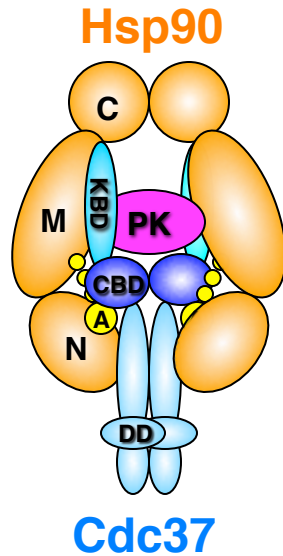


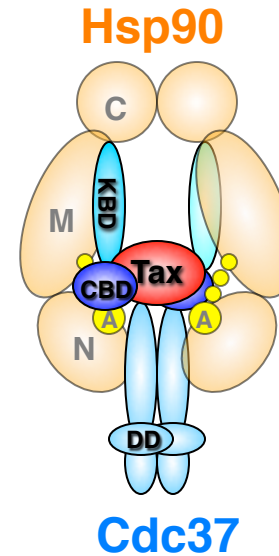
Supplement 3. Schematic representation of Hsp90/Cdc37/Tax ternary complex.

(A)



Hsp90/Cdc37-Kinase complex

(B)



Hsp90/Cdc37-Tax complex

N: ATP binding domain
1-213aa
M: Middle domain
214-432aa
C: Dimerization domain
433-732aa

KBD: Kinase binding domain
40-110aa
CBD: Client binding domain
181-200aa
DD: Dimerization domain
220-250aa