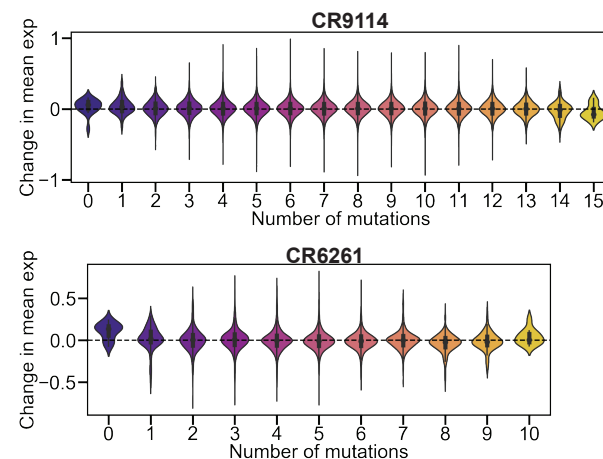
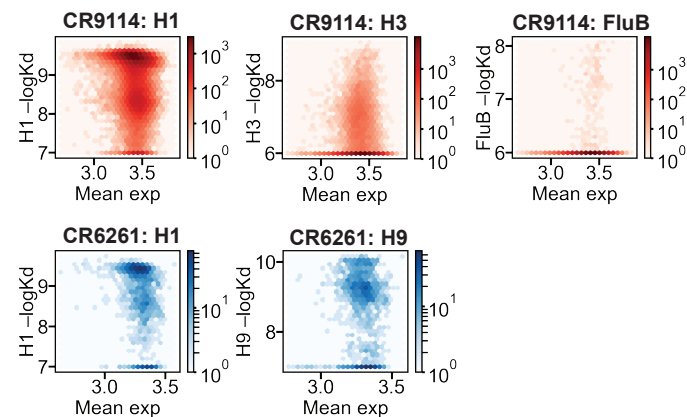


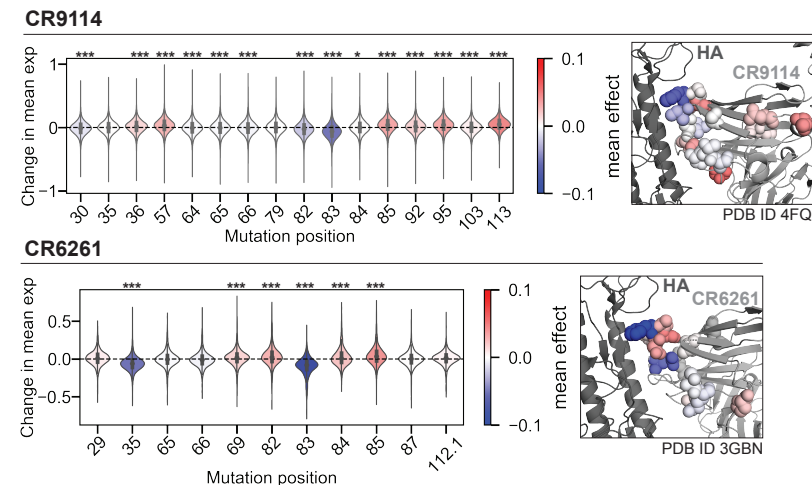
**C** Change in expression by number of mutations



**D** Expression vs. binding affinity by variant



**E** Change in expression by mutation position



**F** Change in expression vs. change in binding affinity by mutation position

