

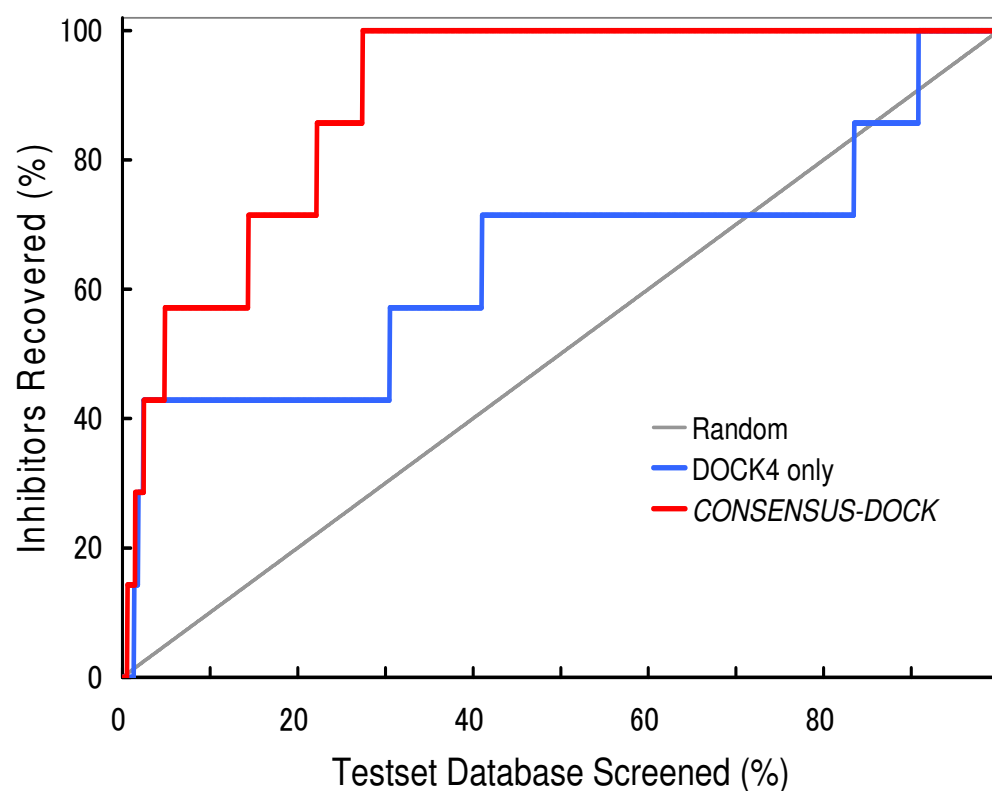
## Supporting Information

### Identification of Death-Associated Protein Kinase Inhibitors Using Structure-Based Virtual Screening

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#### Contents:

Comparison of enrichment curves for *CONSENSUS-DOCK* versus DOCK4 only.



Comparison of enrichment curves for *CONSENSUS-DOCK* versus DOCK4 only. The 'MODEL' structure (in Fig. 2) was used in both calculations. These results obviously indicate that *CONSENSUS-DOCK* has an advantage in the enrichment of inhibitors to DOCK4 only in the Testset Database Screening.