S3 Fig. ATP binding sites of CtCCT subunits

CtCCT5

CtCCT6

CtCCT7

CtCCT8

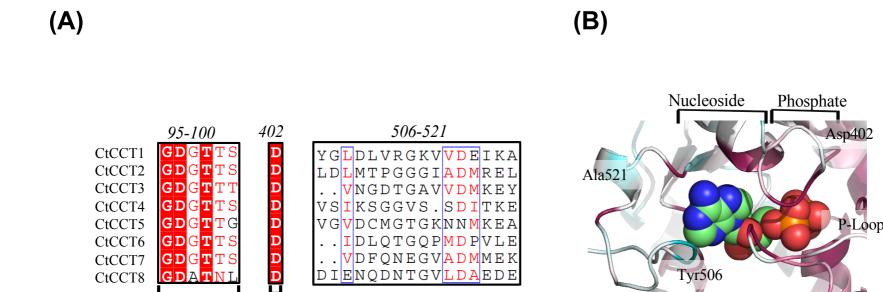
GDGTTG

GDGTTS

GDGTTS

P-Loop

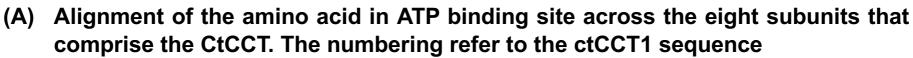
Catalytic



VG<mark>V</mark>DCMGTGKNN<mark>M</mark>KEA

DIENODNTGVLDAEDE

IDLQTGQPMDPVLE VDFQNEGV<mark>ADM</mark>MEK



Tyr506

conserved

variable

CtCCT1 active site colored according to conservation scores calculated for (B) CtCCT1 - CtCCT8