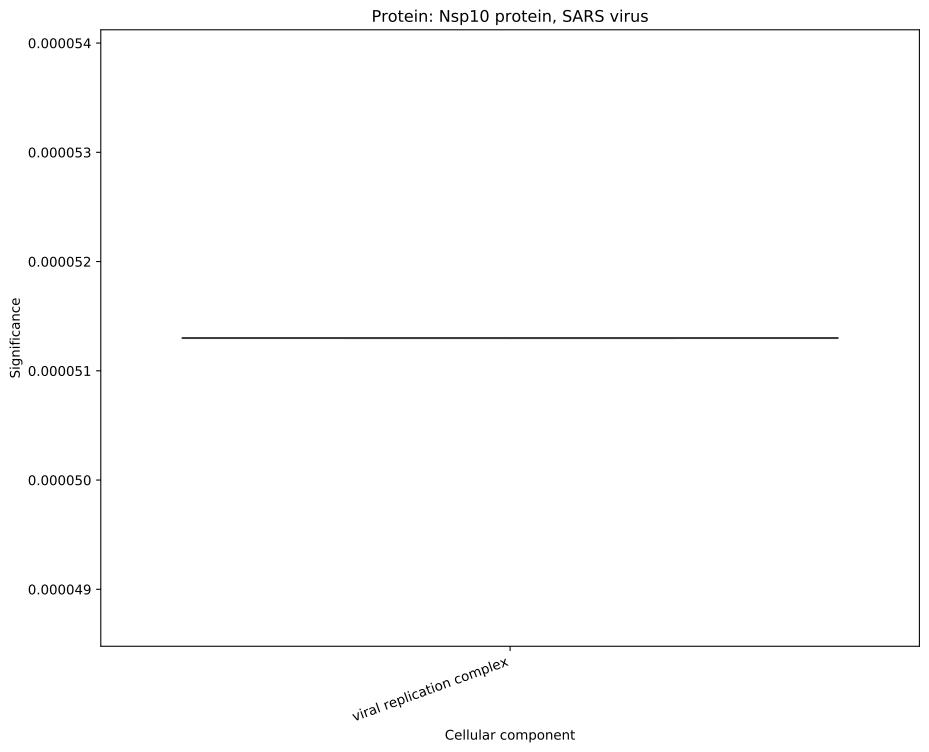
Protein: NRAS 0.00016 0.00014 0.00012 0.00010 Significance 0.00008 0.00006 0.00004 0.00002 0.00000 Genes, ras genes tion genes fenib tation mab Genes Exons imab Rate ptors, p53 ytes ional Neoplastic proto-Oncogenura Mutation and Panitumum Genes Exons imab Rate ptors, p53 ytes ional lastic proto-Oncogenura Mutation Receptors proto-Oncogenura Mutation proto-Oncogenura Mutation Prenylation Receptors proto-Oncogenura Mutation Prenylation Receptors proto-Oncogenura Mutation Receptors proto-Oncogenura Receptors proto MeSH

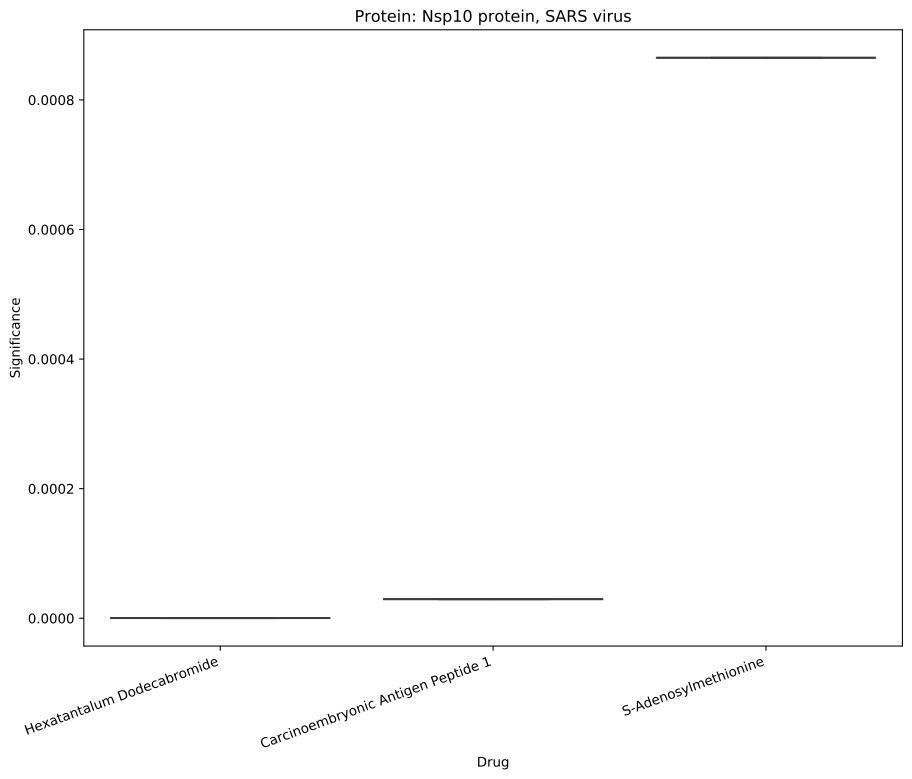


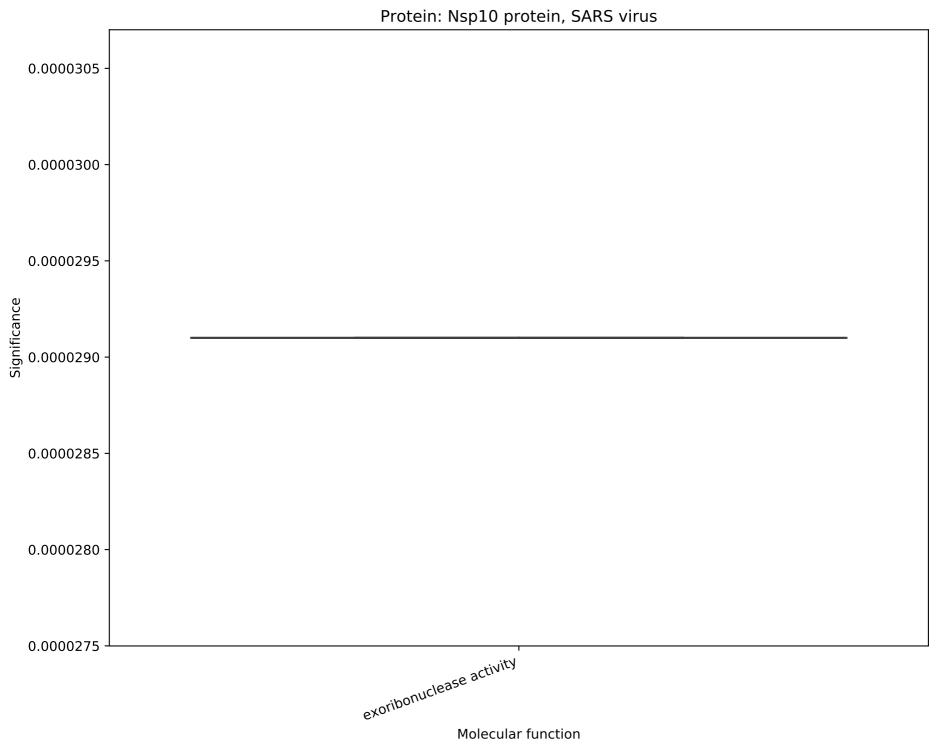


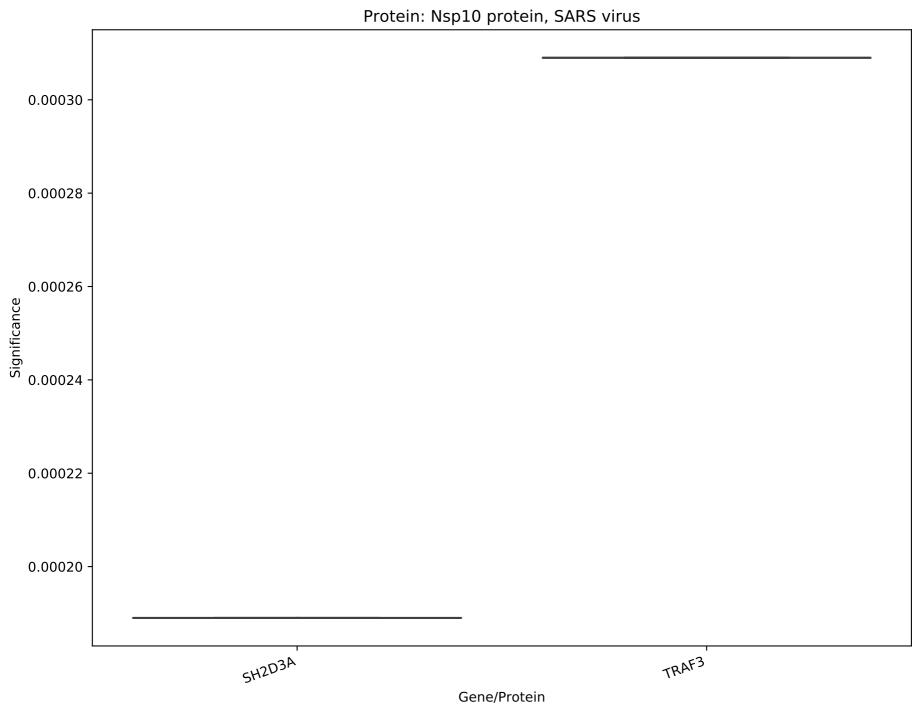


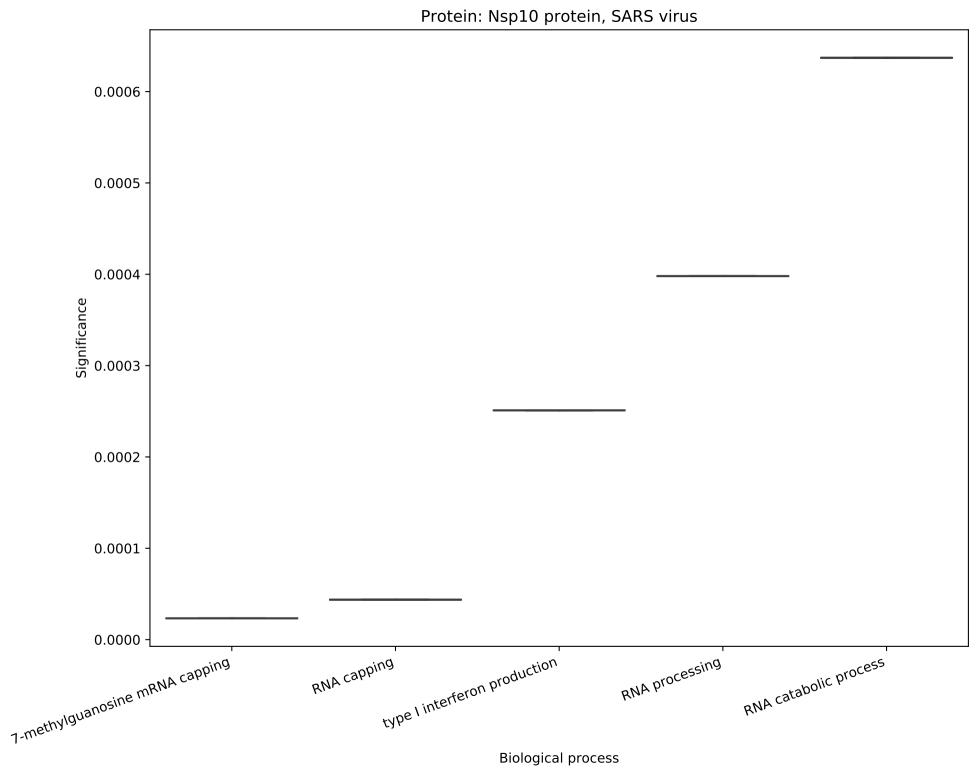


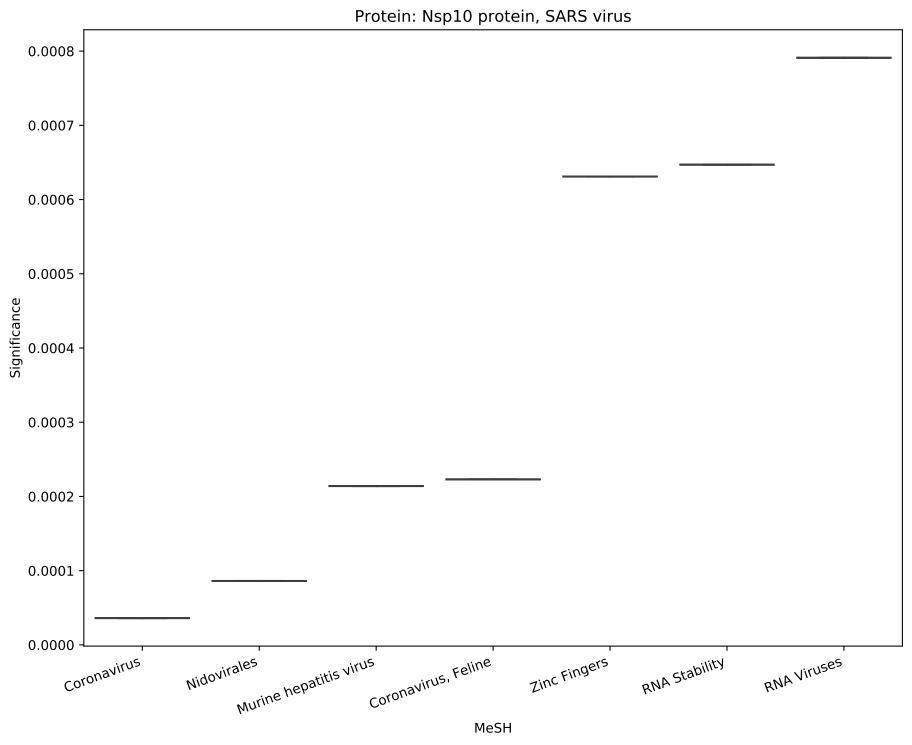


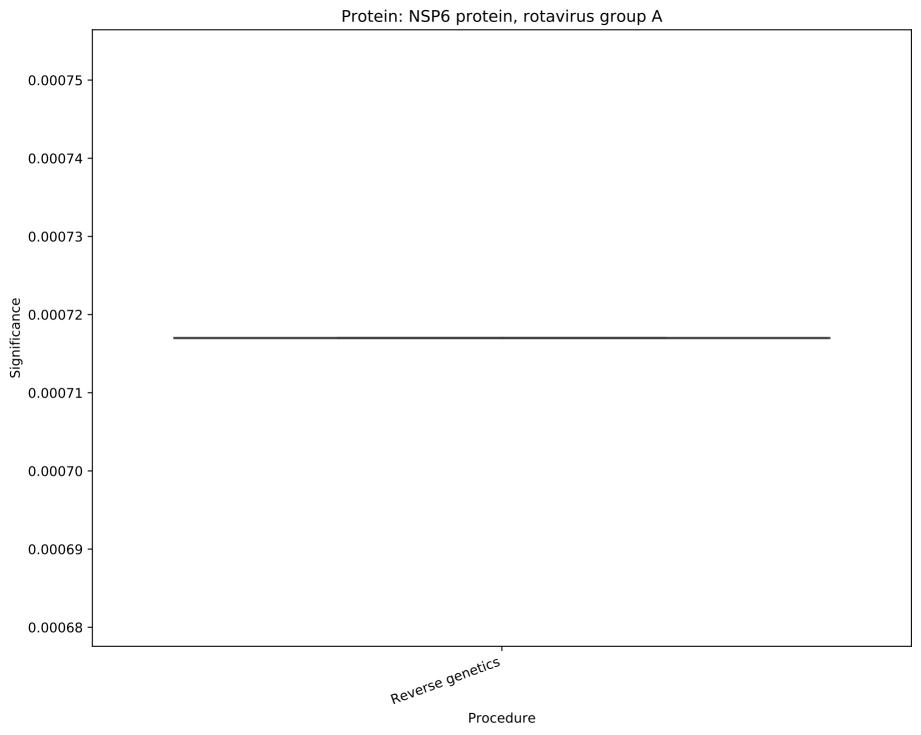


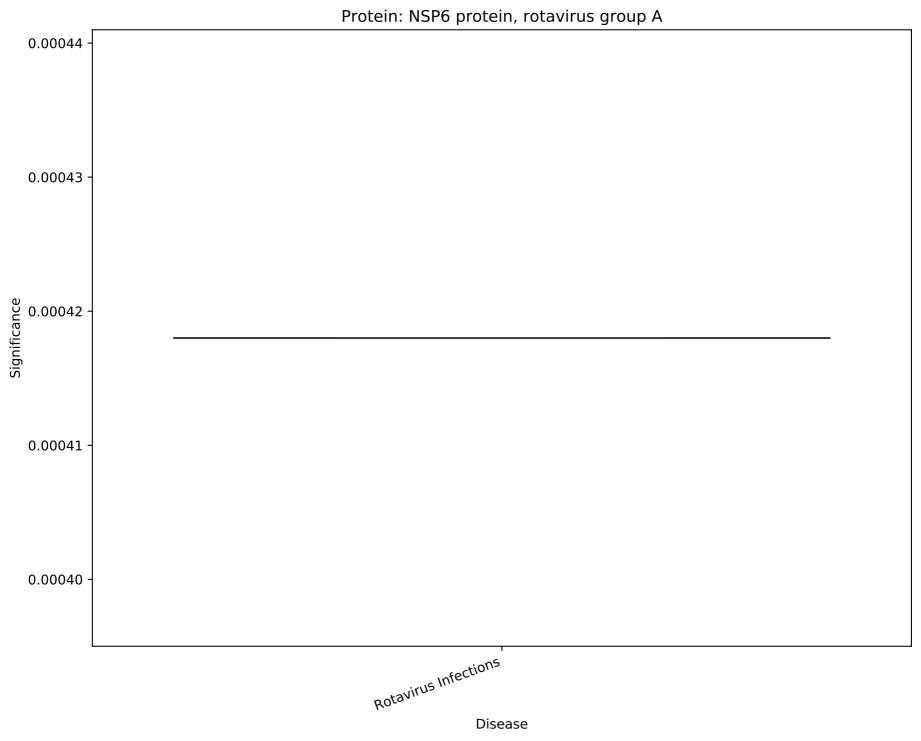


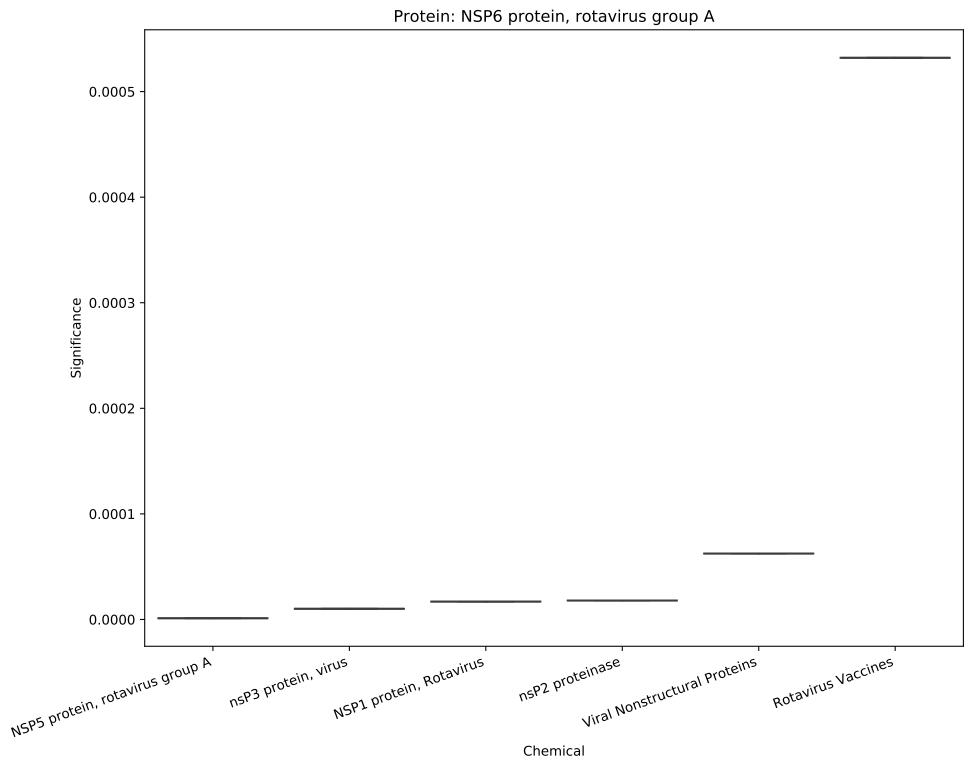


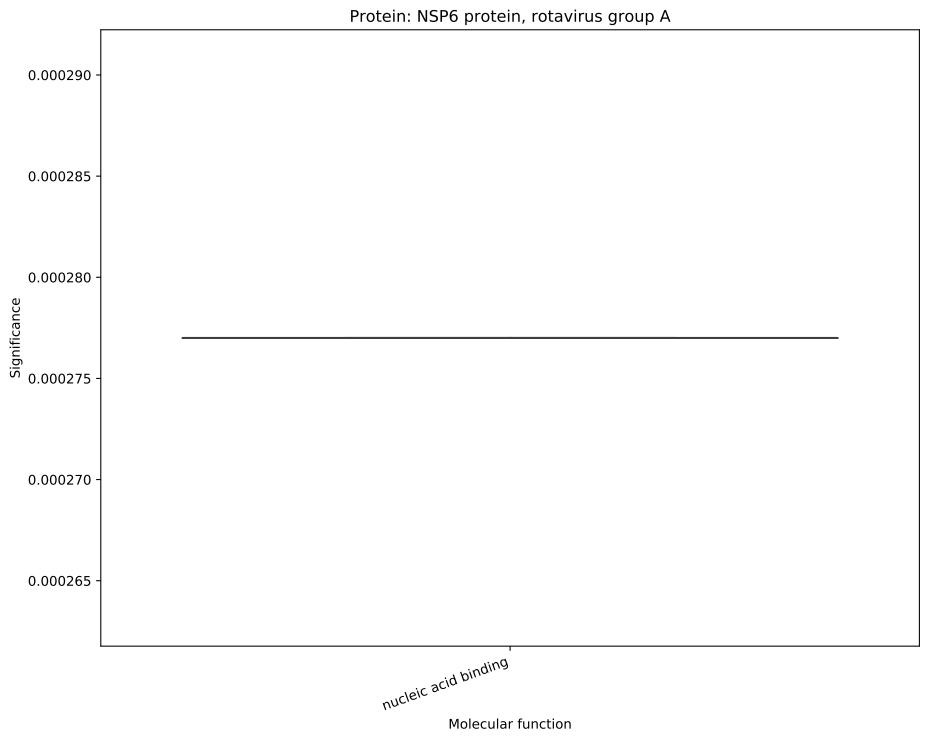


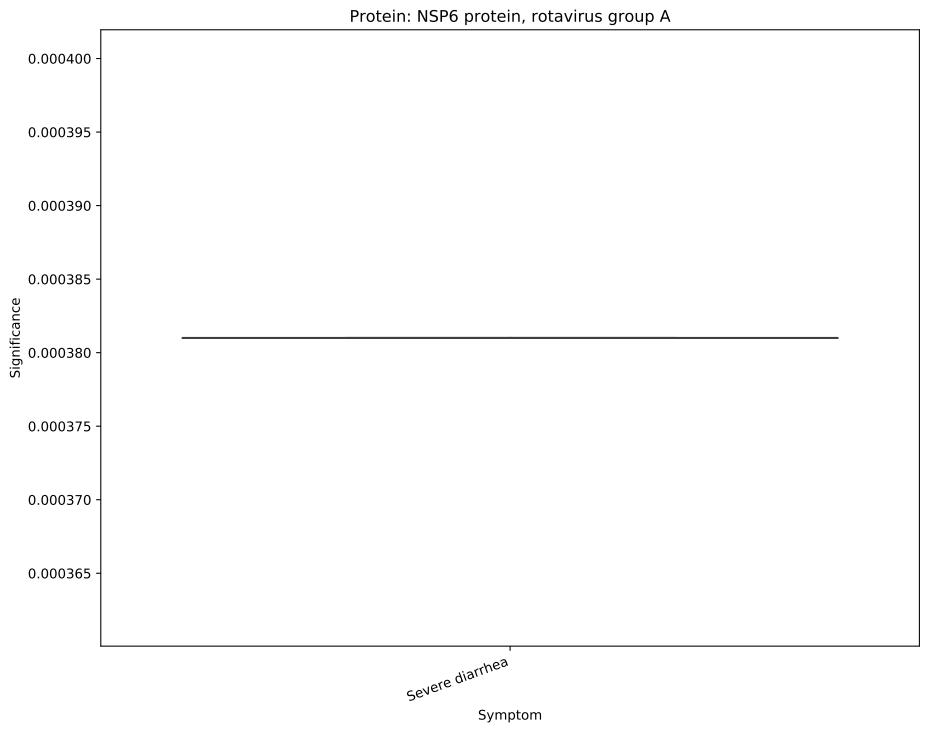


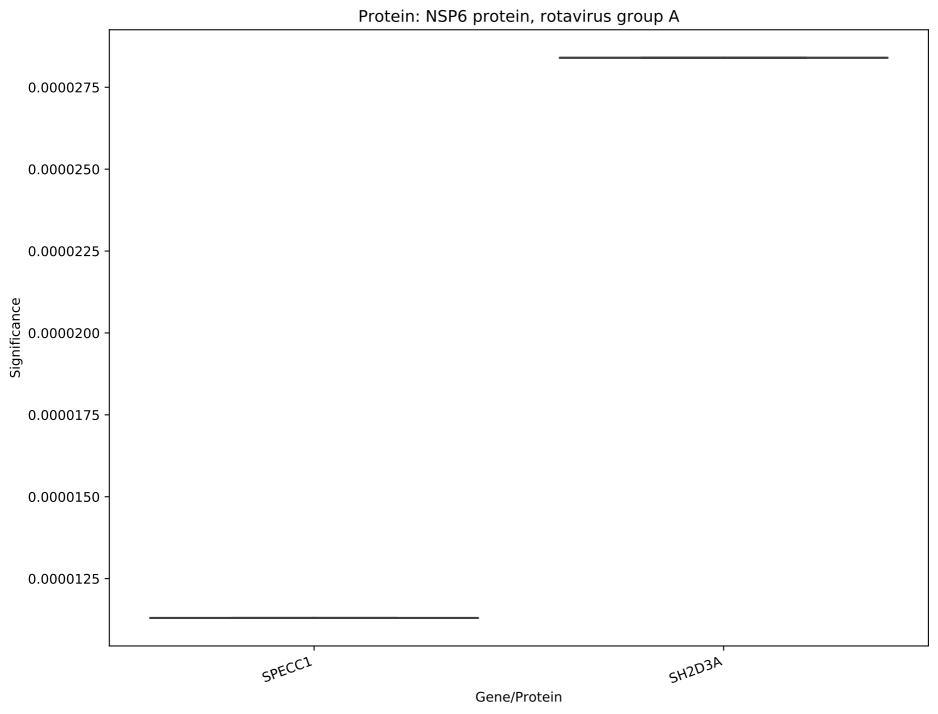


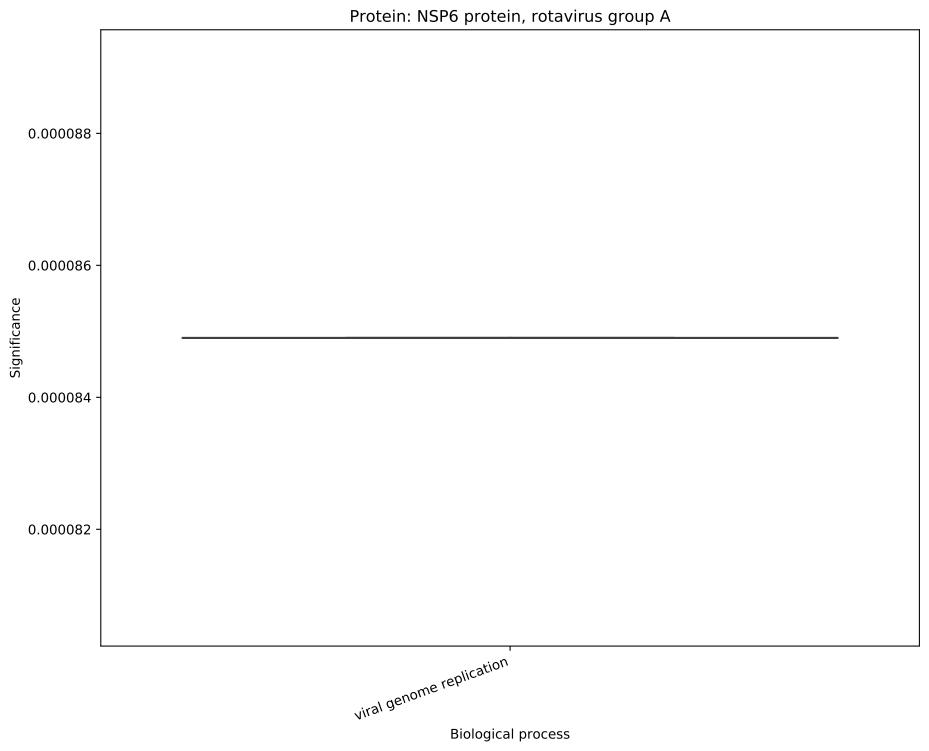


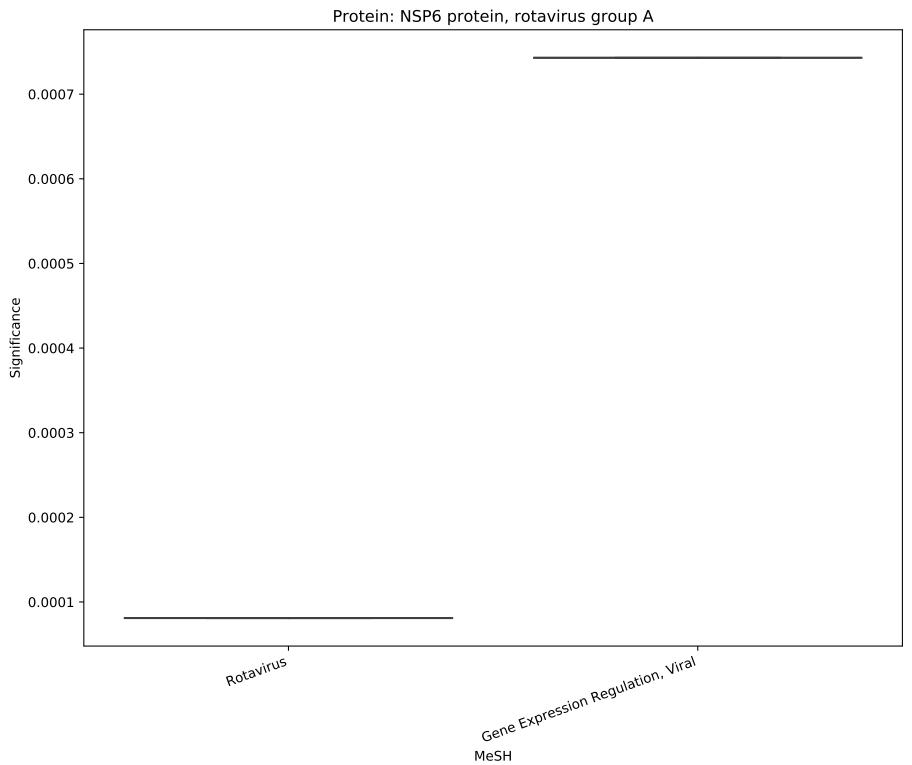


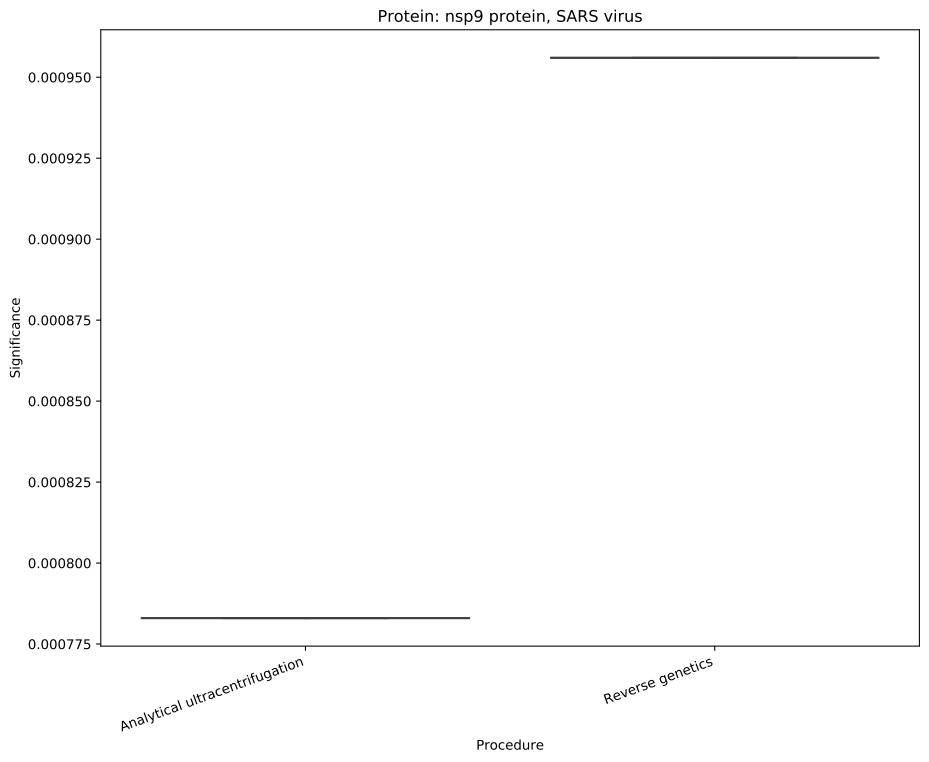


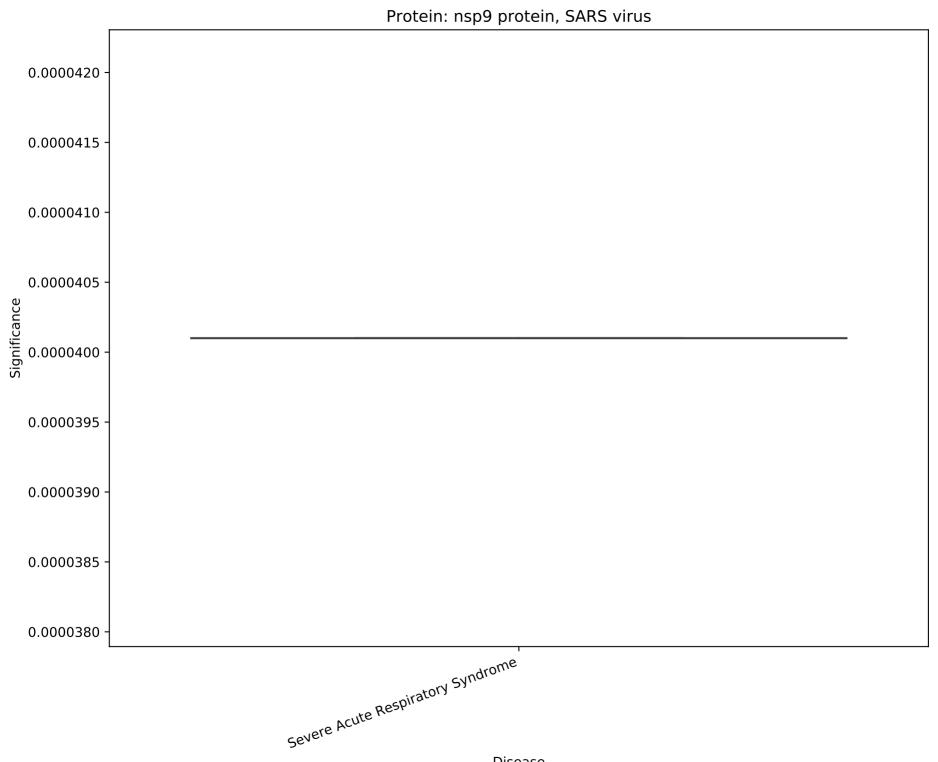


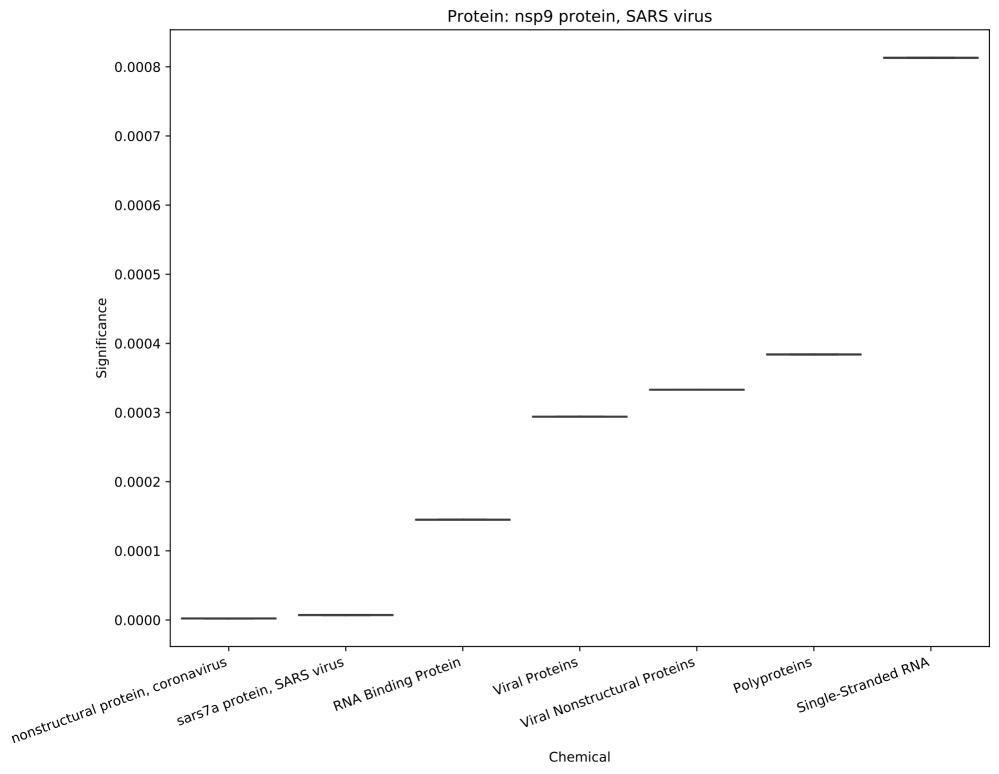


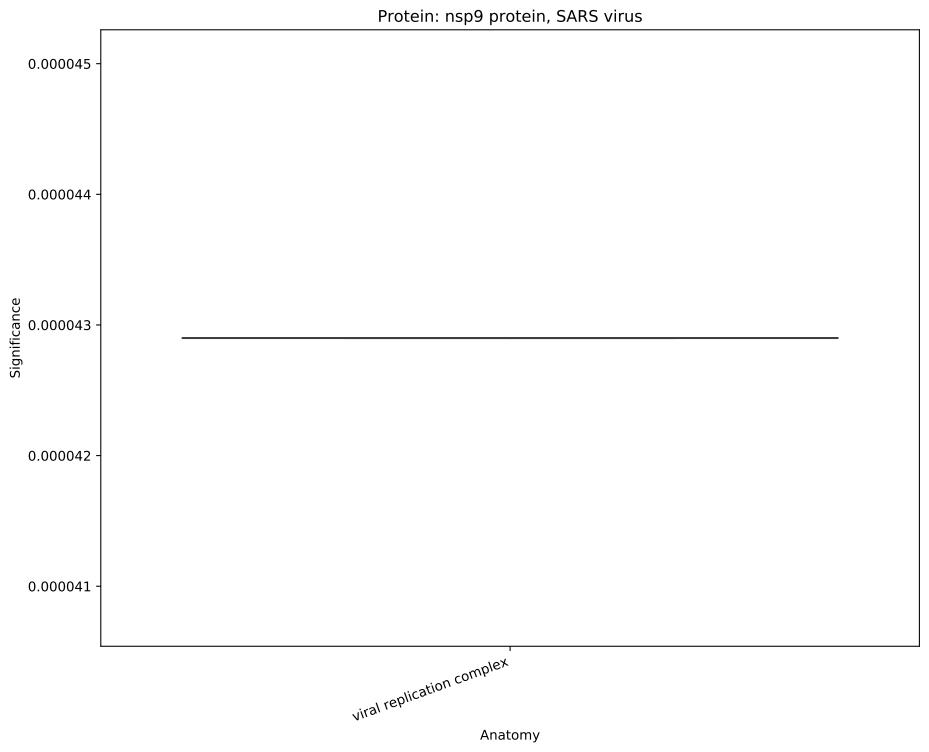


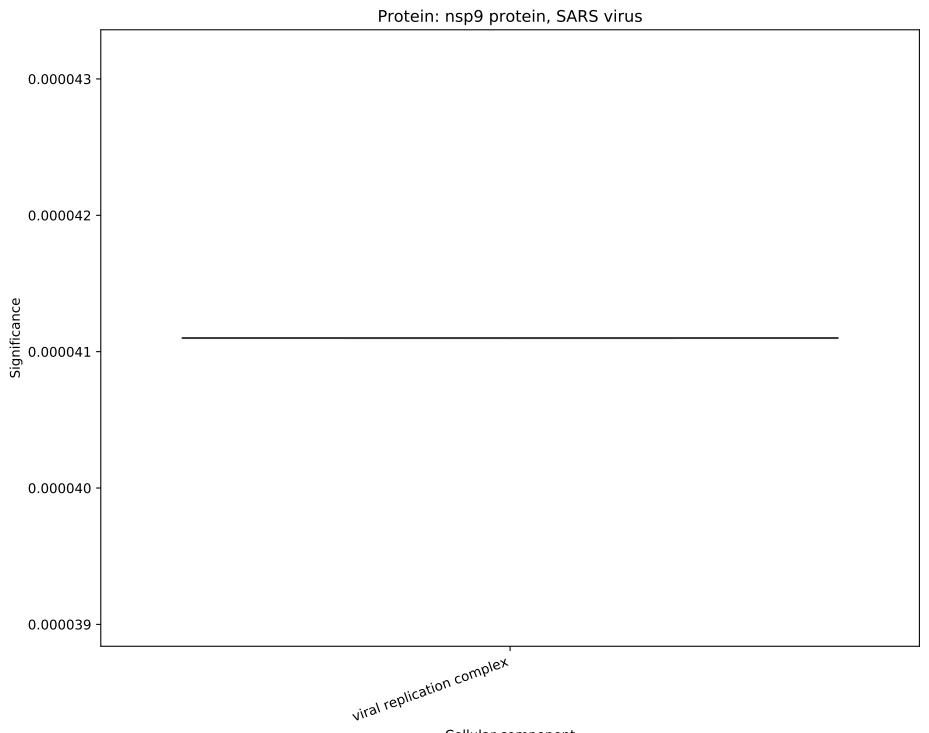




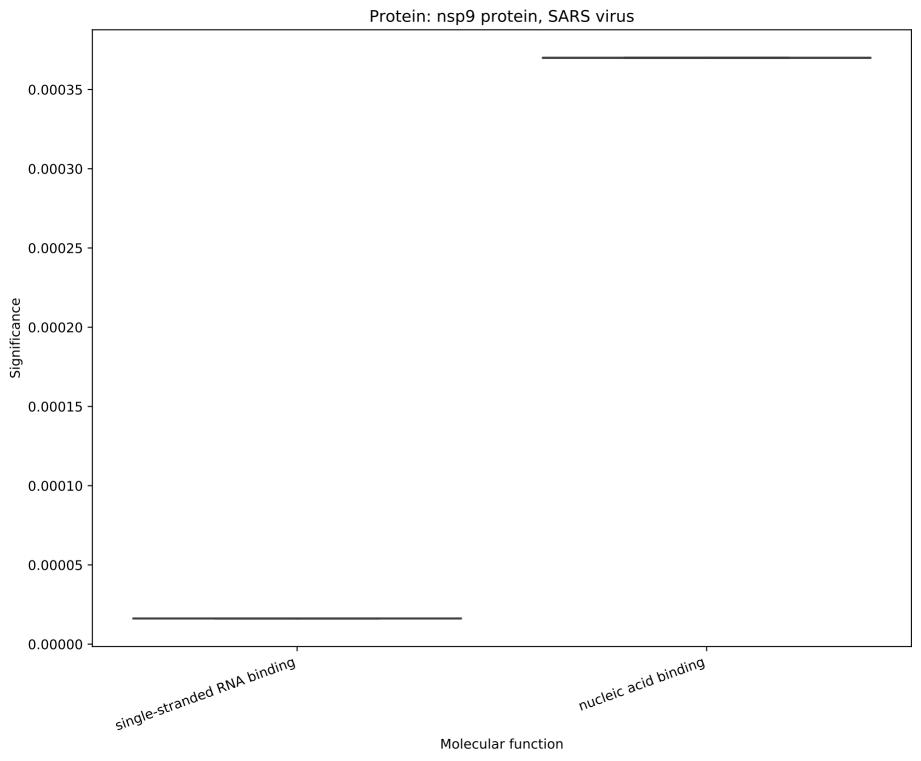


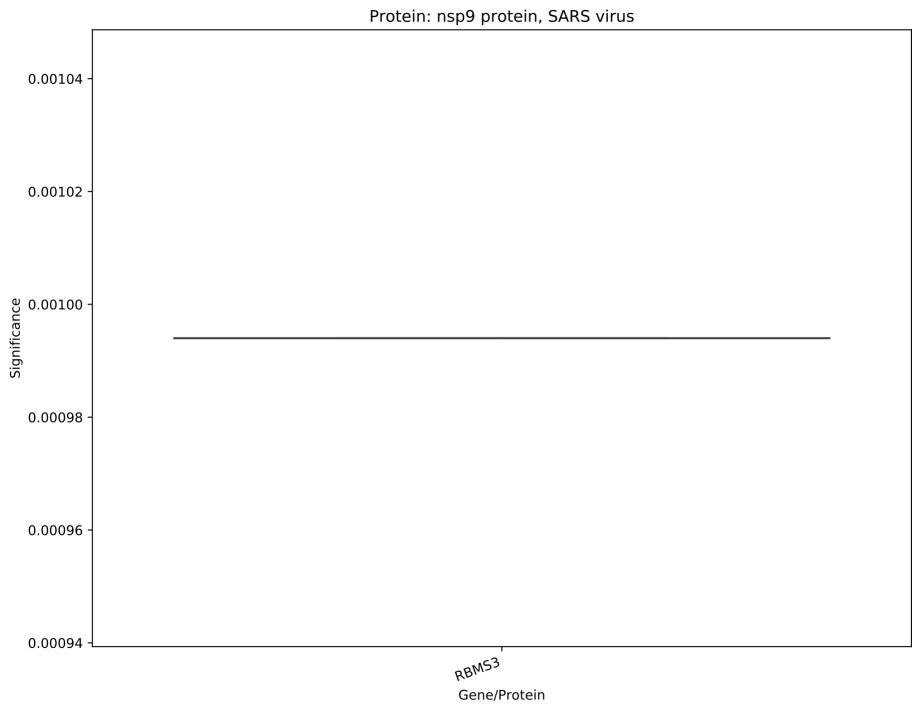


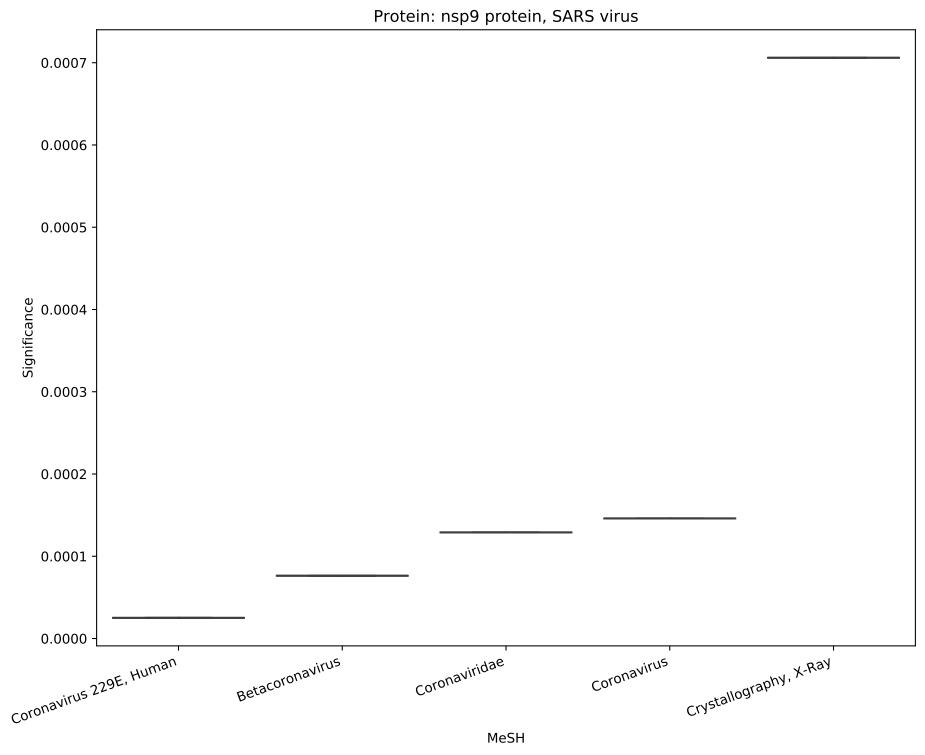


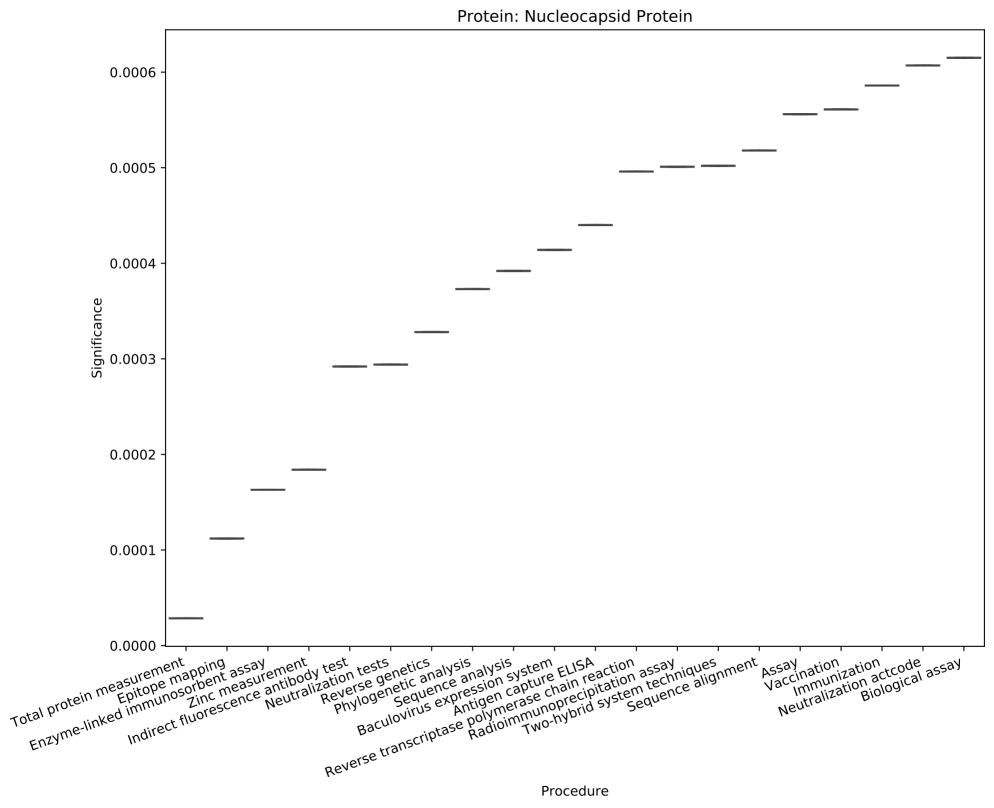


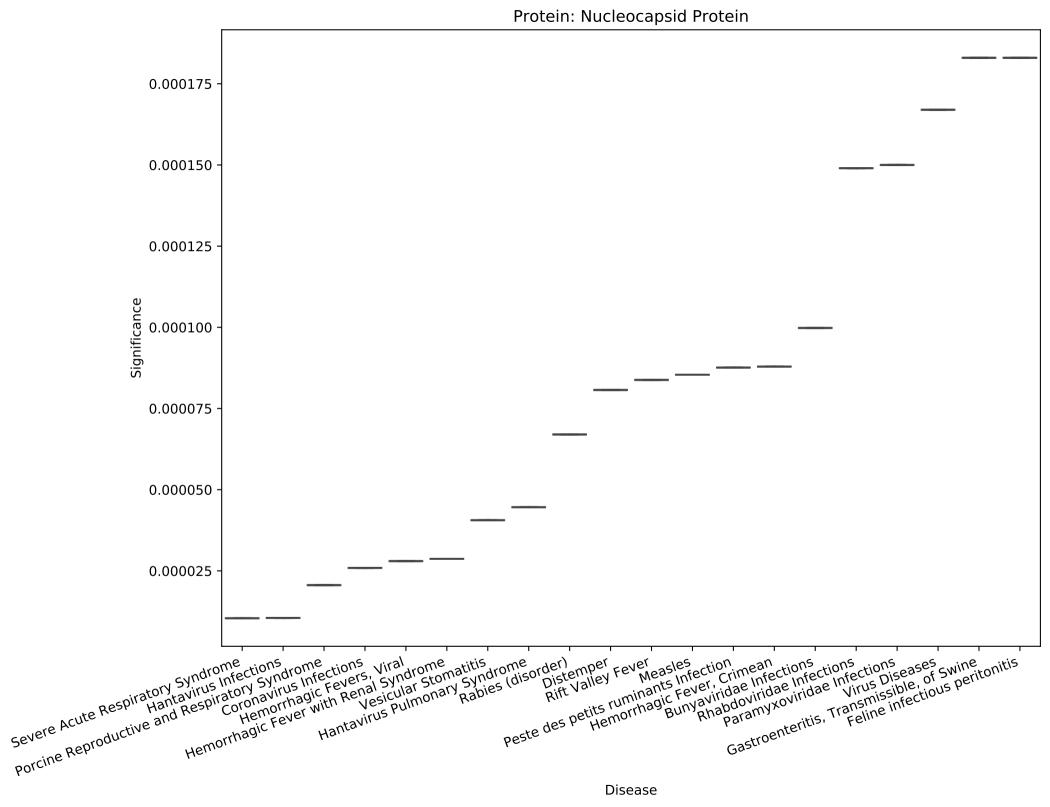
Cellular component

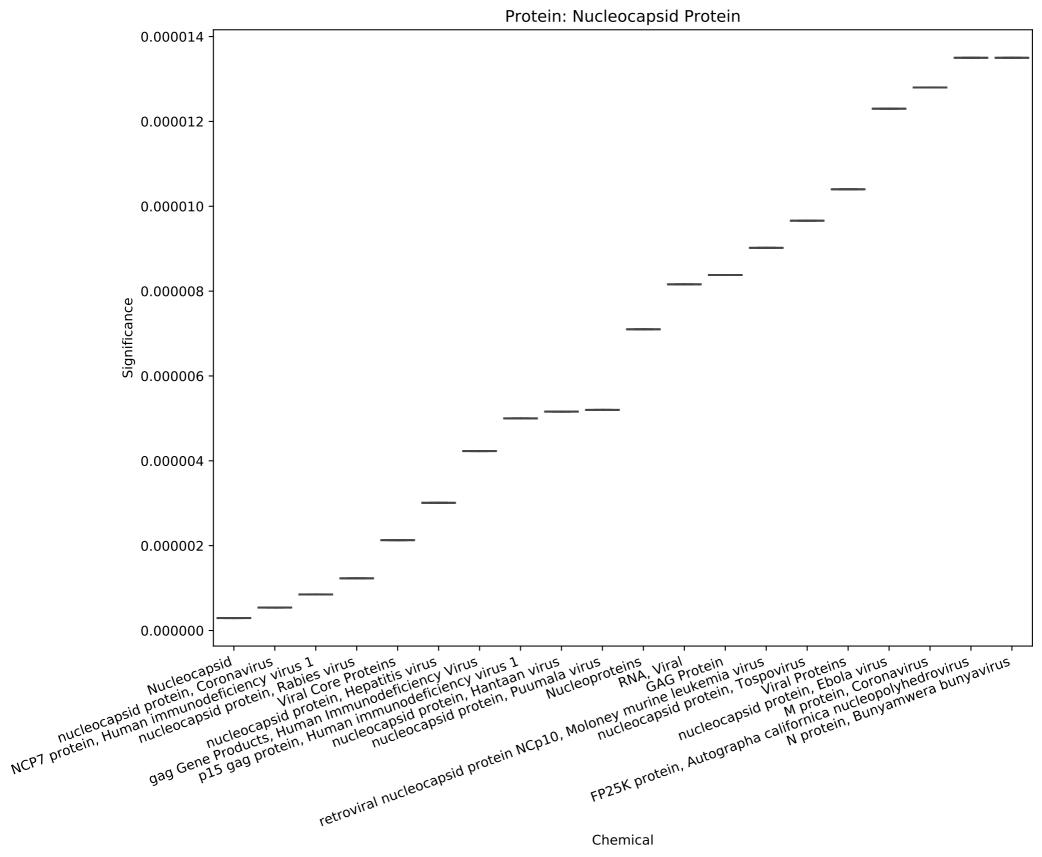


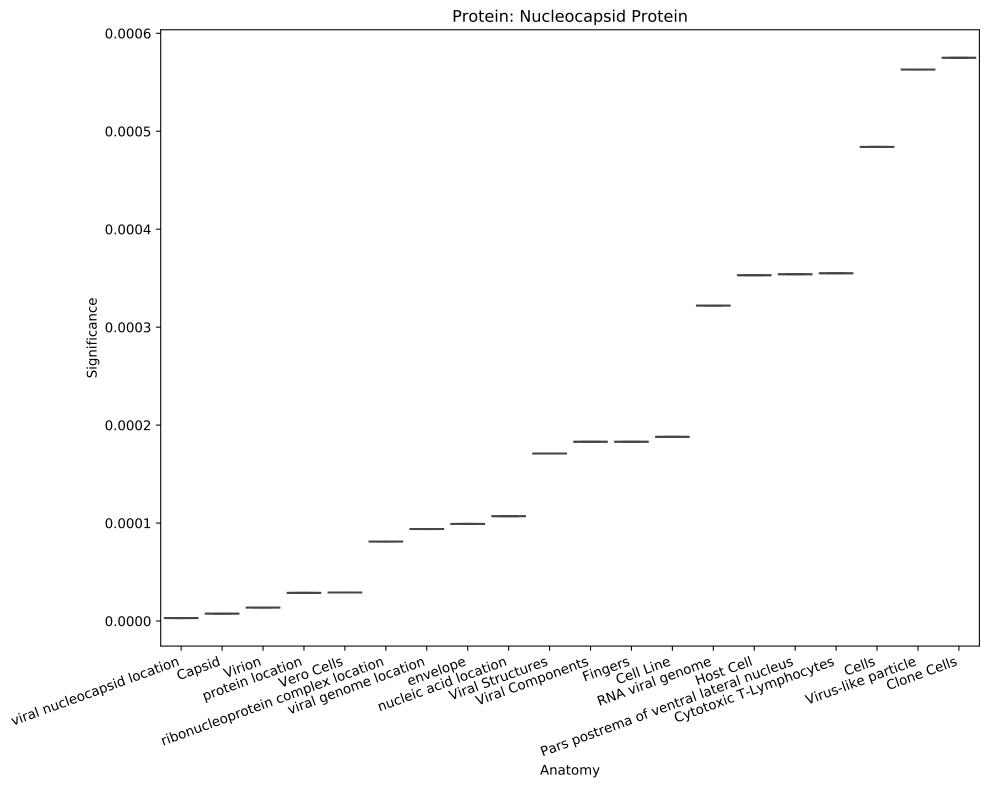


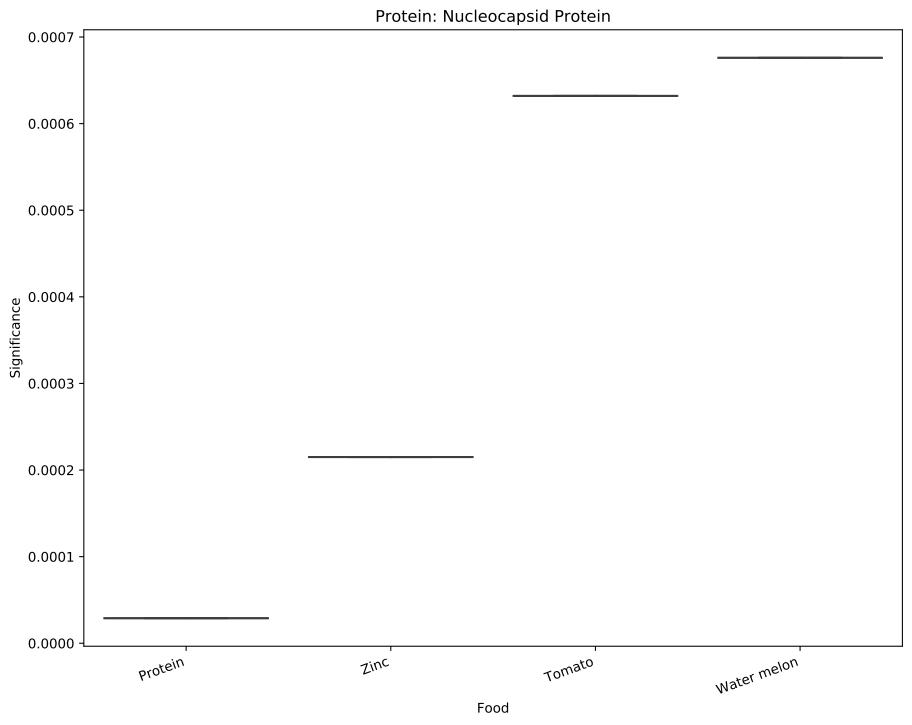


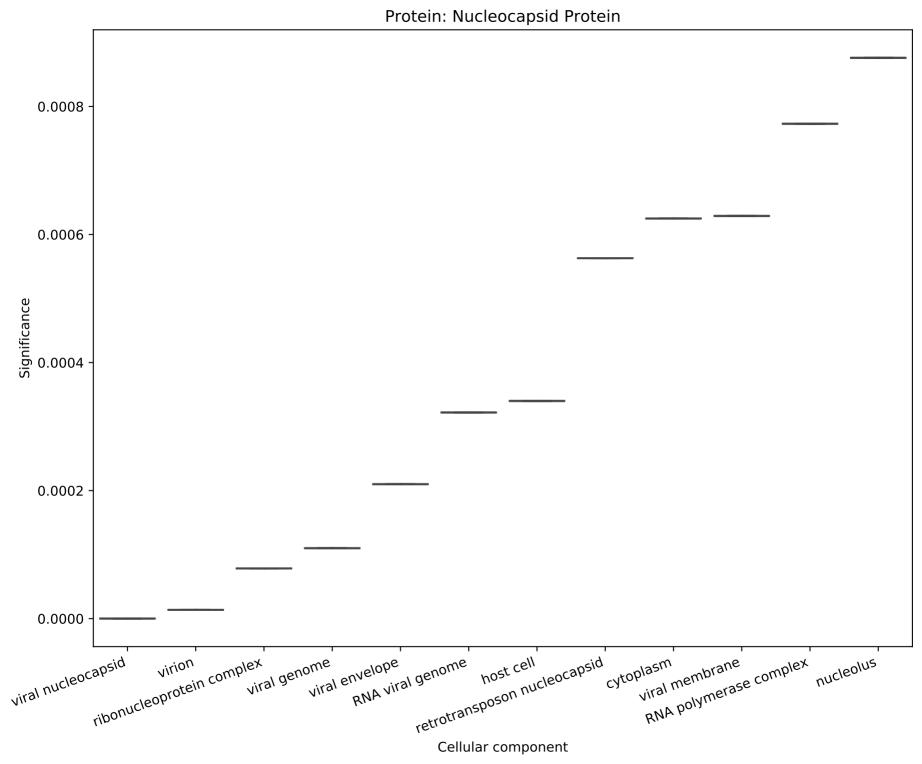


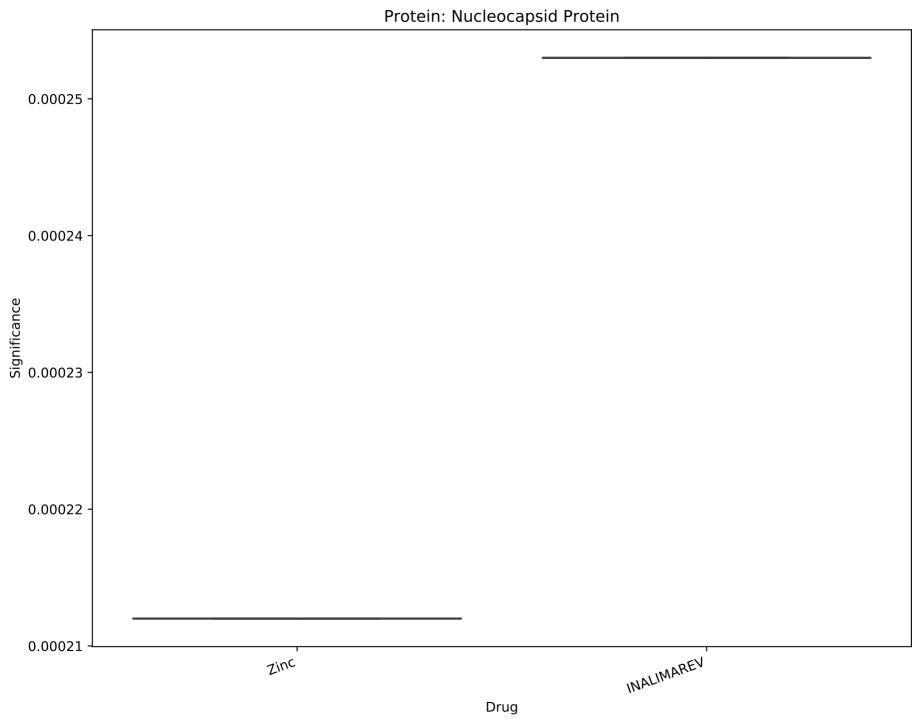


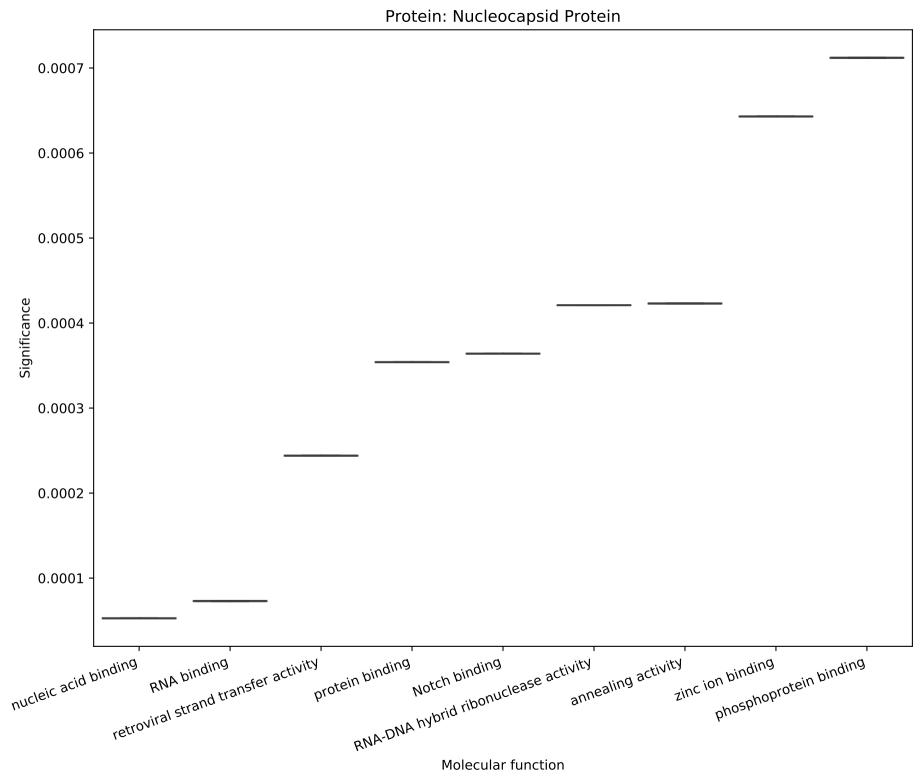


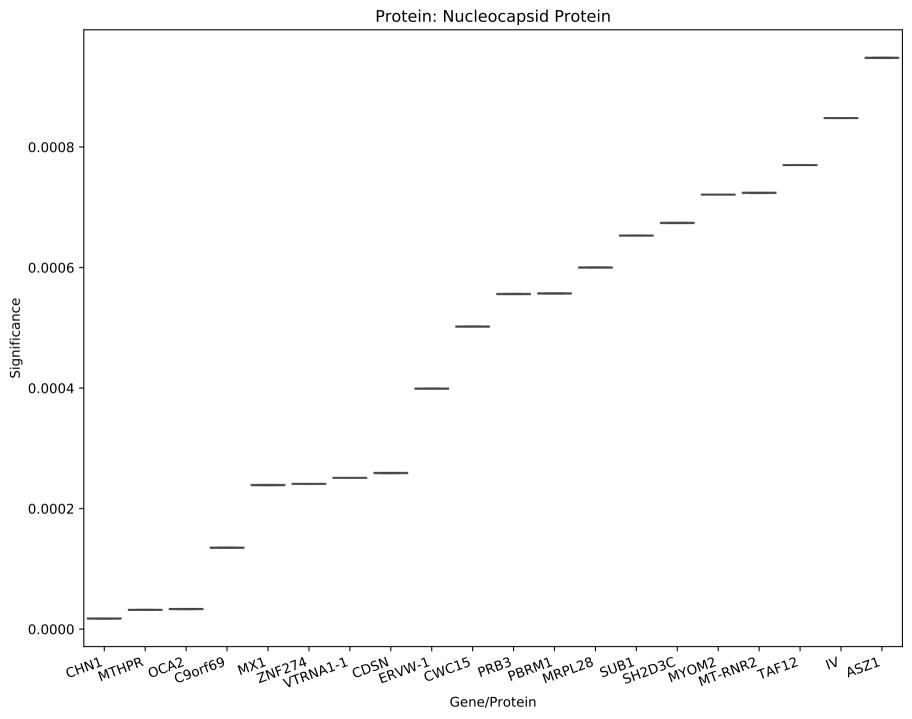


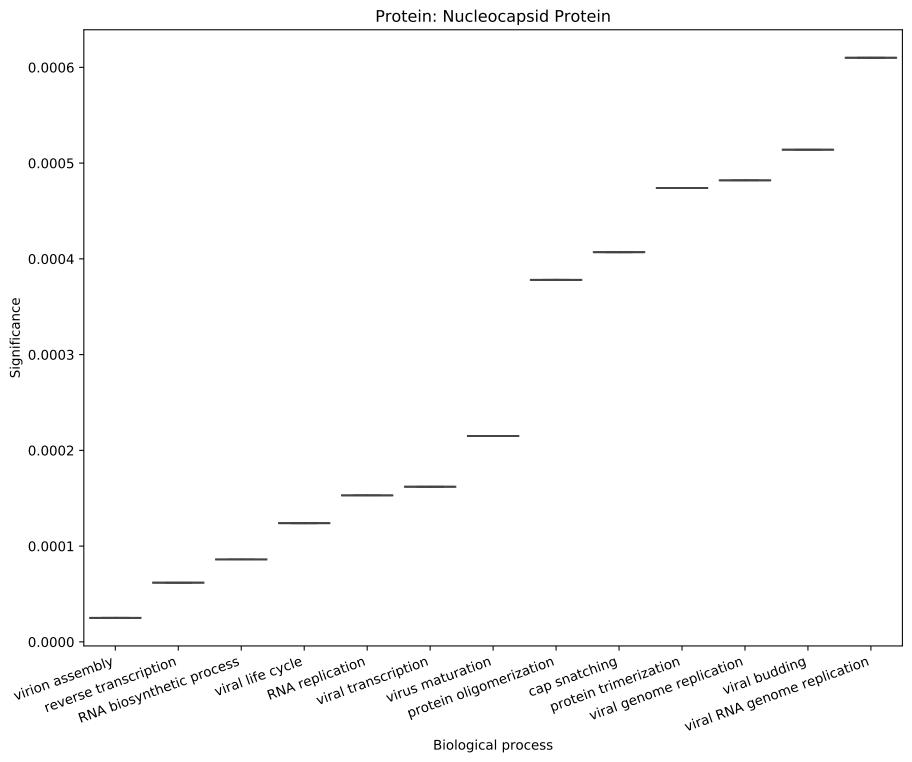


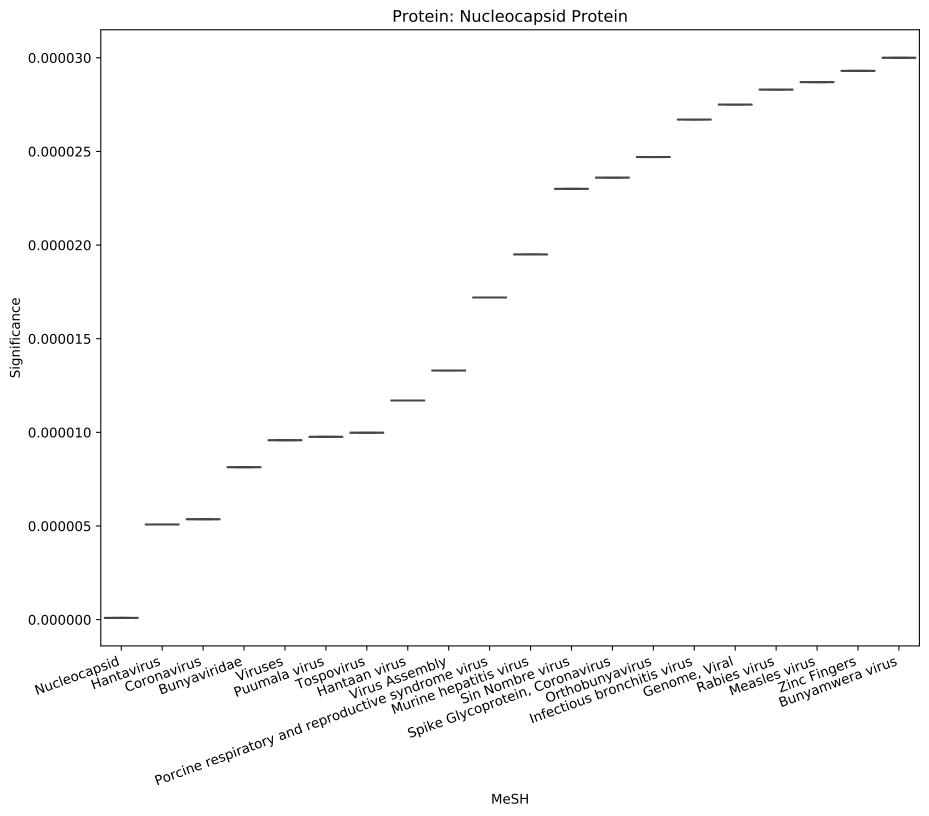


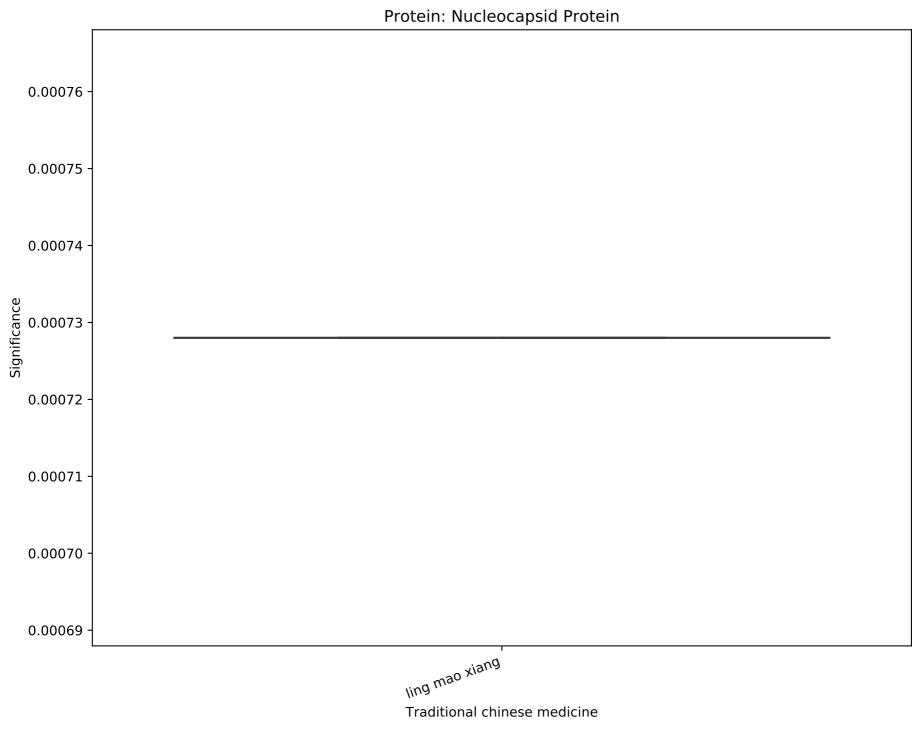


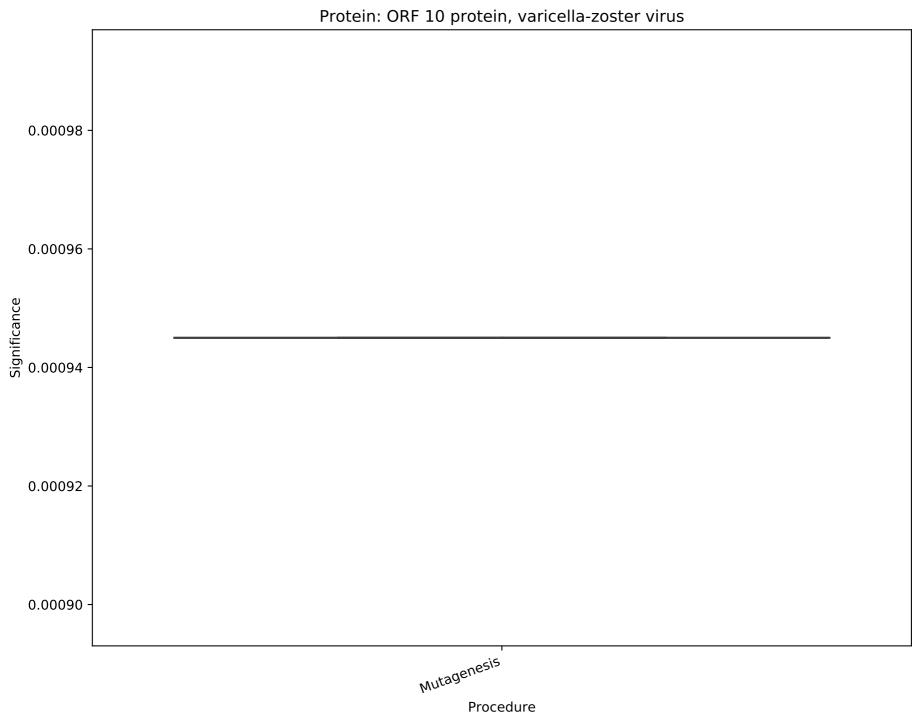


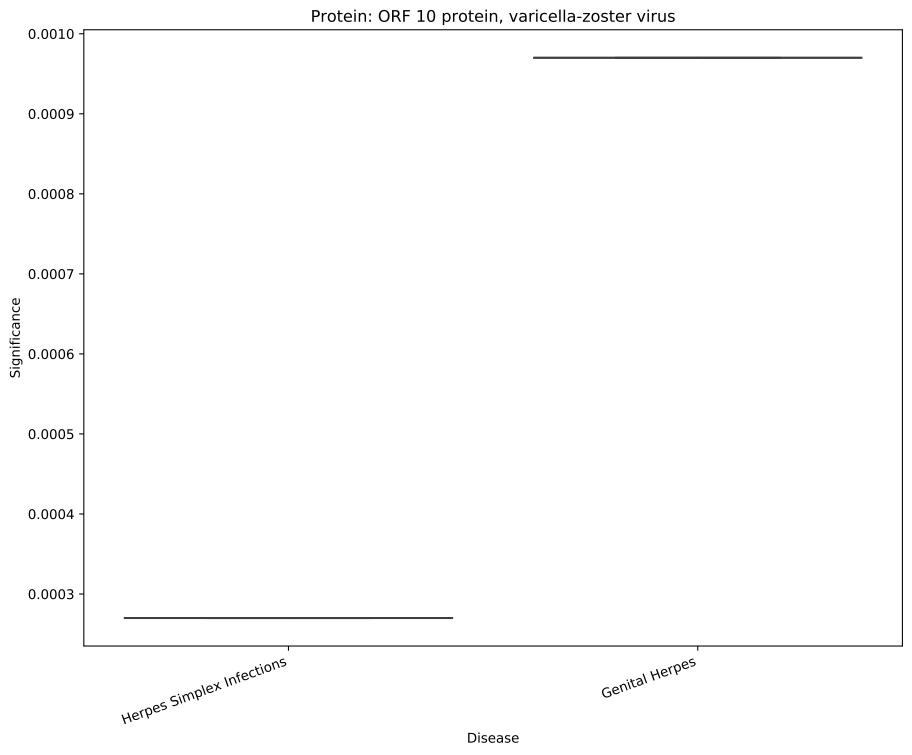


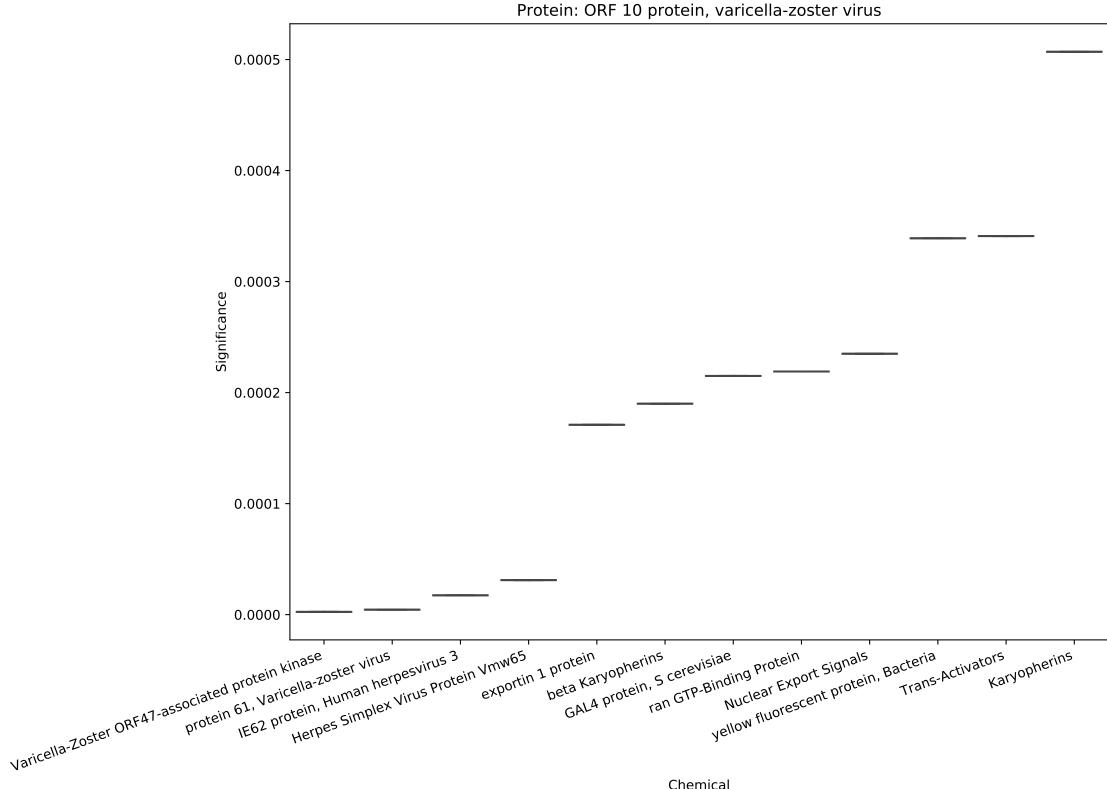


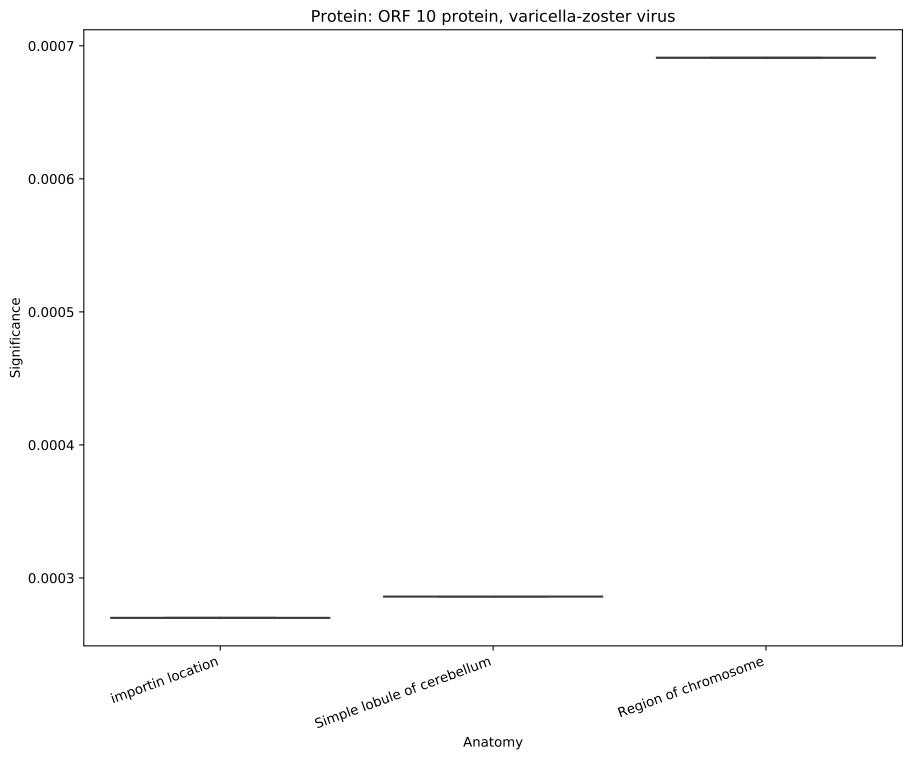


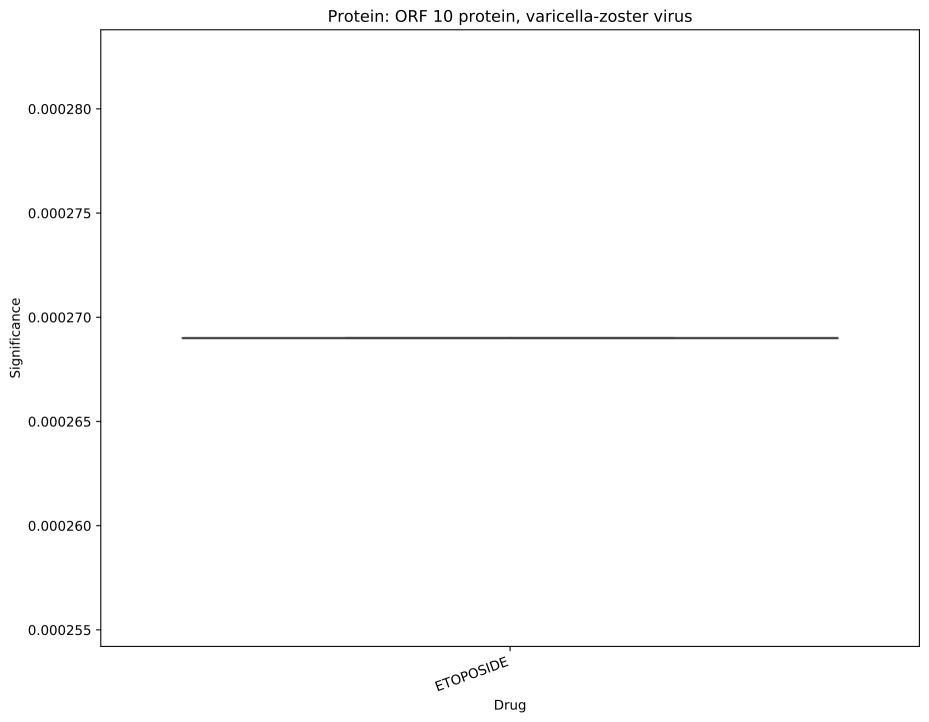


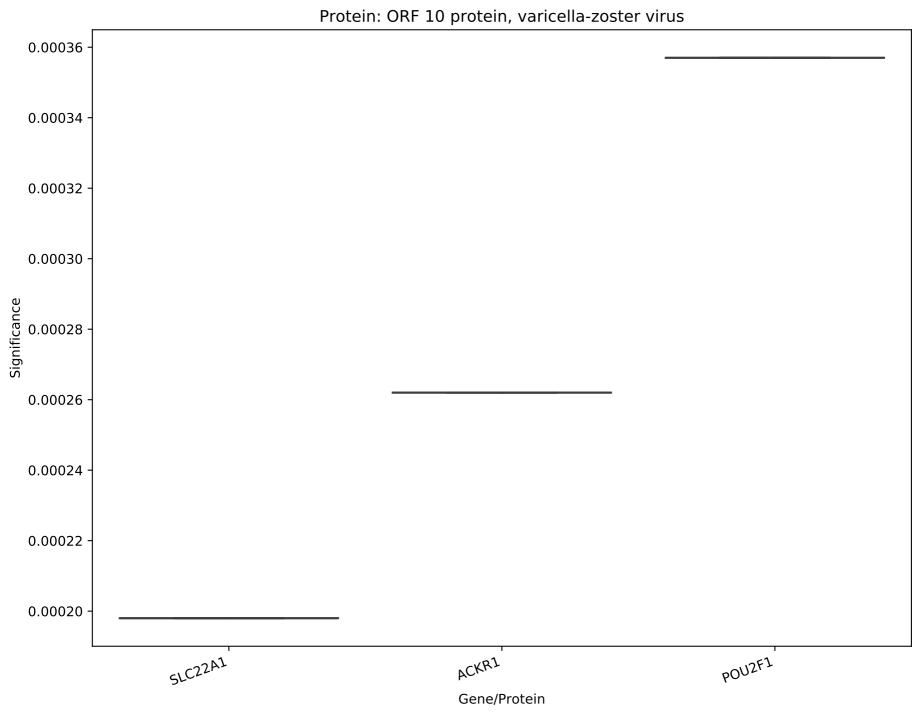


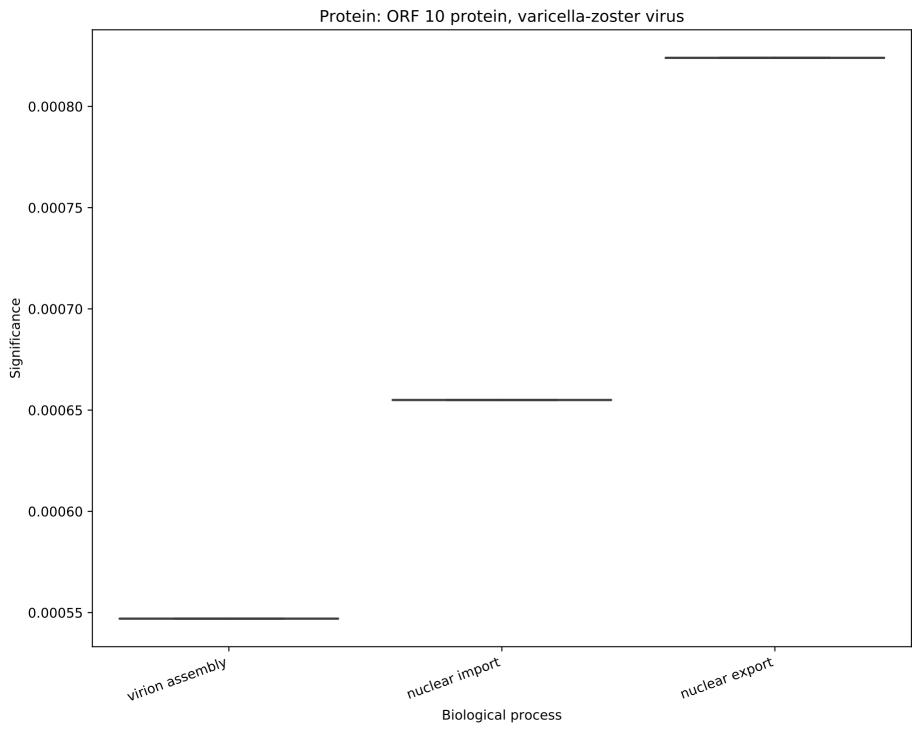


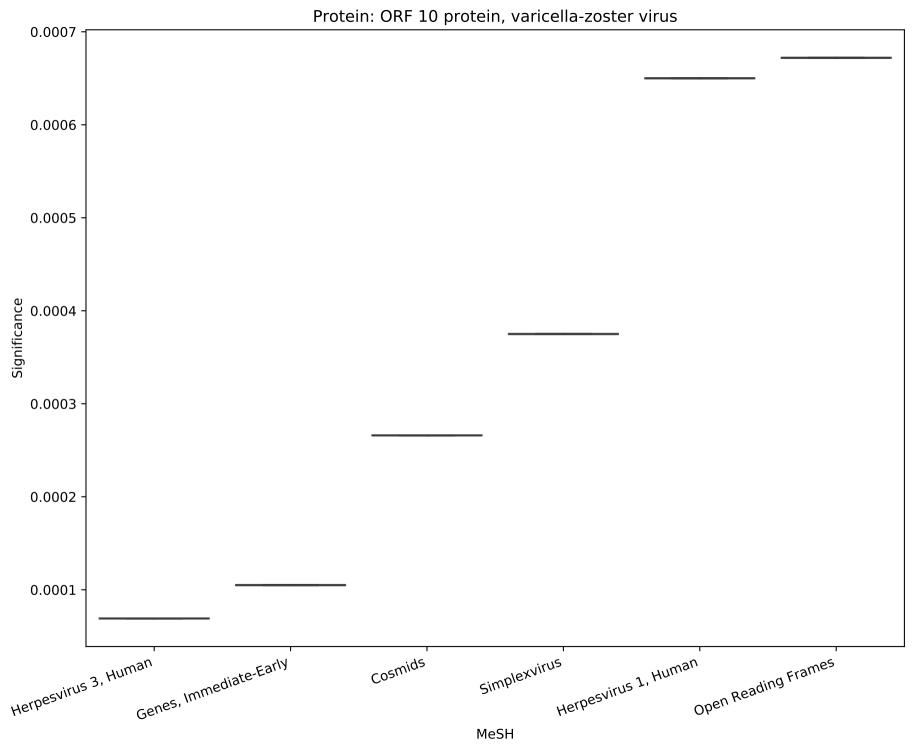


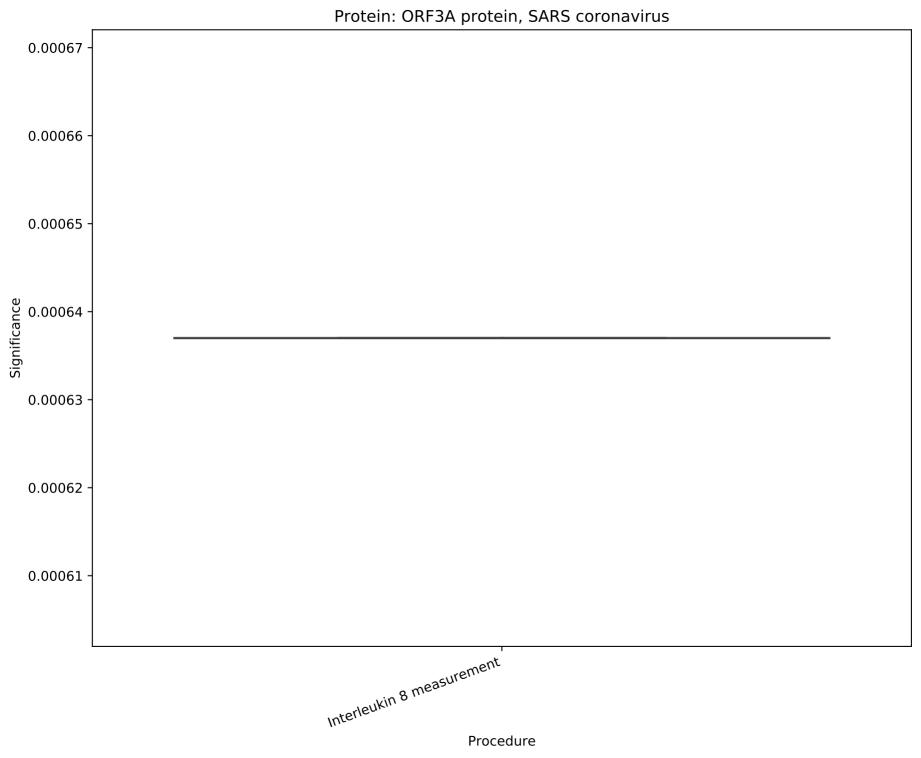


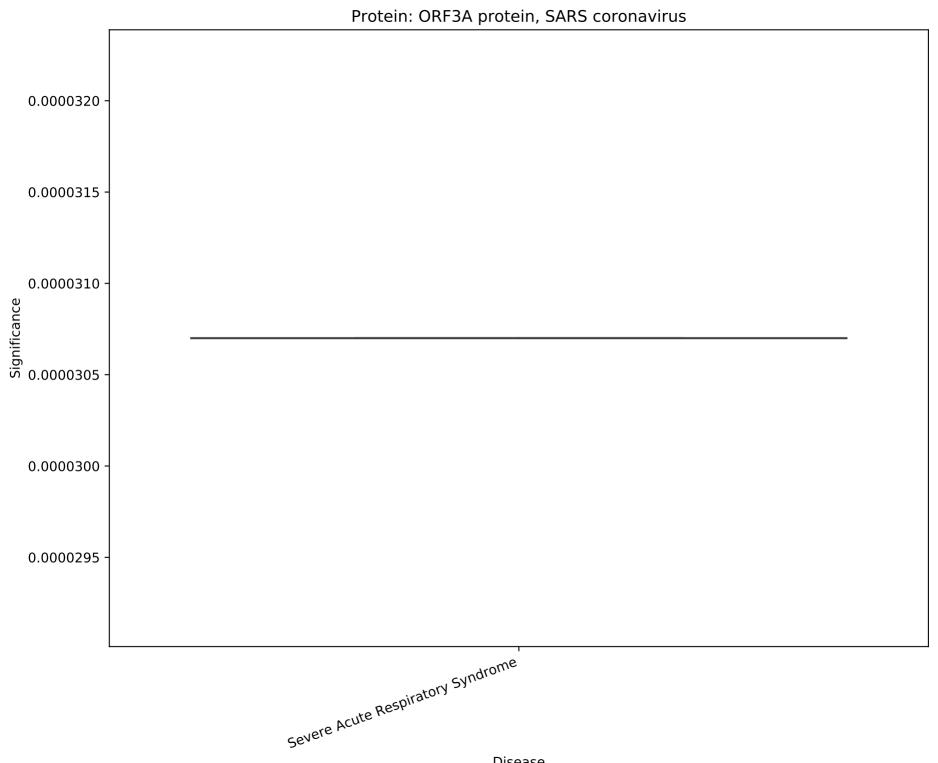


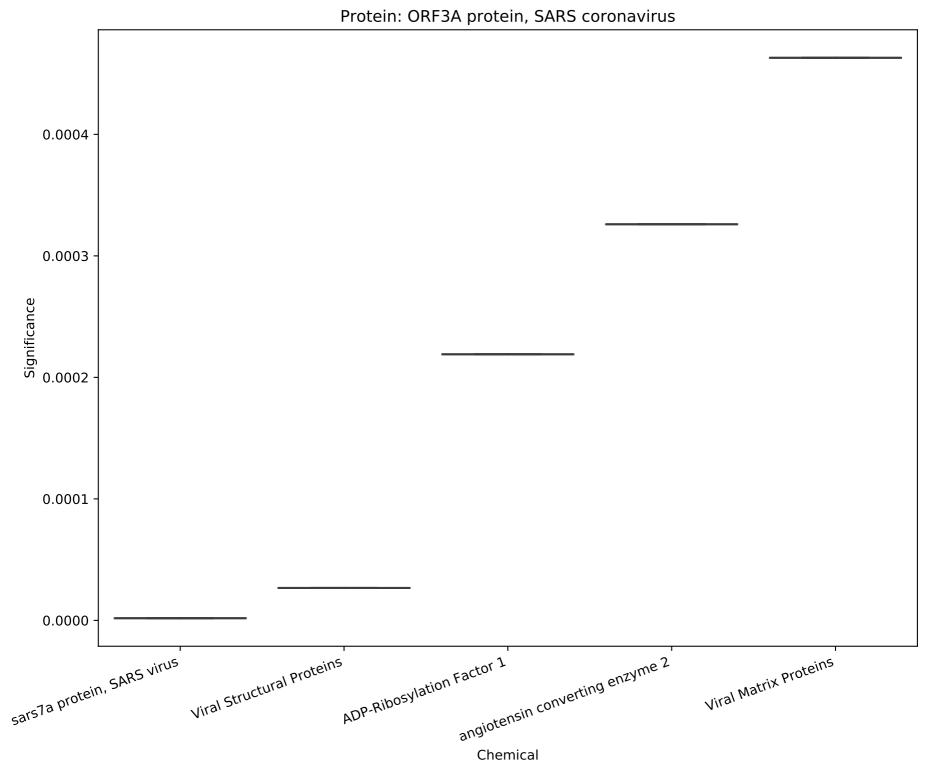


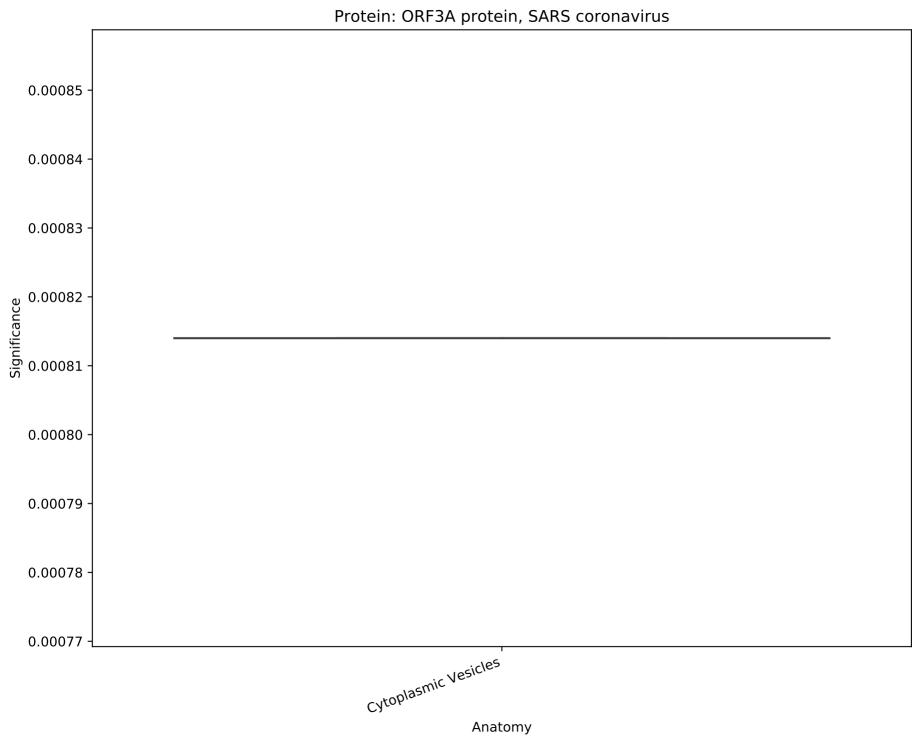


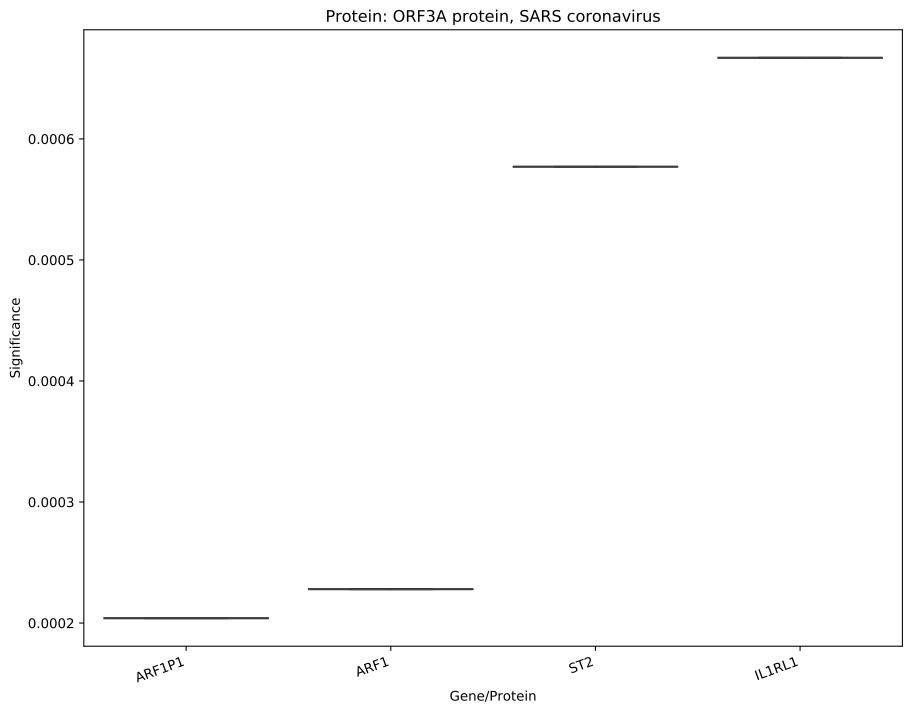


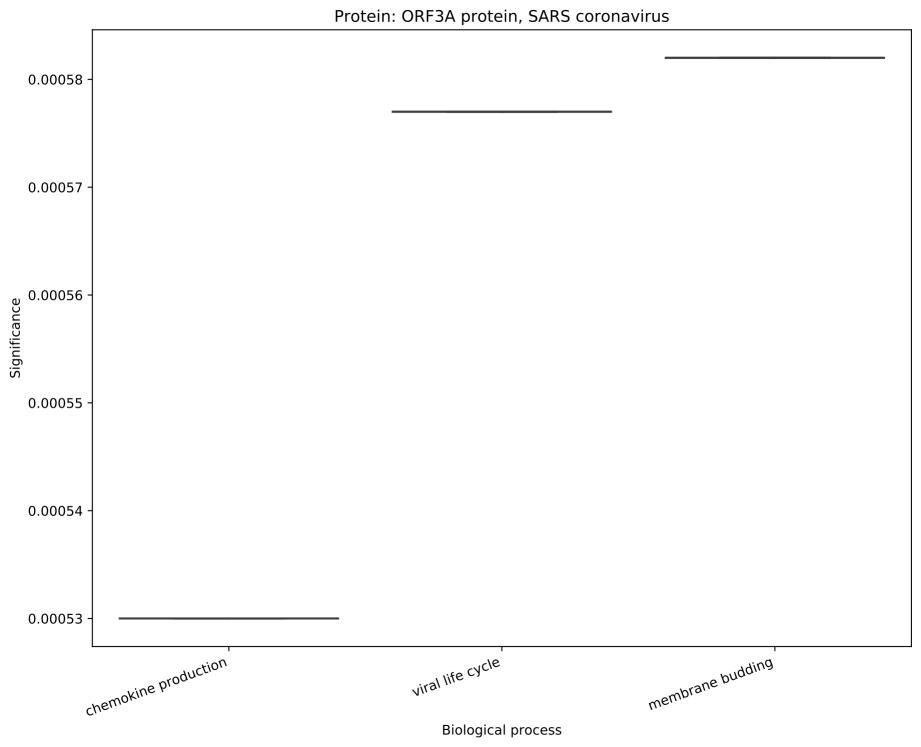


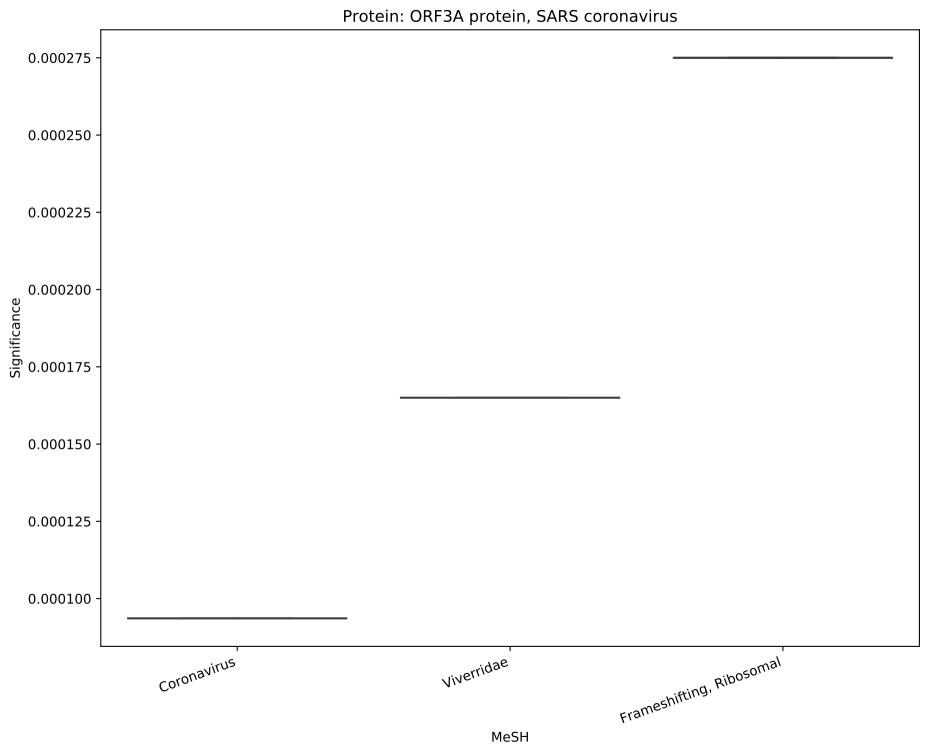


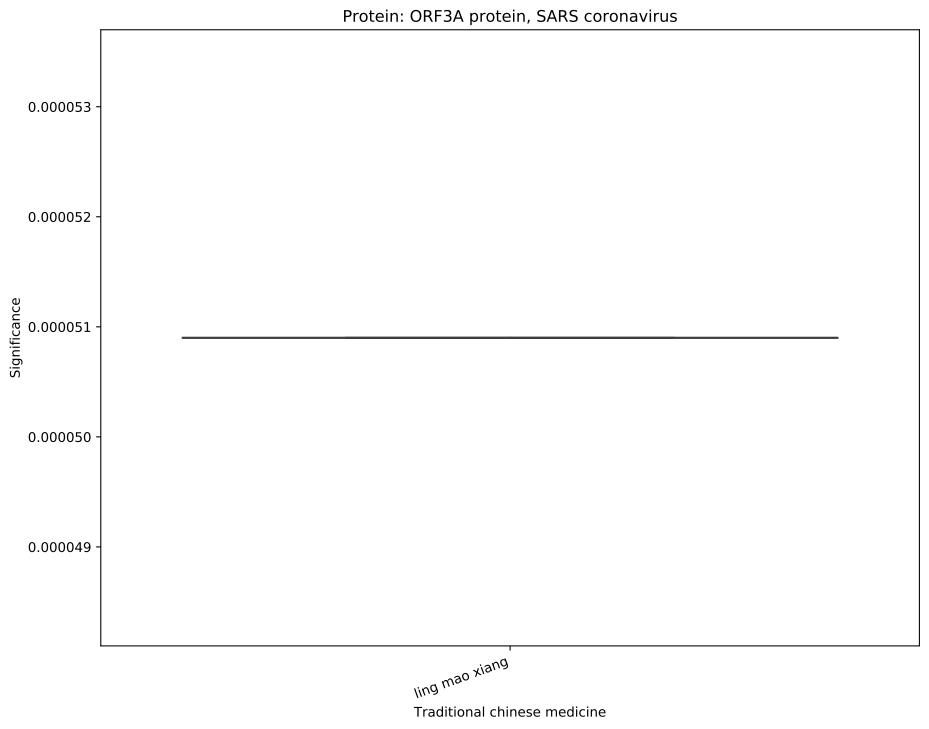


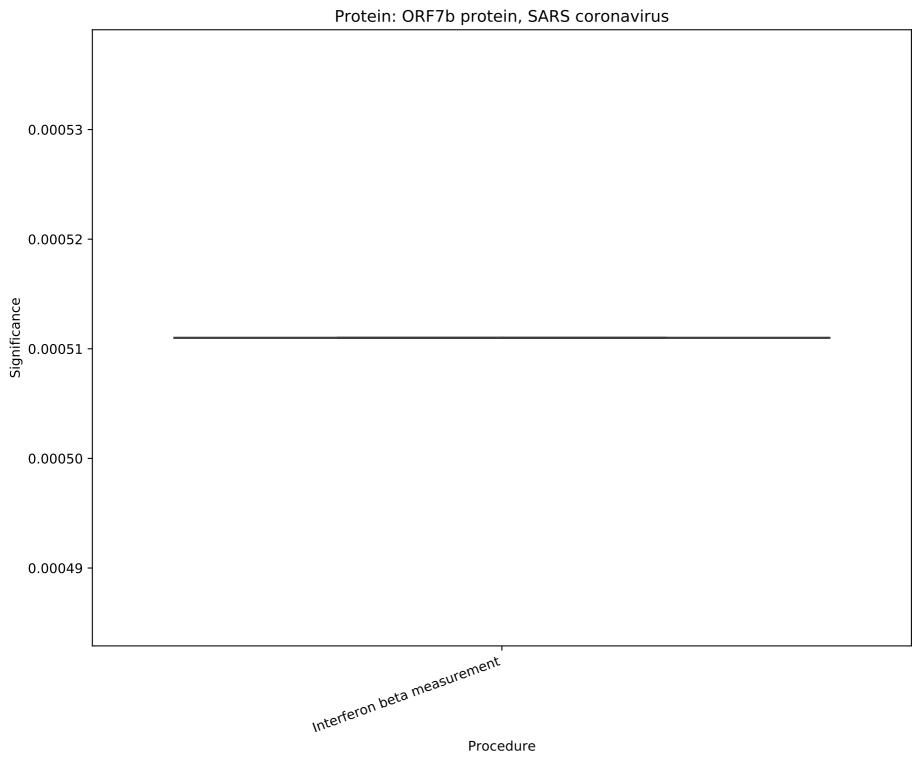


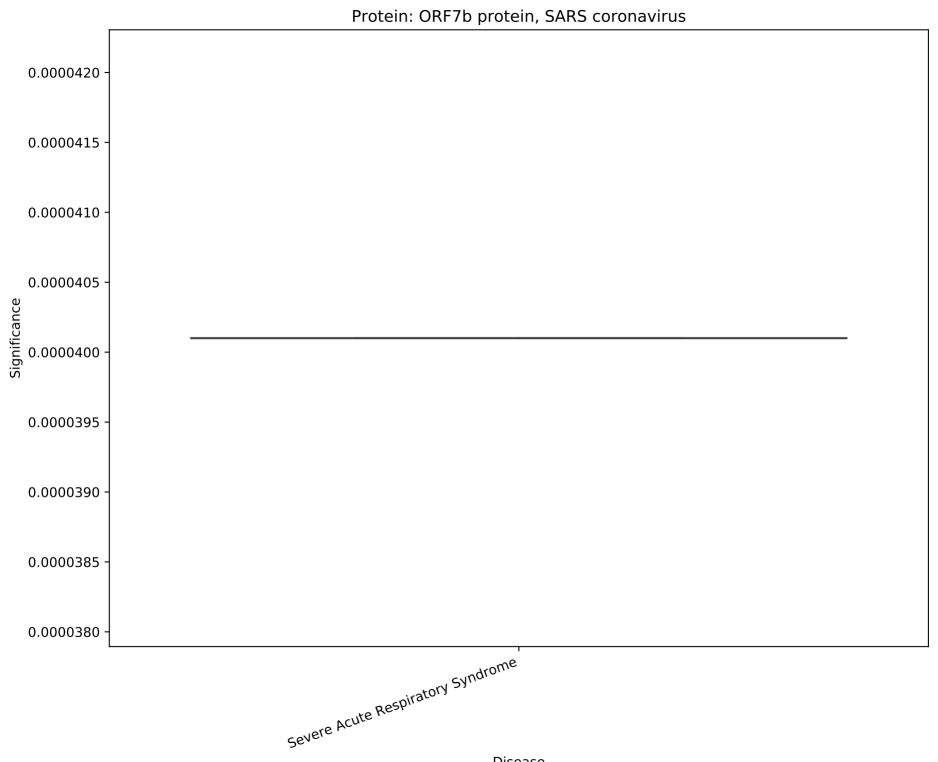


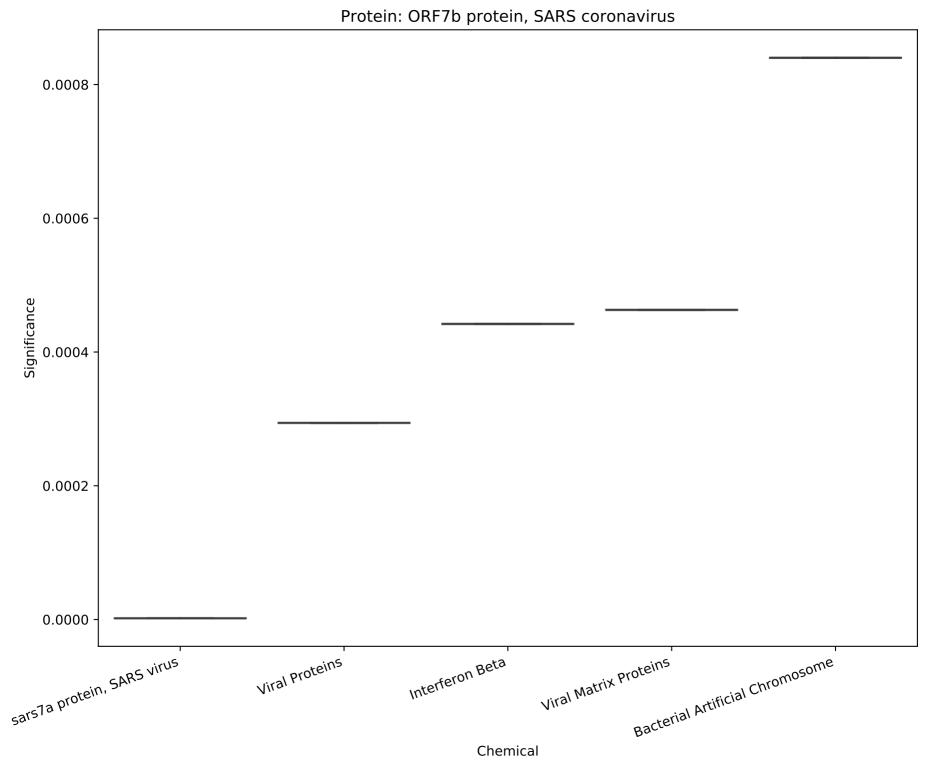


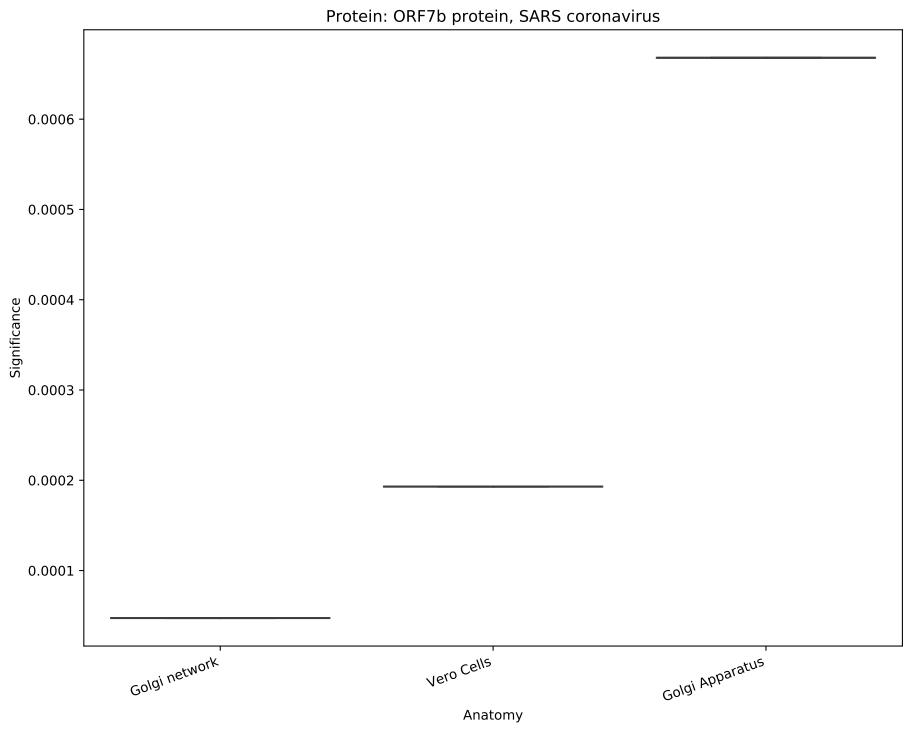


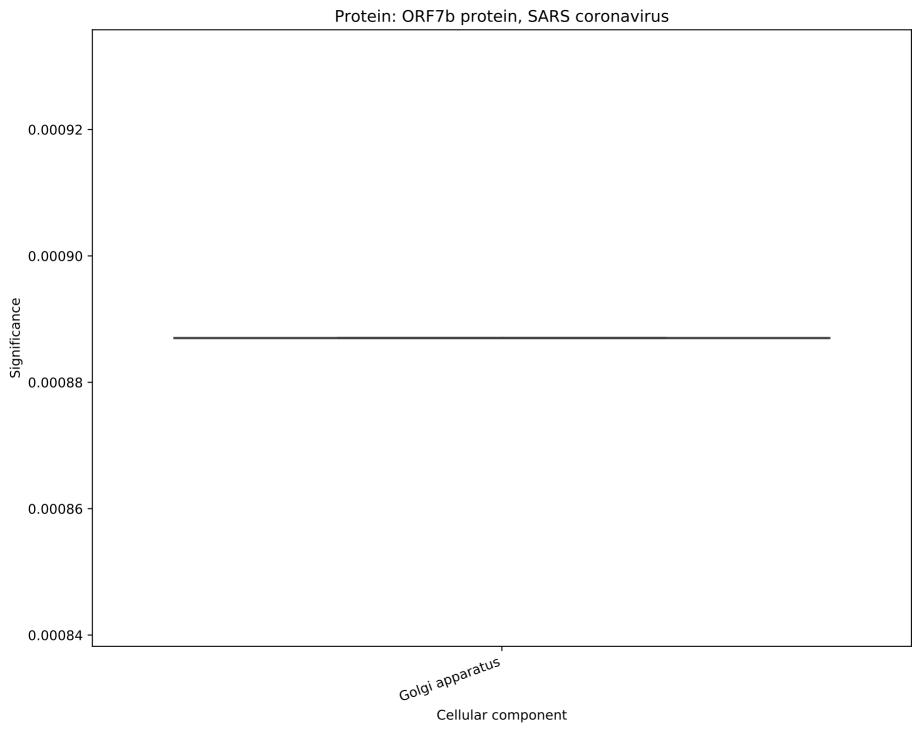


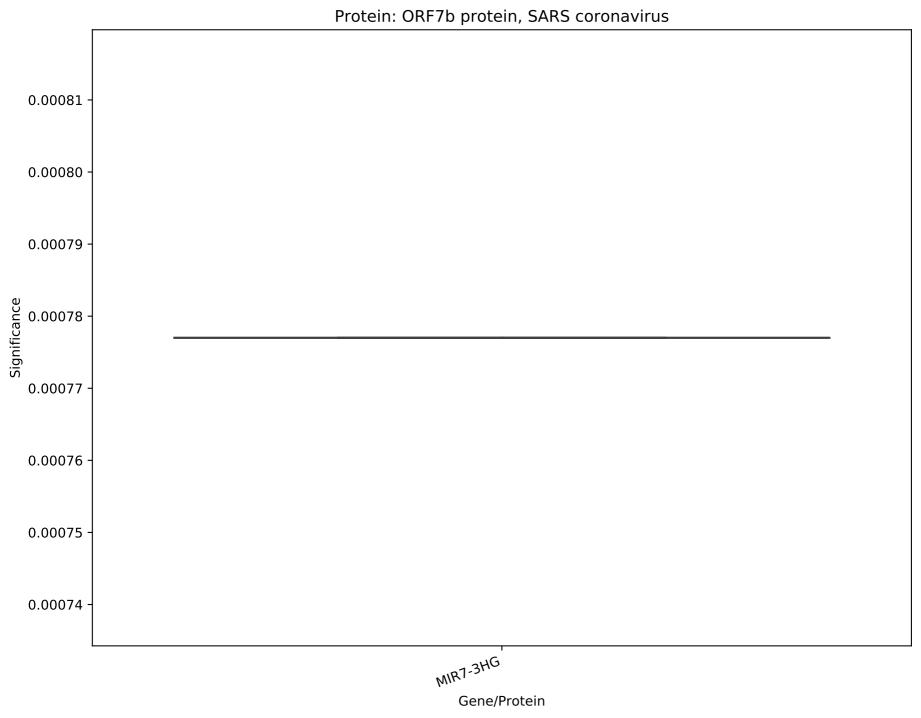


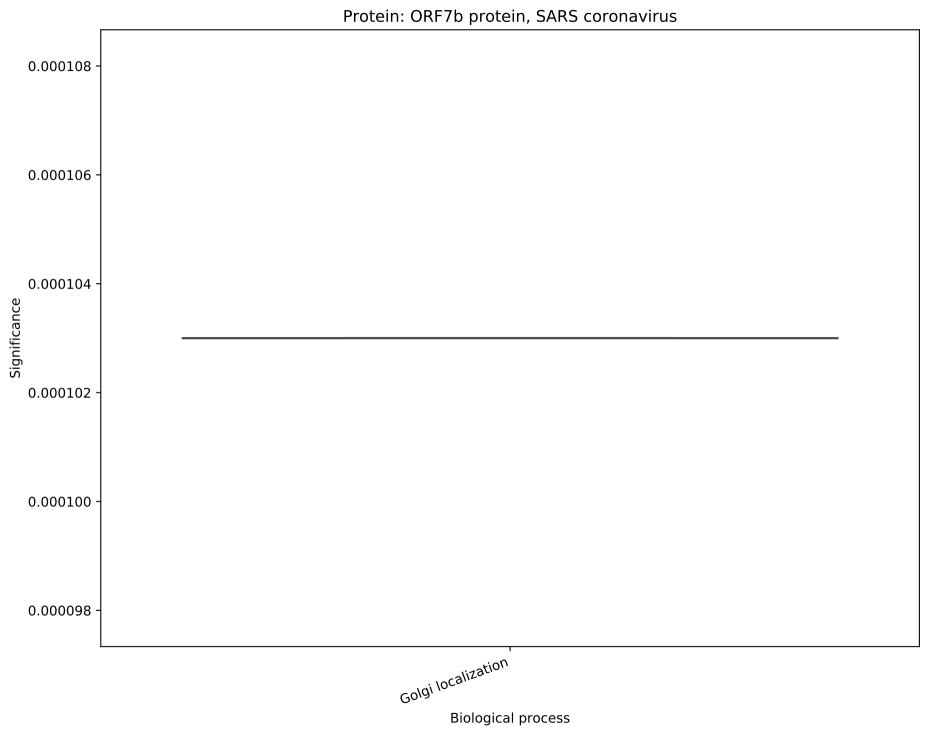


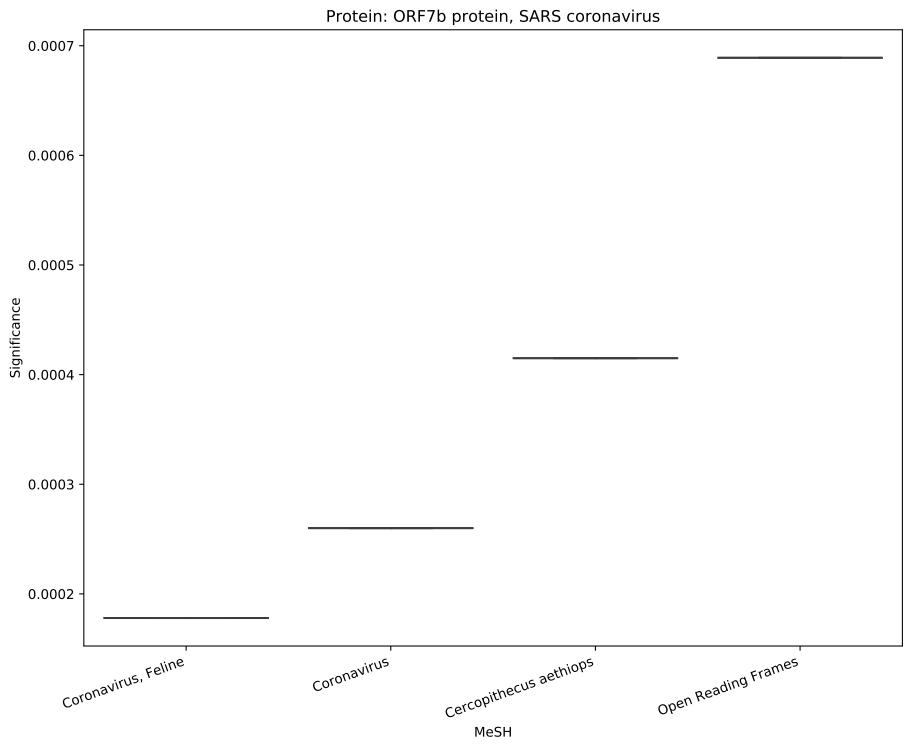


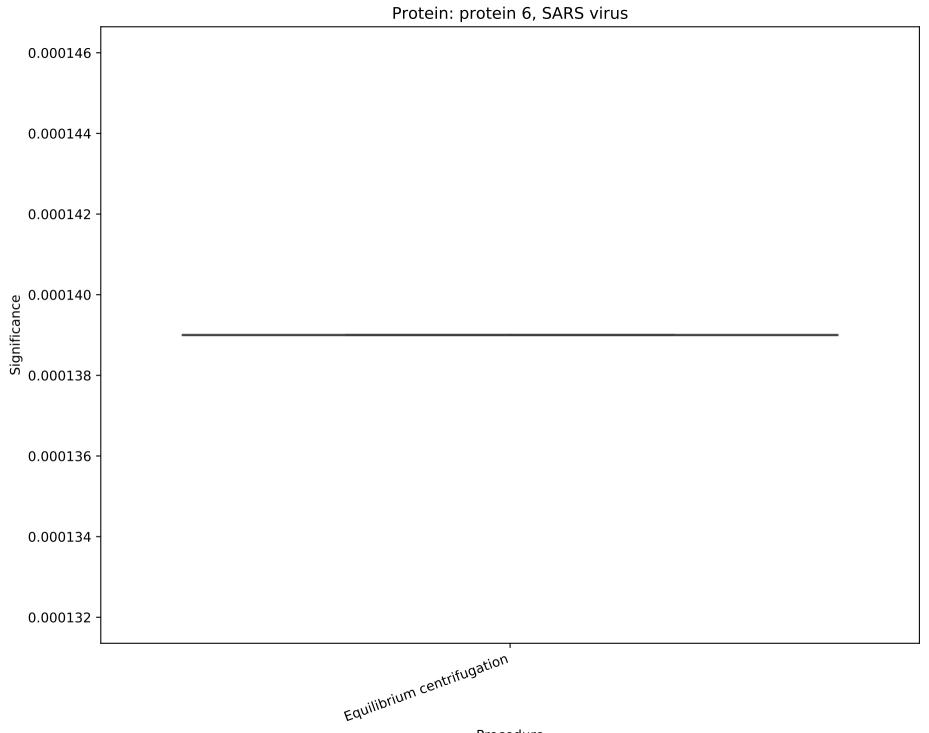




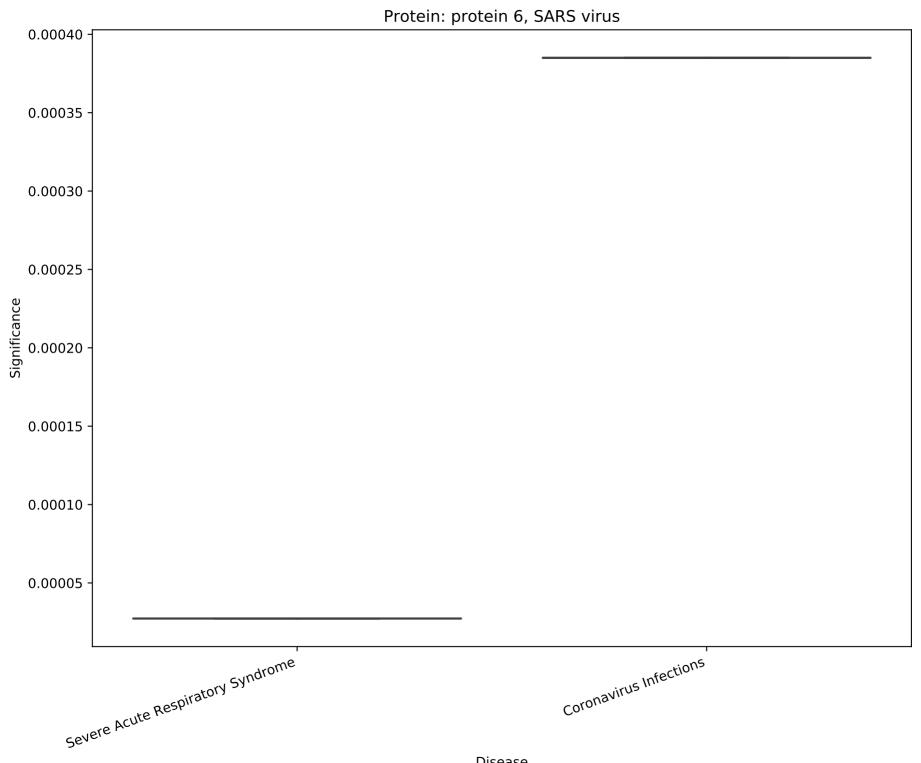


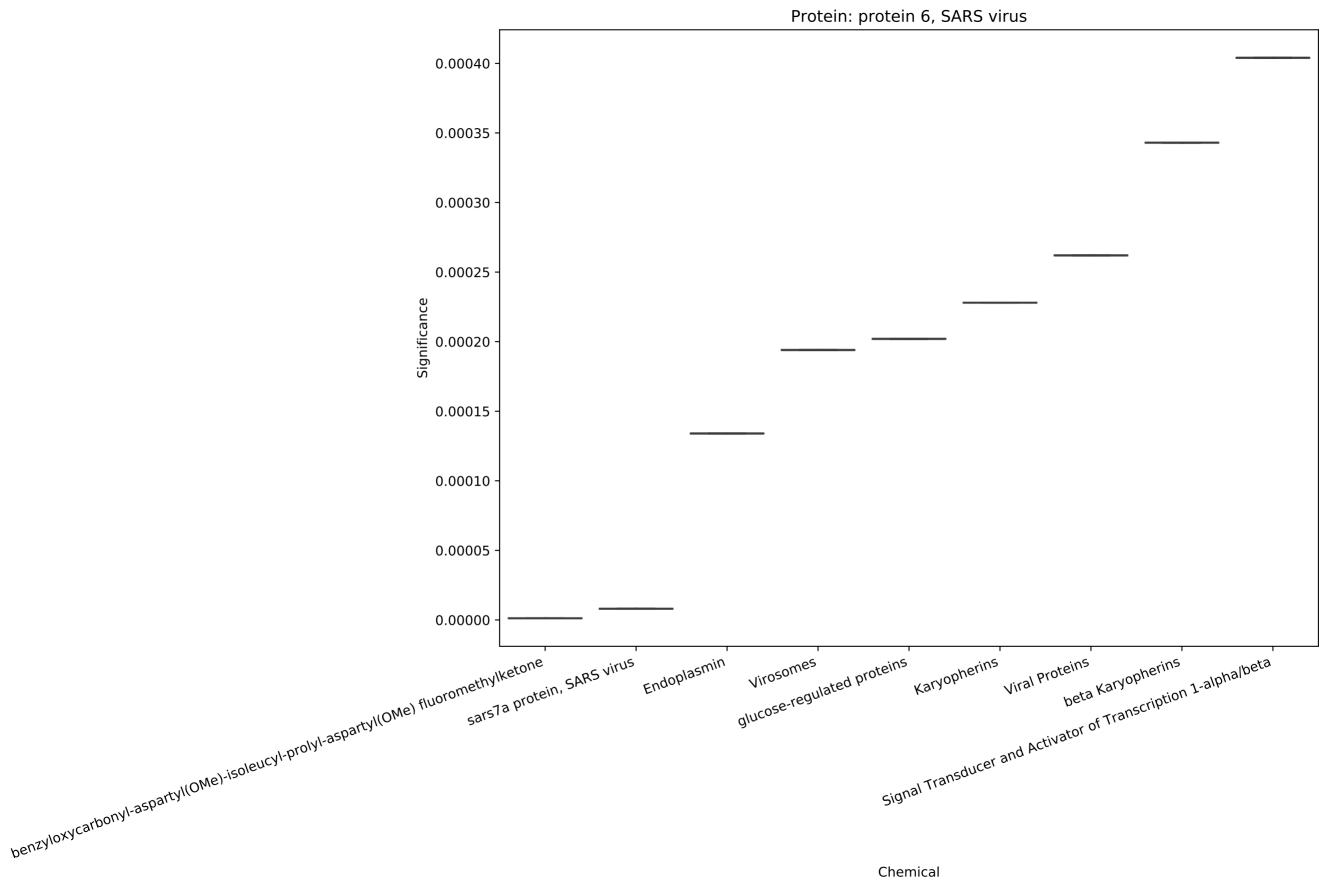


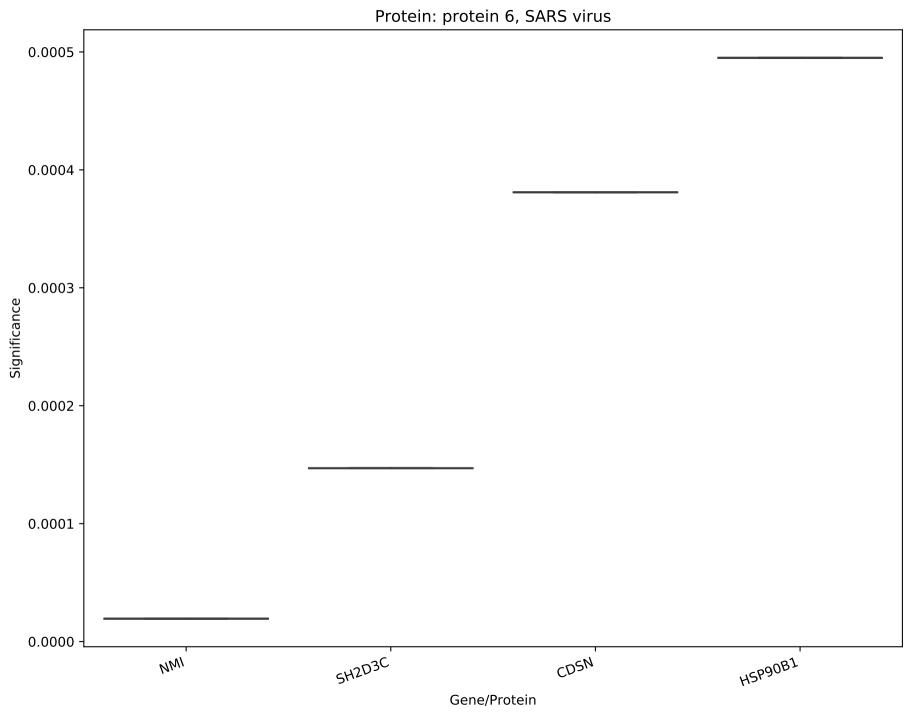


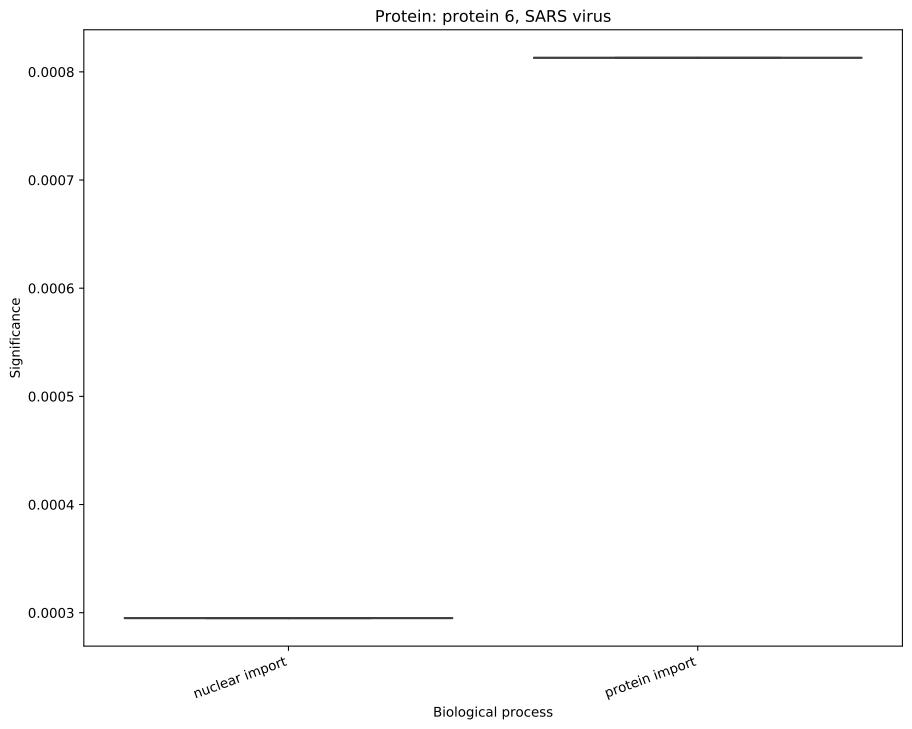


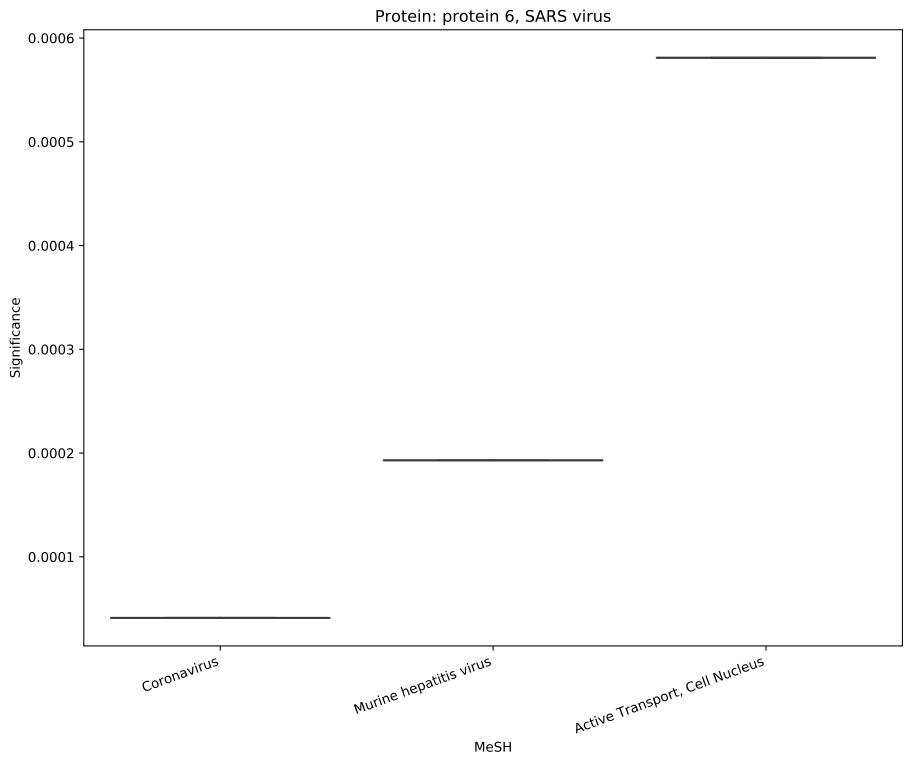
Procedure

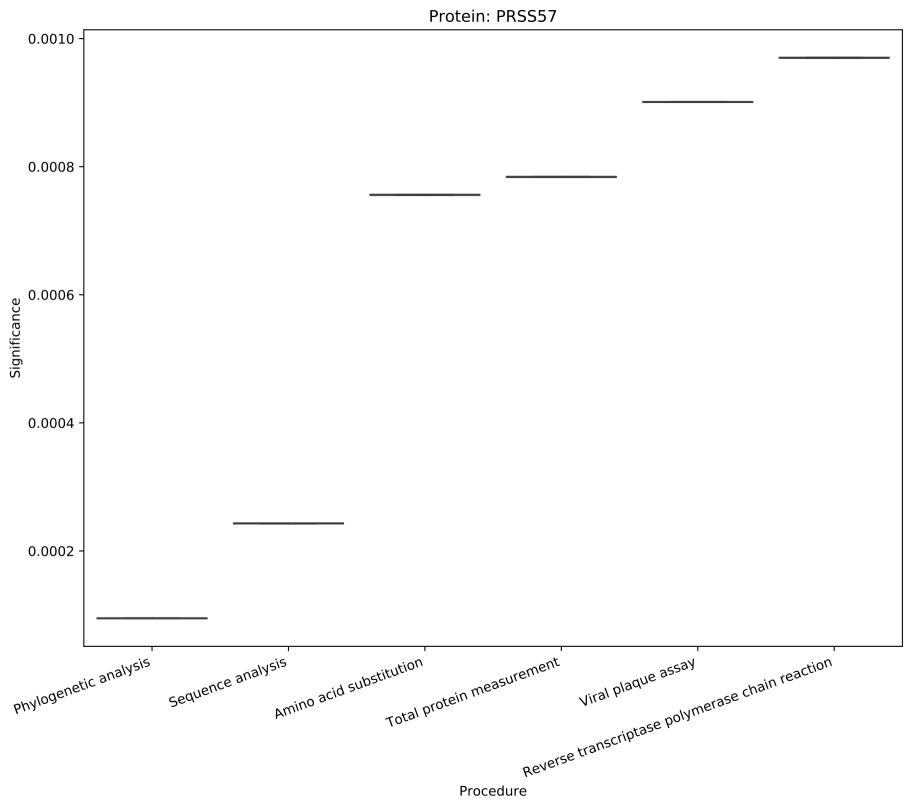


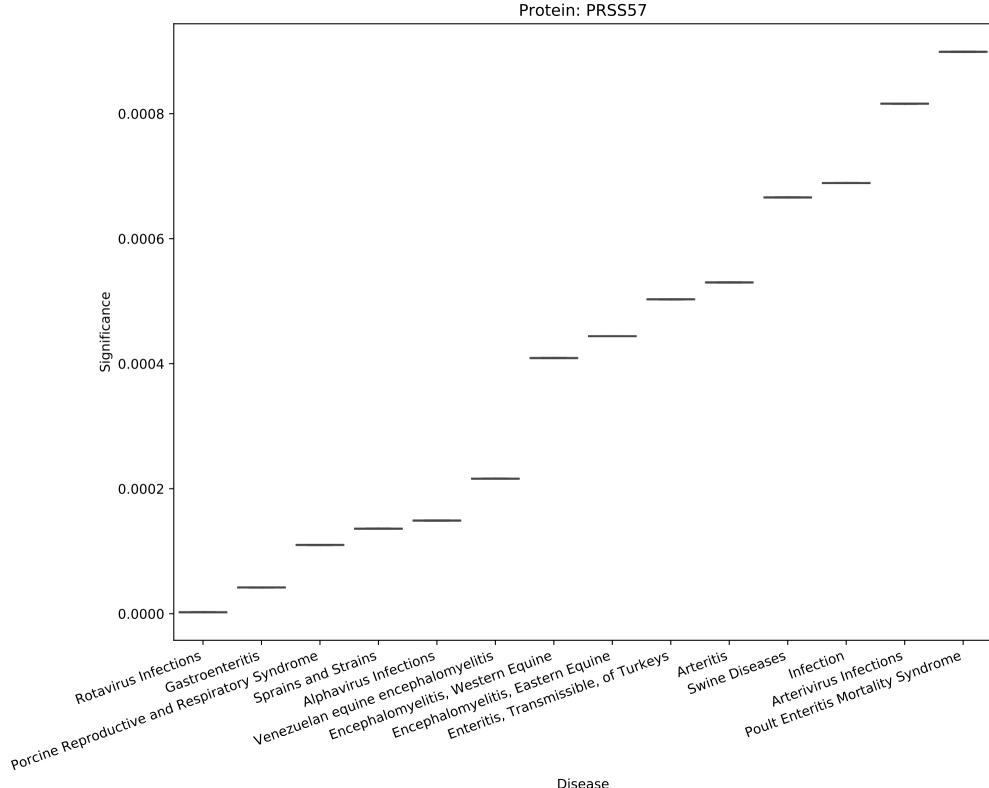


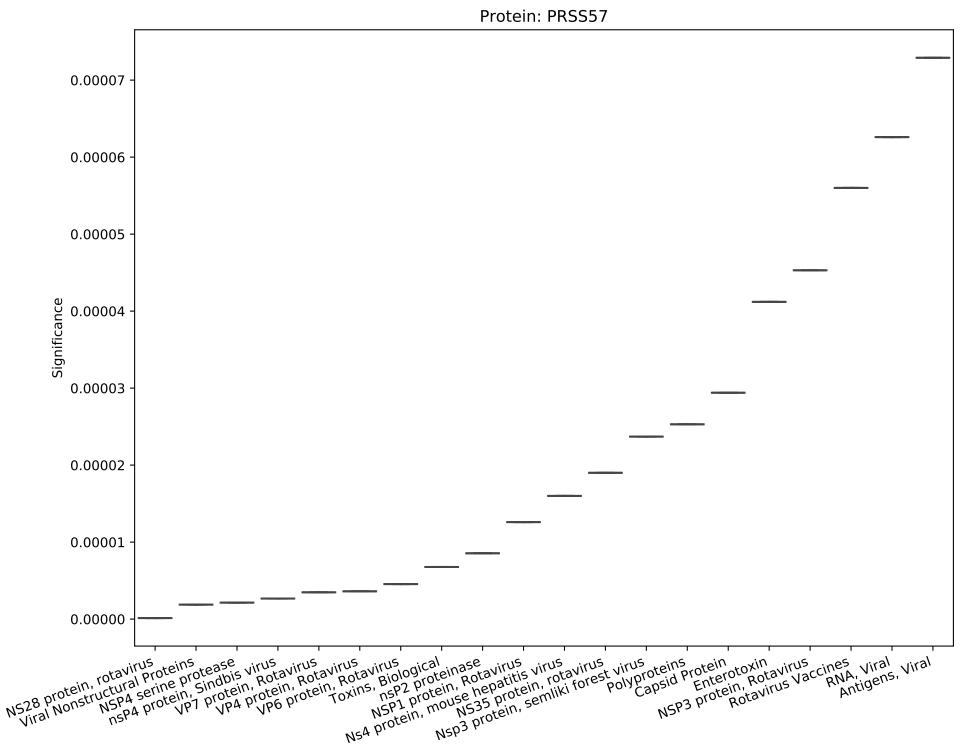




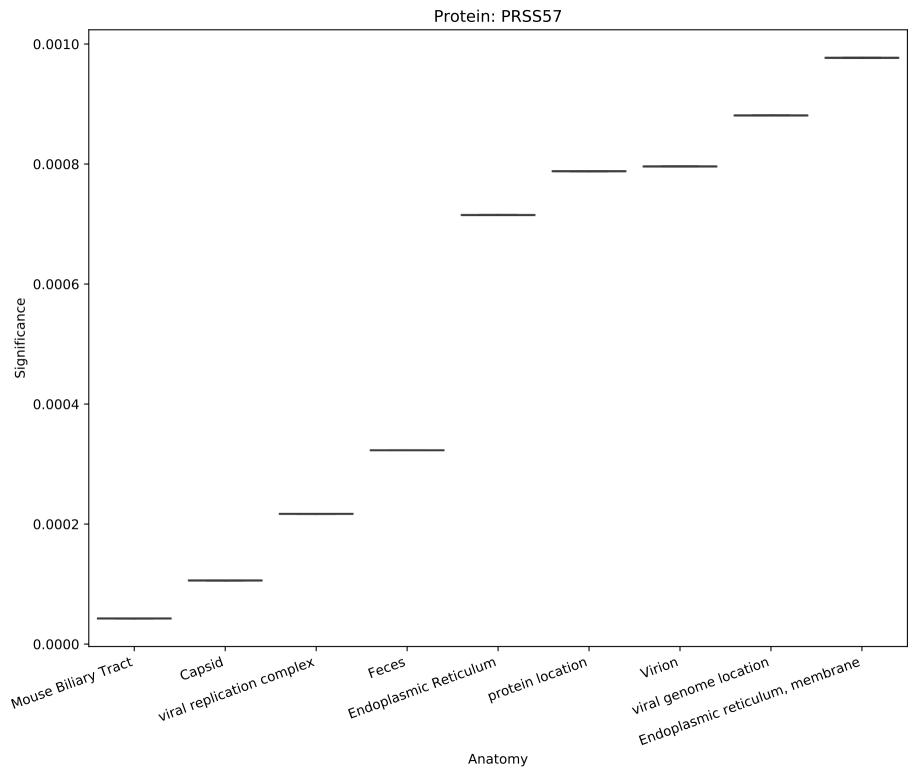


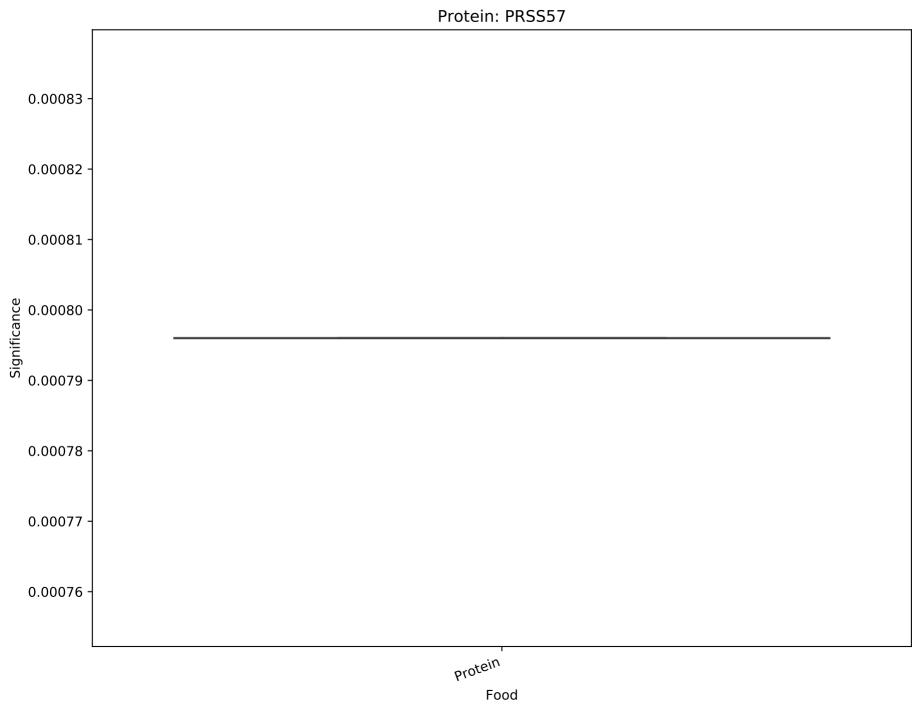


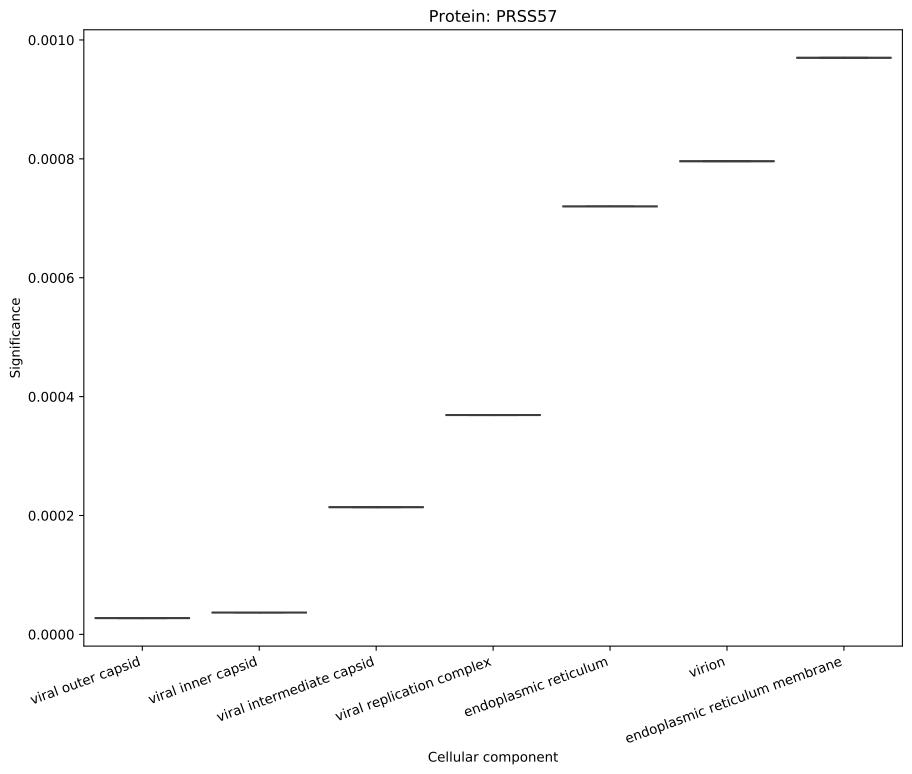


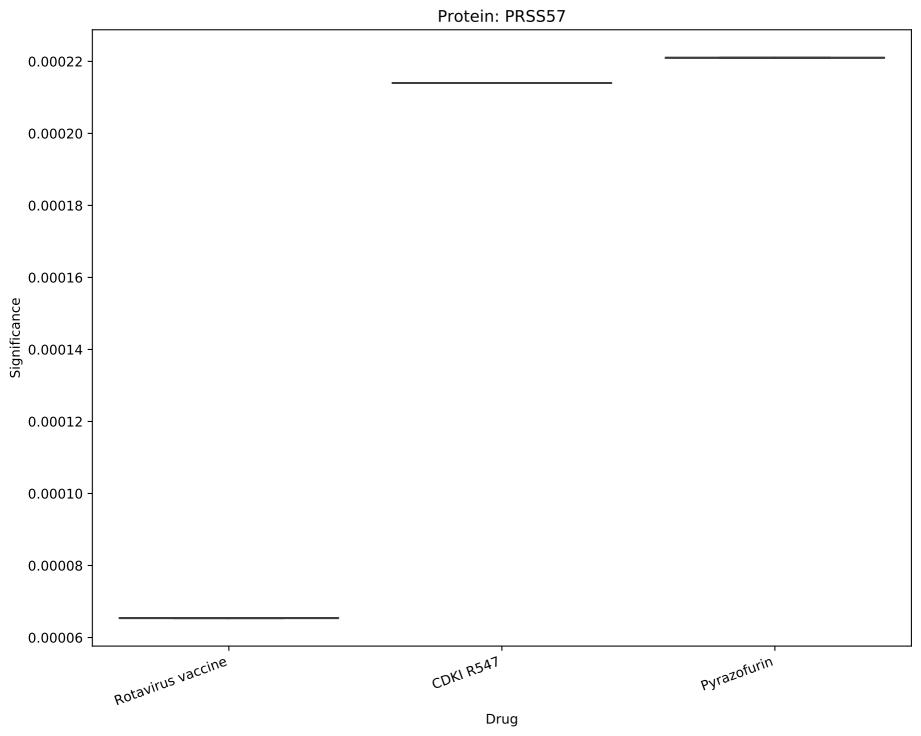


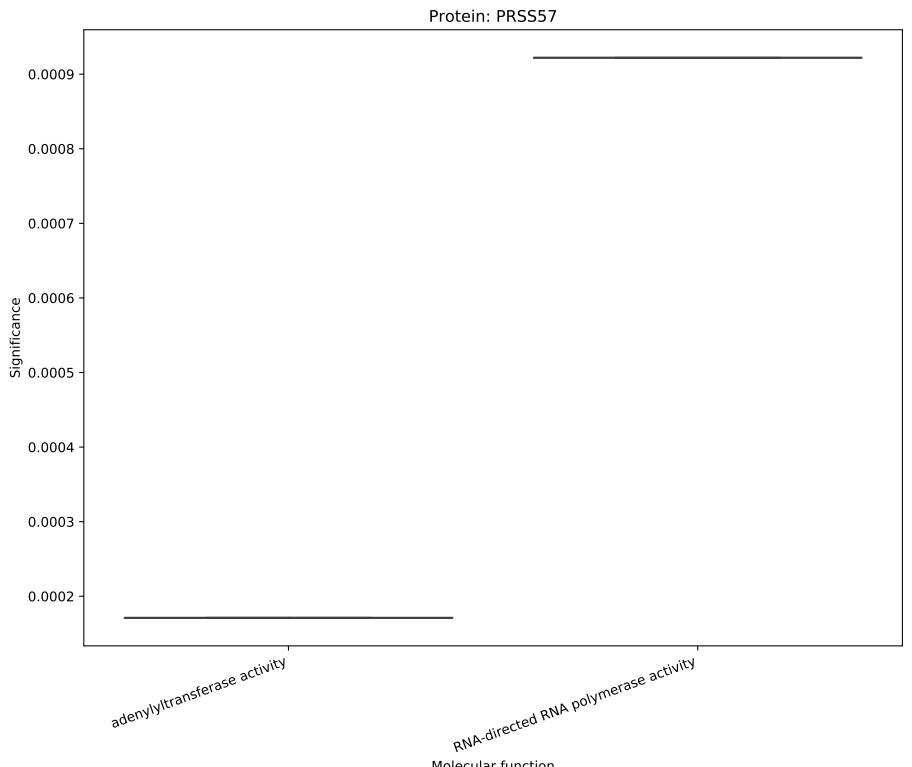
Chemical



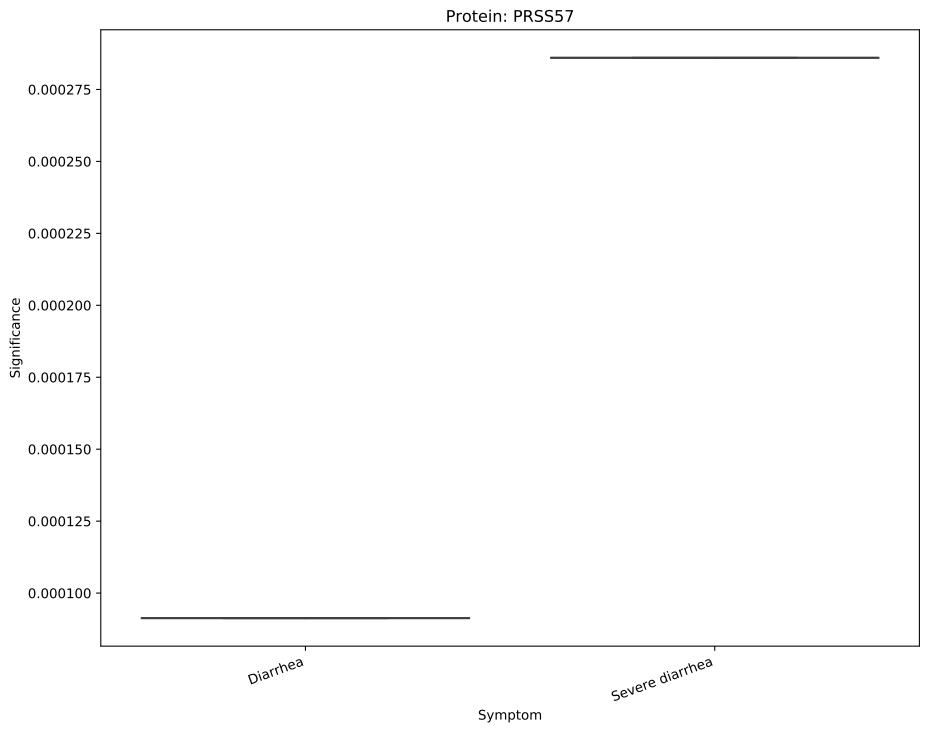


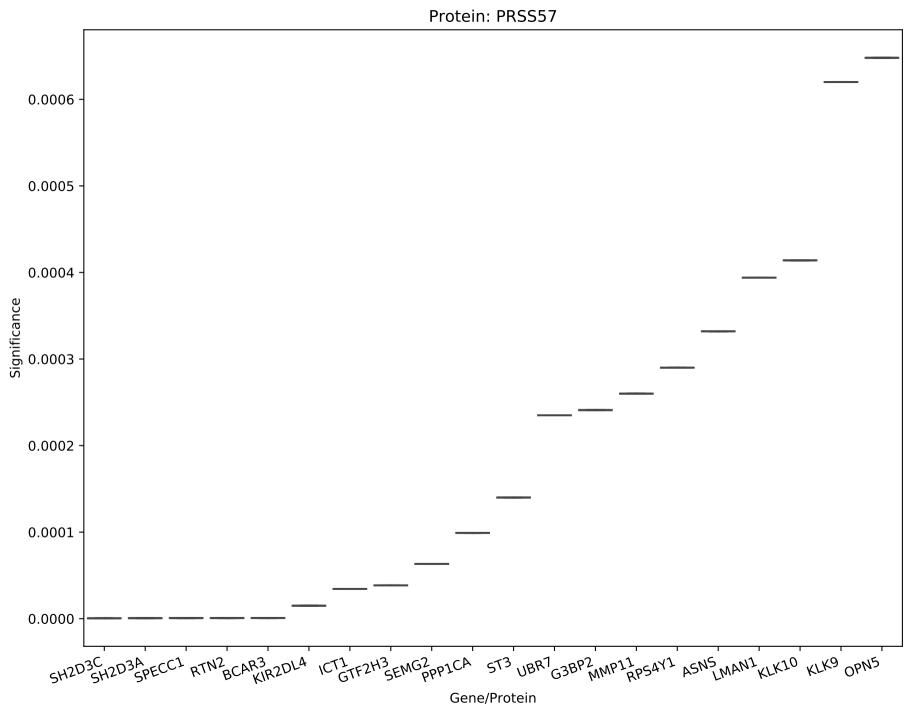


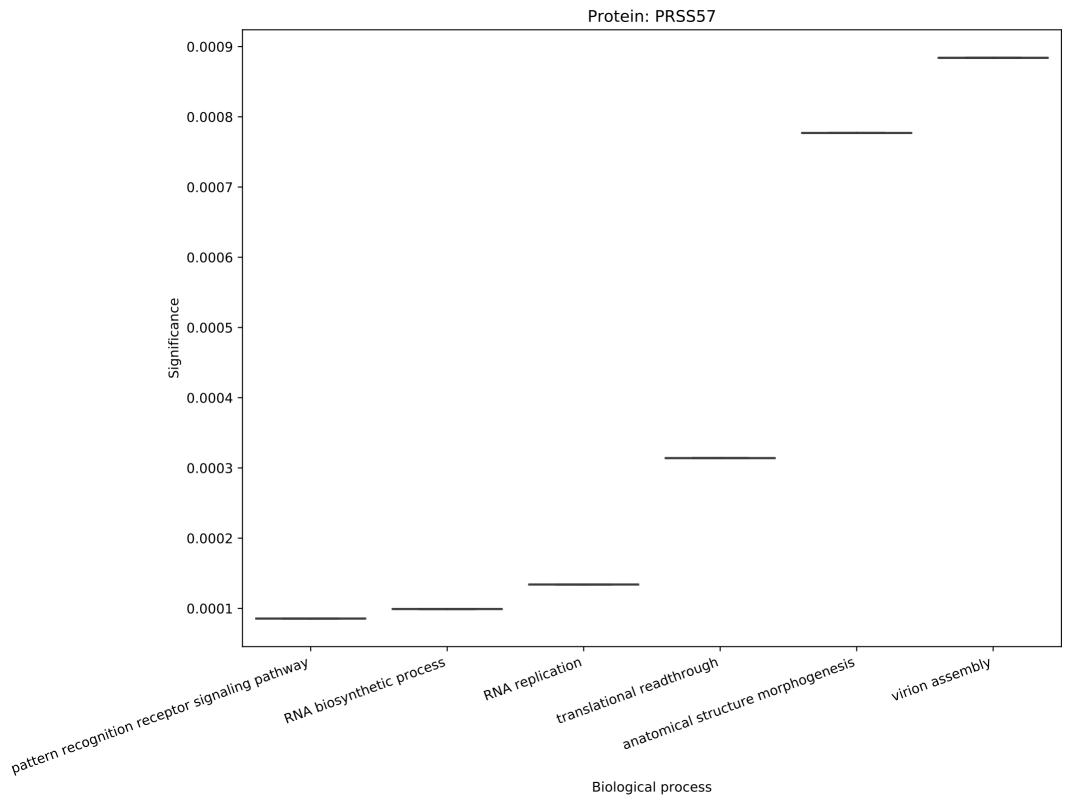




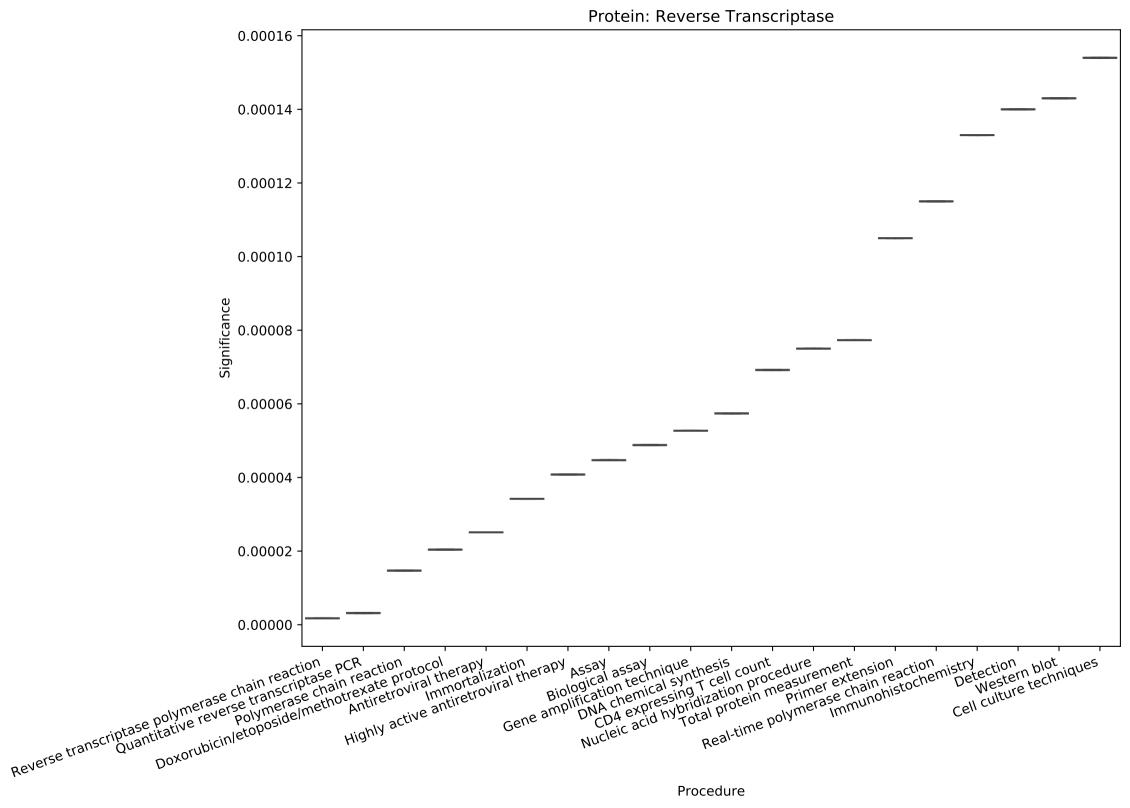
Molecular function

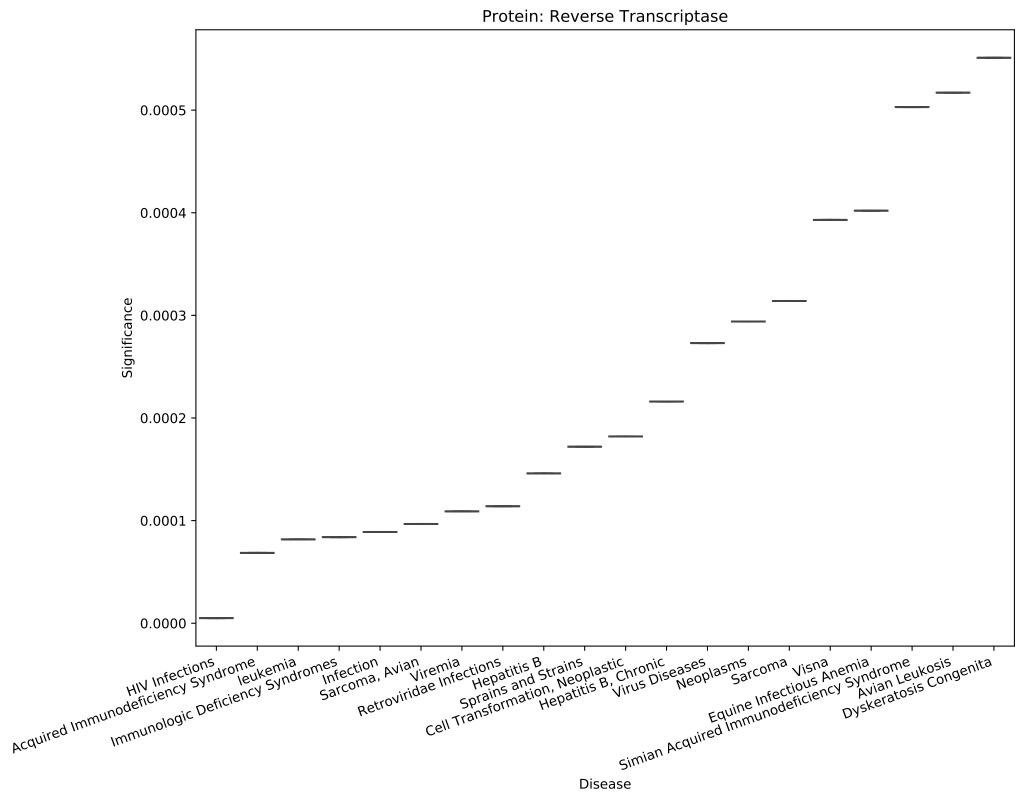


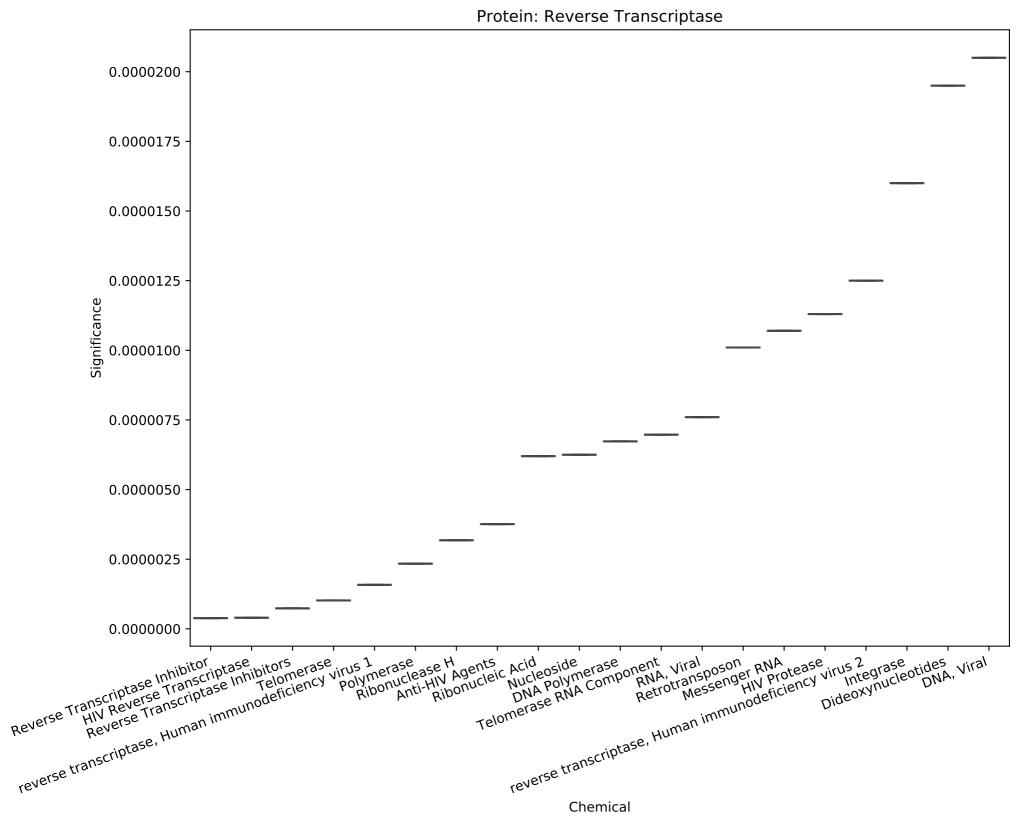


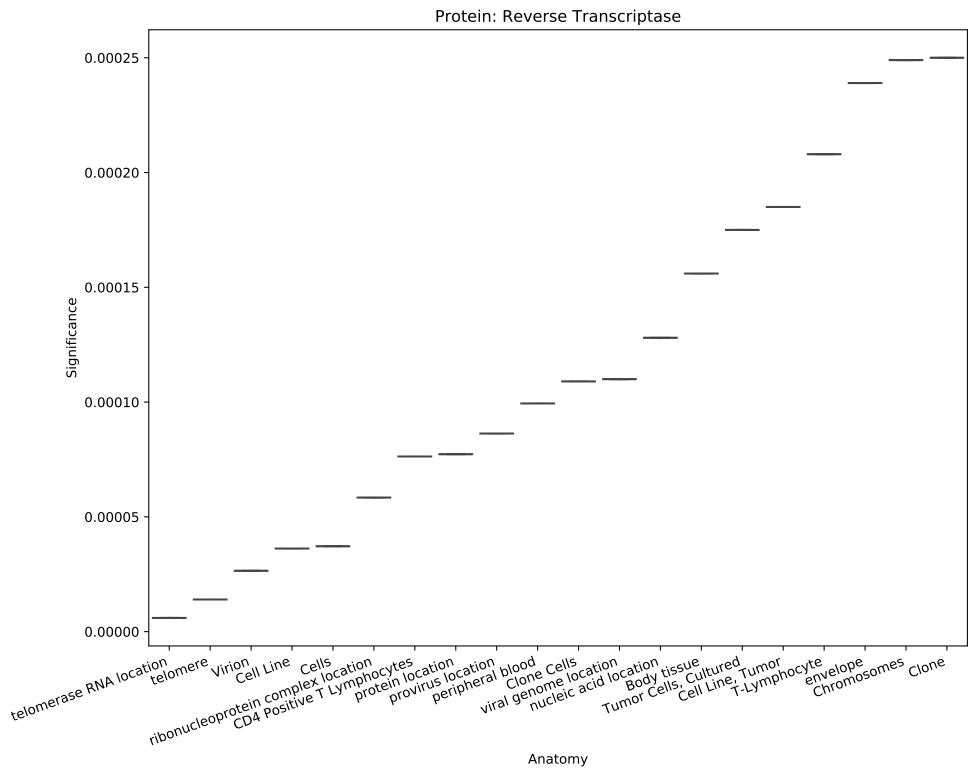


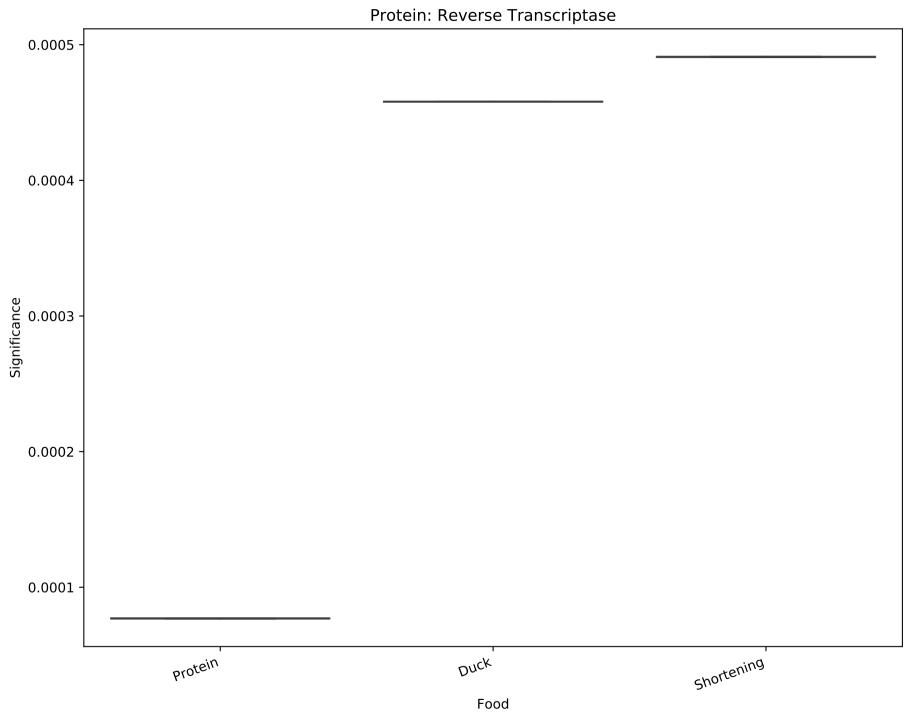
Protein: PRSS57 0.00025 0.00020 0.00015 Significance 0.00010 0.00005 0.00000 Rotavirus virus vi MeSH

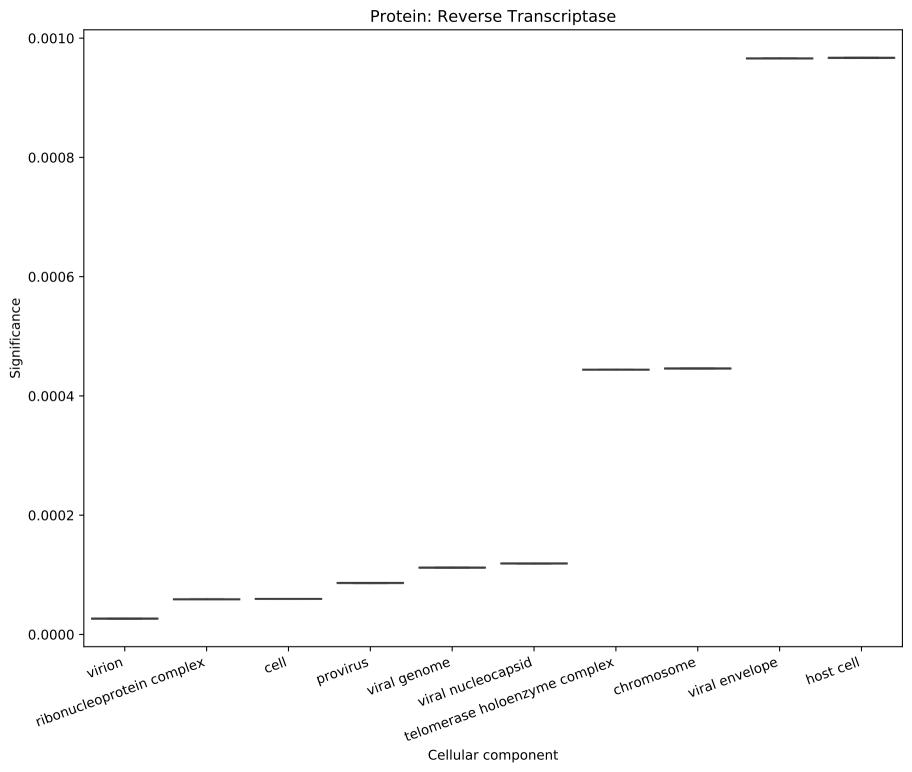


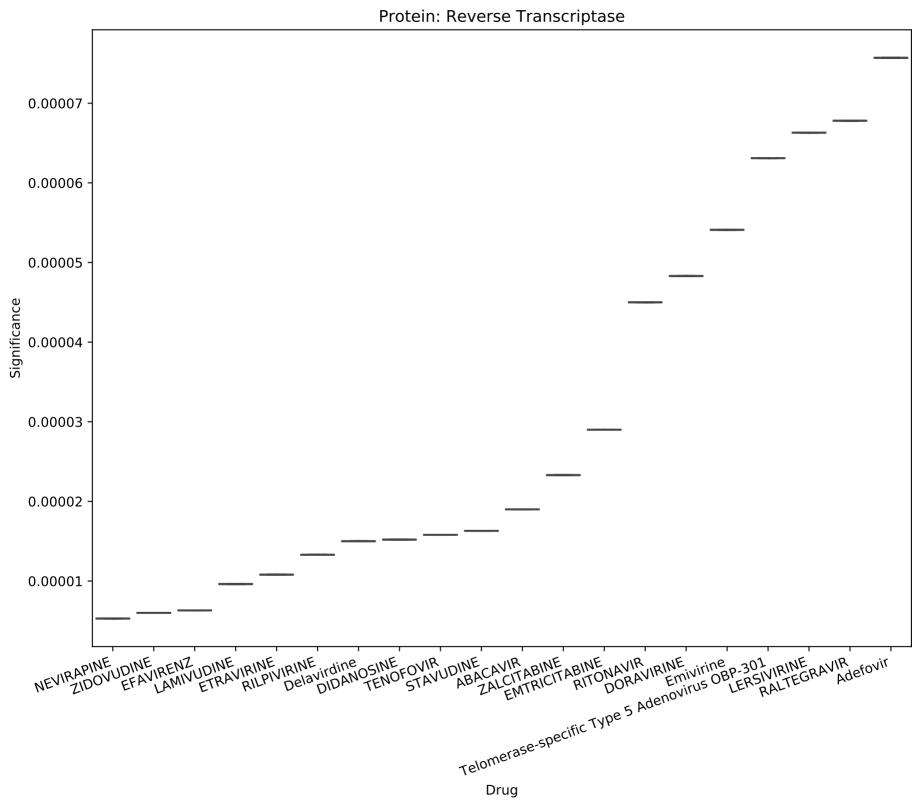


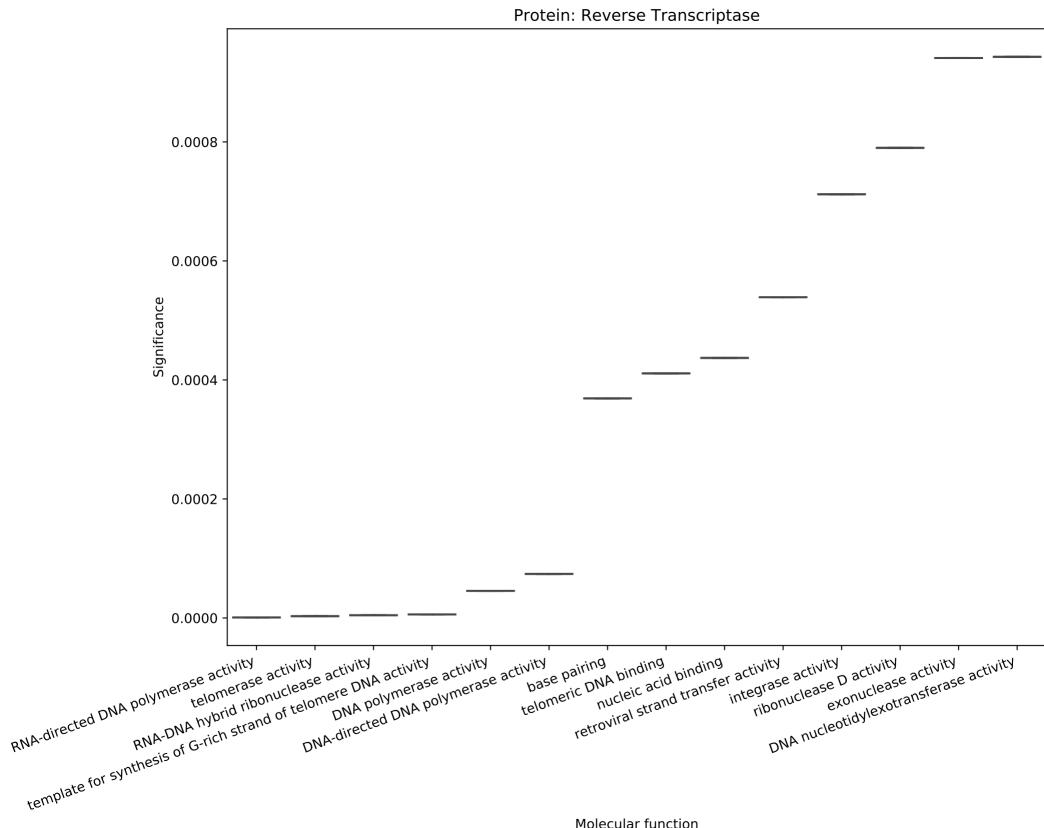


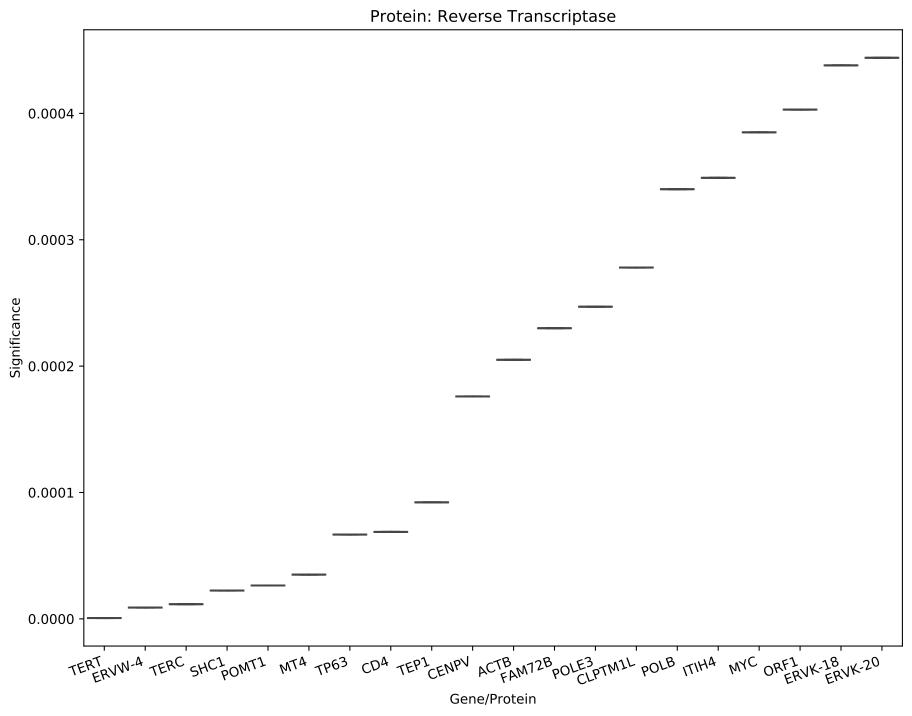


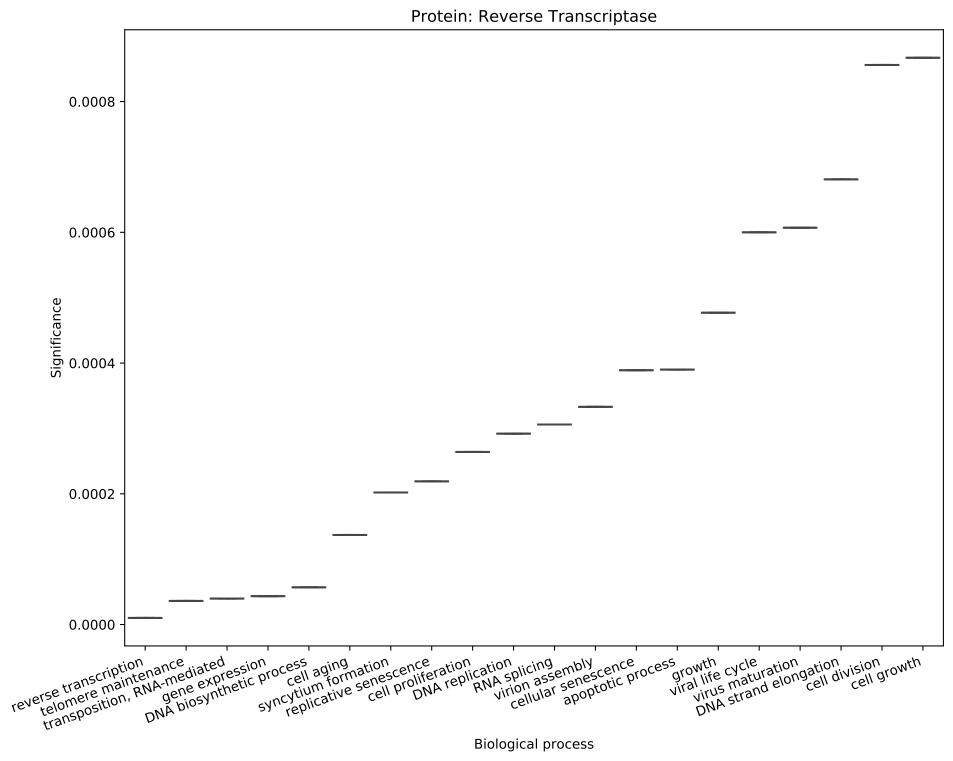












Protein: Reverse Transcriptase 0.000030 -0.000025 -0.000020 -Significance of 0.000015 0.000010 -0.000005 -HIV-1 Viral Virus viridae of polyproses virus estion fovir virus Load ence Viral Active Virus Data Murine HIV-2 of Resistance, Retroviridae of Resistance, Retroviridae of Resistance, Retroviridae of Resistance, Retroviridae of Resistance, Myeloblastosis Retroviridae of Retr

MeSH

