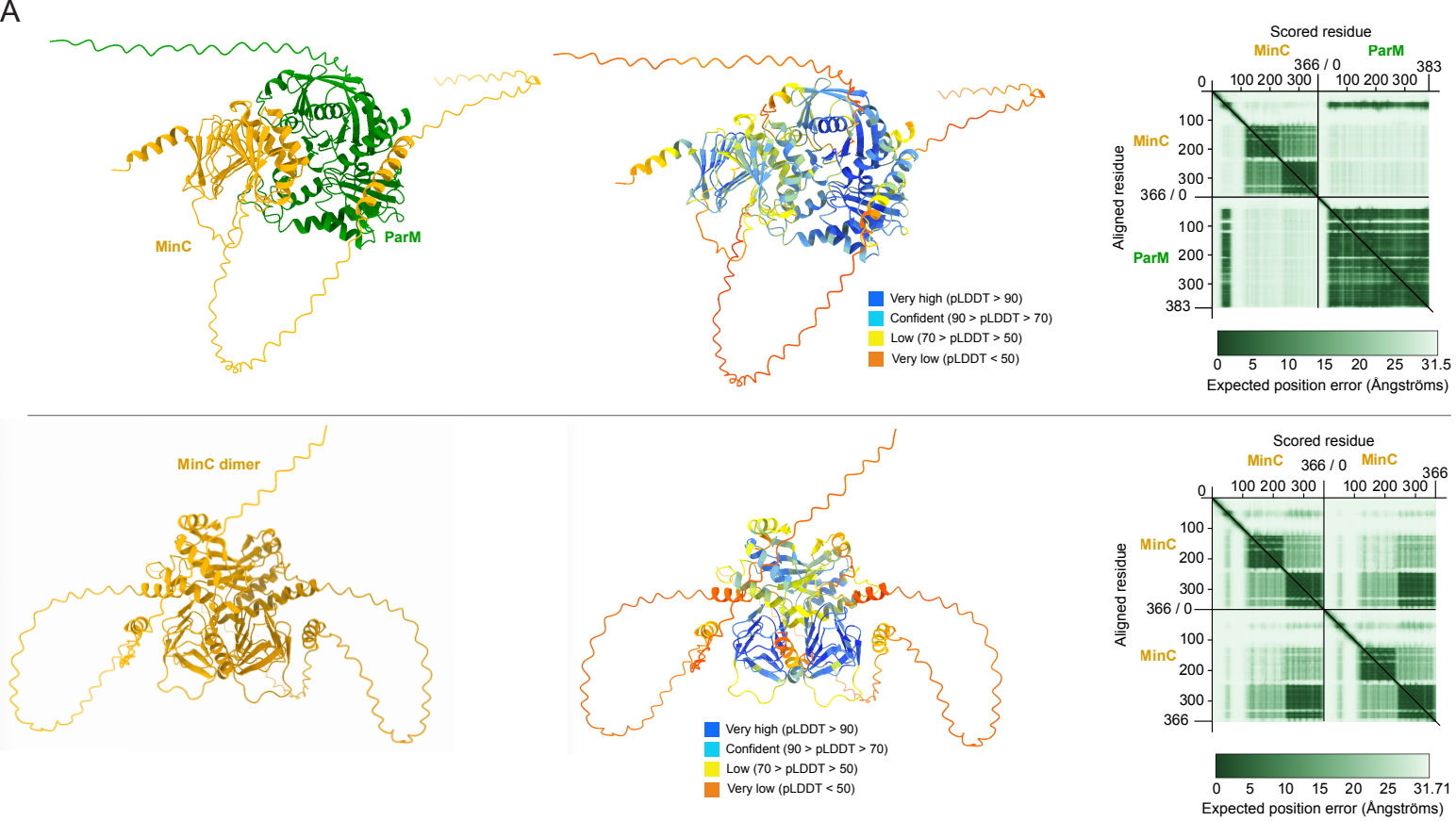
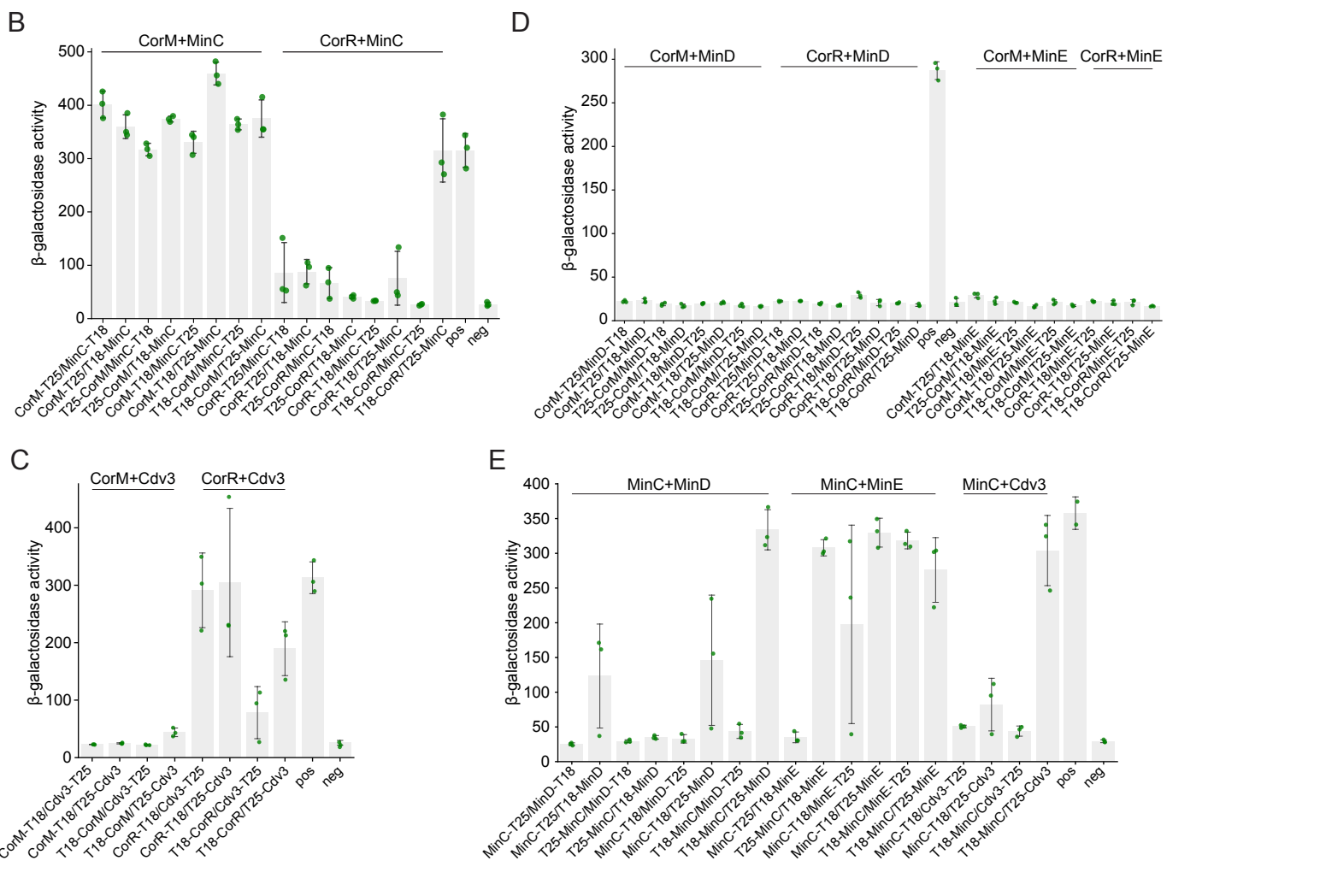

A
MinC
ParM
Very high (pLDDT > 90)
Confident (90 > pLDDT > 70)
Low (70 > pLDDT > 50)
Very low (pLDDT < 50)
Scored residue
MinC
ParM
366 / 0
383
Aligned residue
0 5 10 15 20 25 31.5
Expected position error (Ångströms)
MinC dimer
366
0 5 10 15 20 25 31.71


B
CorM+MinC
CorR+MinC
β-galactosidase activity
CorM-T25/MinC-T18
CorM-T25/T18-MinC
T25-CorM/MinC-T18
T25-CorM/T18-MinC
CorM-T18/MinC-T25
CorM-T18/T25-MinC
T18-CorM/MinC-T25
T18-CorM/T25-MinC
CorR-T25/MinC-T18
CorR-T25/T18-MinC
T25-CorR/MinC-T18
T25-CorR/T18-MinC
CorR-T18/MinC-T25
CorR-T18/T25-MinC
T18-CorR/MinC-T25
T18-CorR/T25-MinC
pos
neg
D
CorM+MinD
CorR+MinD
CorM+MinE
CorR+MinE
CorM-T25/MinD-T18
CorM-T25/T18-MinD
T25-CorM/MinD-T18
T25-CorM/T18-MinD
CorM-T18/MinD-T25
CorM-T18/T25-MinD
T18-CorM/MinD-T25
T18-CorM/T25-MinD
CorR-T25/MinD-T18
CorR-T25/T18-MinD
T25-CorR/MinD-T18
T25-CorR/T18-MinD
CorR-T18/MinD-T25
CorR-T18/T25-MinD
T18-CorR/MinD-T25
T18-CorR/T25-MinD
pos
neg
CorM-T25/T18-MinE
T25-CorM/T18-MinE
CorM-T18/MinE-T25
T18-CorM/MinE-T25
CorM-T18/T25-MinE
T18-CorM/T25-MinE
CorR-T25/T18-MinE
T25-CorR/T18-MinE
CorR-T18/MinE-T25
T18-CorR/MinE-T25
CorR-T18/T25-MinE
T18-CorR/T25-MinE
C
CorM+Cdv3
CorR+Cdv3
β-galactosidase activity
CorM-T18/Cdv3-T25
CorM-T18/T25-Cdv3
T18-CorM/Cdv3-T25
T18-CorM/T25-Cdv3
CorR-T18/Cdv3-T25
CorR-T18/T25-Cdv3
T18-CorR/Cdv3-T25
T18-CorR/T25-Cdv3
pos
neg
E
MinC+MinD
MinC+MinE
MinC+Cdv3
β-galactosidase activity
MinC-T25/MinD-T18
MinC-T25/T18-MinD
T25-MinC/MinD-T18
T25-MinC/T18-MinD
MinC-T18/MinD-T25
MinC-T18/T25-MinD
T18-MinC/MinD-T25
T18-MinC/T25-MinD
MinC-T25/T18-MinE
T25-MinC/T18-MinE
MinC-T18/MinE-T25
MinC-T18/T25-MinE
T18-MinC/MinE-T25
T18-MinC/T25-MinE
MinC-T18/Cdv3-T25
MinC-T18/T25-Cdv3
T18-MinC/Cdv3-T25
T18-MinC/T25-Cdv3
pos
neg