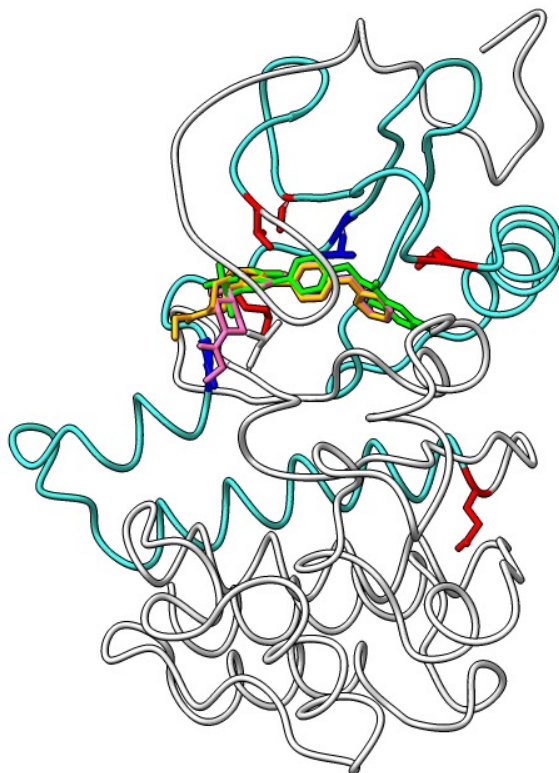


# **Evaluating ibrutinib, pirtobrutinib, remibrutinib, rilzabrutinib, and zanubrutinib in 34 BTK-variants at 7 inhibitor resistance sites**

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**Supplementary table 1.** The BTK codons and the corresponding nucleotide alterations in the respective plasmids

V416A	GTG > GCG	E513G	GAG > GGG
V416E	GTG > GAG	E513A	GAG > GCG
V416G	GTG > GGG	L528F	TTG > TTC
V416L	GTG > CTG	L528M	TTG > ATG
V416M	GTG > ATG	L528S	TTG > TCG
A428D	GCC > GAC	L528V	TTG > GTG
A428G	GCC > GGC	L528W	TTG > TGG
A428P	GCC > CCC	T316A	ACA > GCA
A428S	GCC > TCC	T316I	ACA > ATA
A428T	GCC > ACC	T316K	ACA > AAA
A428V	GCC > GTC	T316P	ACA > CCA
M437I	ATG > ATC	T316R	ACA > AGA
M437K	ATG > AAG	T316S	ACA > TCA
M437R	ATG > AGG		
M437T	ATG > ACG		
M437V	ATG > GTG		
M477I	ATG > ATC		
M477K	ATG > AAG		
M477R	ATG > AGG		
M477T	ATG > ACG		
M477V	ATG > GTG		



### Supplementary Figure S1.

Three-dimensional structure of the BTK kinase domain including ibrutinib (PDB ID 5p9i) (Bender *et al.* 2017). The region containing the investigated amino acids (416-528) is indicated with a cyan ribbon. The structure was superimposed with complexes for zanubrutinib (6j6m) (Guo *et al.* 2019) and pirtobrutinib (8fl1) (Gomez *et al.* 2023). Ibrutinib is in yellow, zanubrutinib in pink, and pirtobrutinib in green. The investigated residues are in red, and residues 474 and 481, studied earlier, are indicated in blue.

### References

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Guo Y, Liu Y, Hu N, Yu D, Zhou C, Shi G *et al.* Discovery of Zanubrutinib (BGB-3111), a Novel, Potent, and Selective Covalent Inhibitor of Bruton's Tyrosine Kinase. *J Med Chem.* 2019; 62:7923-7940.