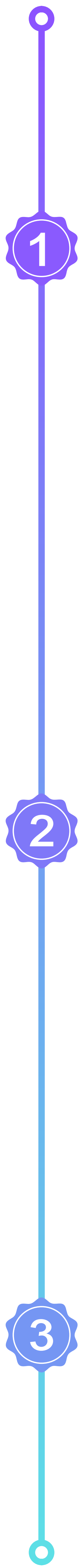


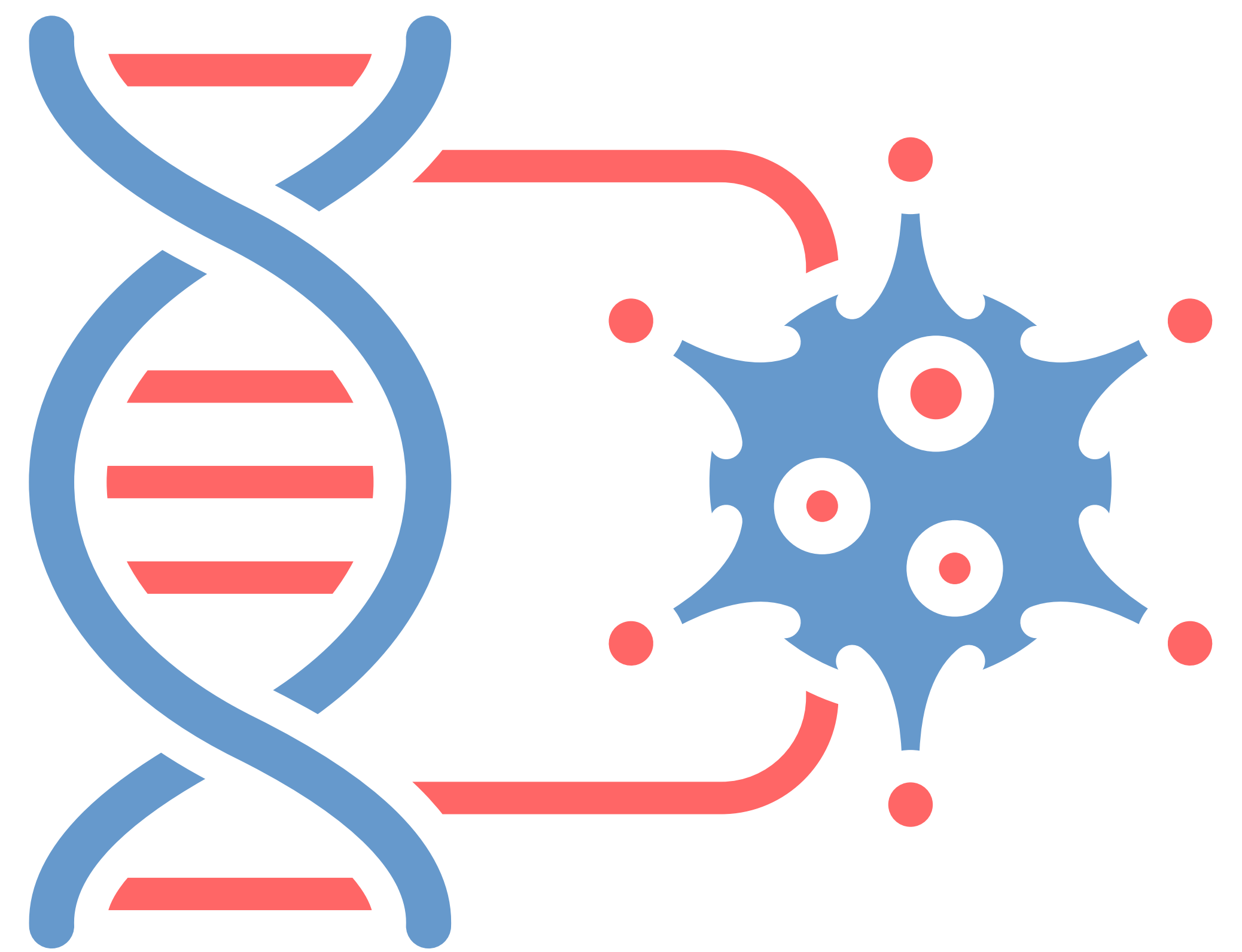
Methodology



1 Use cBioPortal and COSMIC database to identify most common mutations within the protein which contributes to LUAD.



2 Observing and Analyzing Wildtype vs Mutant Proteins. How do mutations change the protein and its functions.



3 Analyze drug binding affinities of both mutated proteins and how do they compare to each other.

