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Figure S1. The point mutation of T208D results in the conformational change in β 2 region of PI3K α -RBD, disrupting the KRas4B–RBD interface.

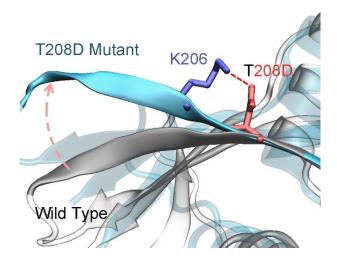


Figure S2. The distance profiles for backbone hydrogen bonds at the antiparallel β-sheet interface in mutated KRas4B–PI3K α -RBD (R230E) complex.

