Supplementary Table 1. Statistics of Cryo-EM Datasets

Sample	Micrographs (N)	Reconstruction	Particles (N)	Resolution (Å)
CBF3−*CEN3* (Supplementary Figure 2)	3,537		987,484 Auto-picked particles	
		CBF3^{core}	146,673 (30.6%)	-
		Monomeric CBF3−*CEN3*	175,447 (36.6%)	-
		Dimeric CBF3−*CEN3*	157,883 (32.8%)	4.4
		CBF3^{msk}	198,010	3.0
		Others (Hybrid and bad particles not included in the 3D classification)	507,481	
CBF3^{core}−Ndc10^{DBD} -/+*CEN3* (Supplementary Figure 3)			757,433 Auto-picked particles	
CBF3^{core}−Ndc10^{DBD} CBF3^{core}−Ndc10^{DBD}−*CEN3*	724 1,171	CBF3^{core}−Ndc10^{DBD}	104,938 (31.8%)	3.6
		CBF3^{core}	184,356 (55.8%)	3.9
		Monomeric CBF3^{core}−Ndc10^{DBD}−*CEN3*	27,669 (8.4%)	-
		Dimeric CBF3^{core}−Ndc10^{DBD}−*CEN3*	13,140 (4.0%)	-
		Others (Hybrid and bad particles not included in the 3D classification)	427,330	
CBF3−*CEN3*$^{biotin-avidin}$	400		113,800 Auto-picked particles	
		CBF3−*CEN3*$^{biotin-avidin}$	10,889 (15.2%)	12.5
		CBF3^{core}	19,626 (27.5%)	-
		CBF3^{core}−Ndc10^{DBD}	40,919 (57.3%)	-
		Others (Hybrid and bad particles not included in the 3D classification)	42,366	