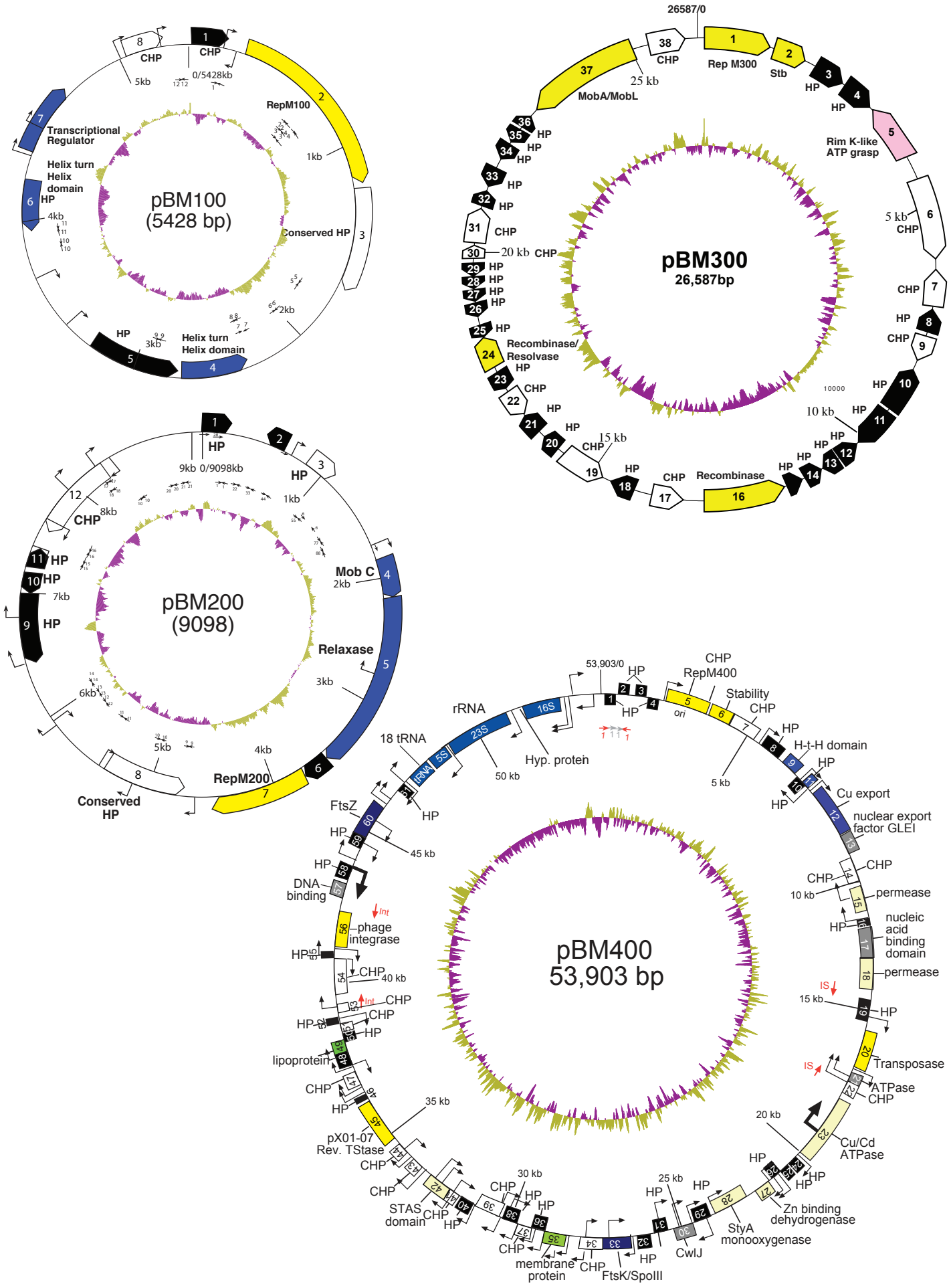
Figure S1


pBM100
(5428 bp)
0/5428kb
RepM100
Conserved HP
Transcriptional Regulator
Helix turn Helix domain
HP
CHP
pBM300
26,587bp
26587/0
Rep M300
Stb
Rim K-like ATP grasp
MobA/MobL
Recombinase/ Resolvase
Recombinase
pBM200
(9098)
9kb 0/9098kb
Mob C
Relaxase
RepM200
Conserved HP
pBM400
53,903 bp
53,903/0
RepM400
Stability
ori
rRNA
16S
23S
5S
tRNA
18 tRNA
Hyp. protein
H-t-H domain
Cu export
nuclear export factor GLEI
permease
nucleic acid binding domain
Transposase
ATPase
Cu/Cd ATPase
Zn binding dehydrogenase
StyA monooxygenase
CwlJ
FtsK/SpoIII
membrane protein
STAS domain
pX01-07 Rev. TStase
lipoprotein
DNA binding
phage integrase
FtsZ
Int
IS