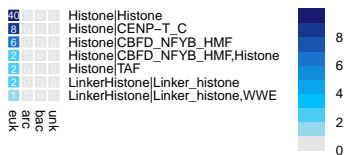
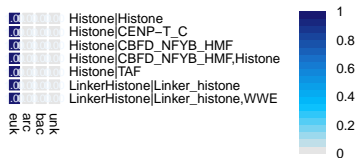


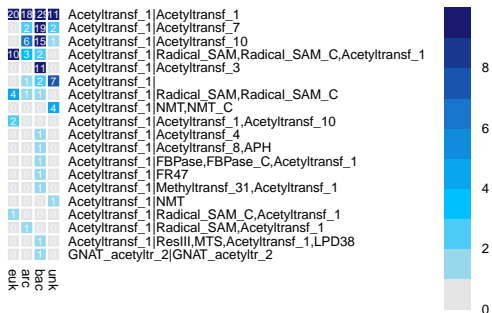
Histone architectures, per closest taxon
n=61 viral genes



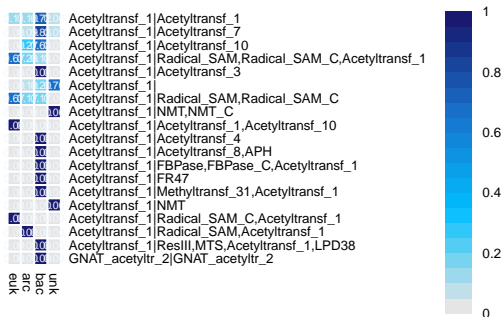
Histone architectures, per closest taxon n=61 viral genes



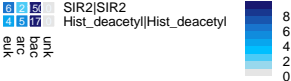
Acetyltransf_1 architectures, per closest taxon
n=281 viral genes



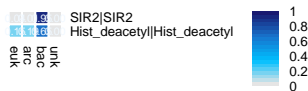
n=281 viral genes



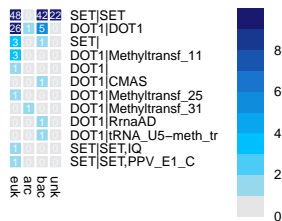
Hist_deacetyl architectures, per closest taxon
n=184 viral genes



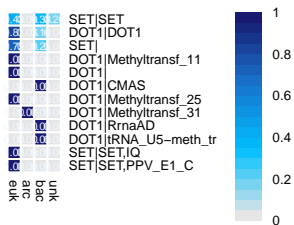
Hist_deacetyl architectures, per closest taxon **n=184 viral genes**



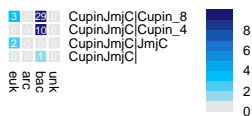
DOT1 architectures, per closest taxon
n=159 viral genes



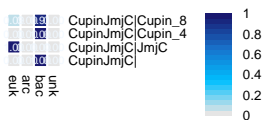
DOT1 architectures, per closest taxon
n=159 viral genes



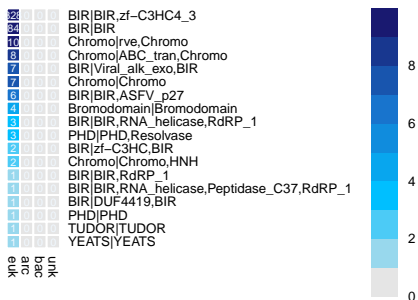
SWIRM architectures, per closest taxon
n=45 viral genes



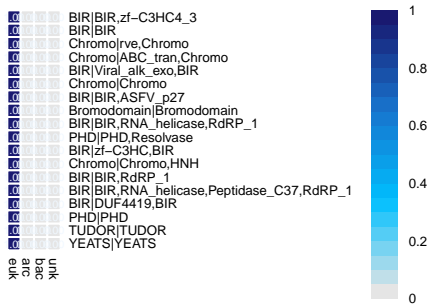
SWIRM architectures, per closest taxon
n=45 viral genes



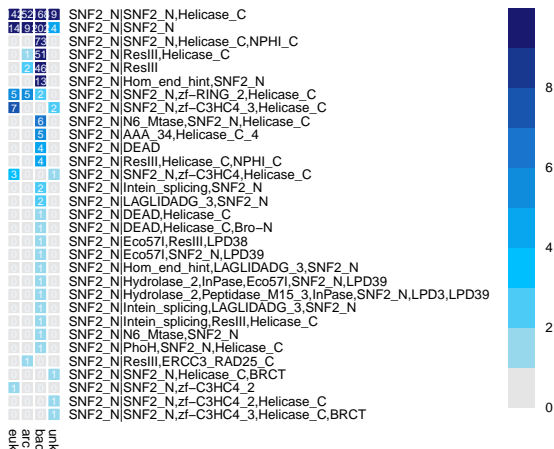
ADD_ATRX architectures, per closest taxon
n=470 viral genes



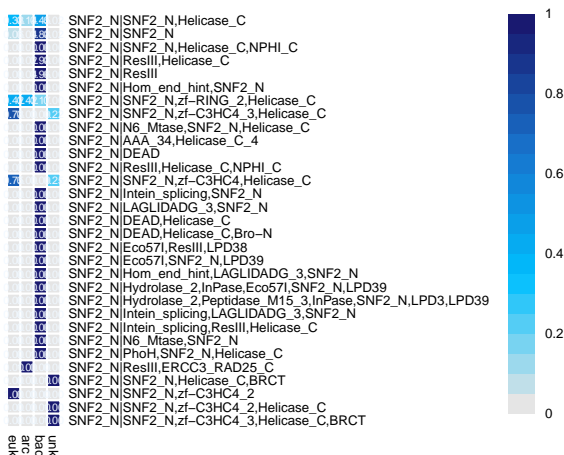
ADD_ATRX architectures, per closest taxon
n=470 viral genes



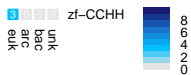
SNF2_N architectures, per closest taxon
n=850 viral genes



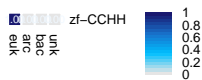
SNF2_N architectures, per closest taxon **n=850 viral genes**



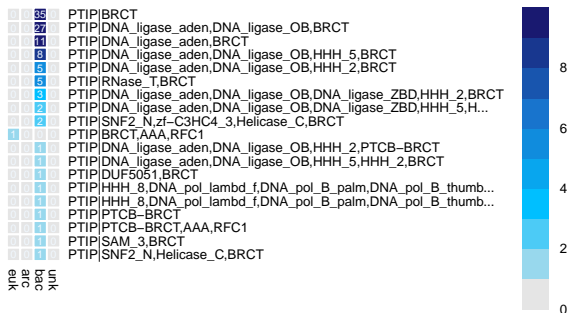
ASF1_hist_chap architectures, per closest taxon
n=3 viral genes



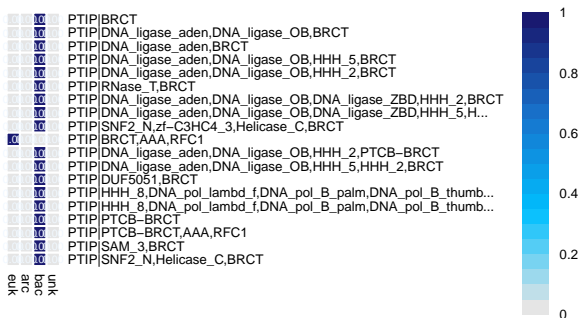
ASF1_hist_chap architectures, per closest taxon
n=3 viral genes



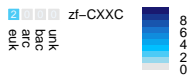
PA1 architectures, per closest taxon
n=108 viral genes



PA1 architectures, per closest taxon
n=108 viral genes



Kelch architectures, per closest taxon
n=2 viral genes



Kelch architectures, per closest taxon n=2 viral genes

