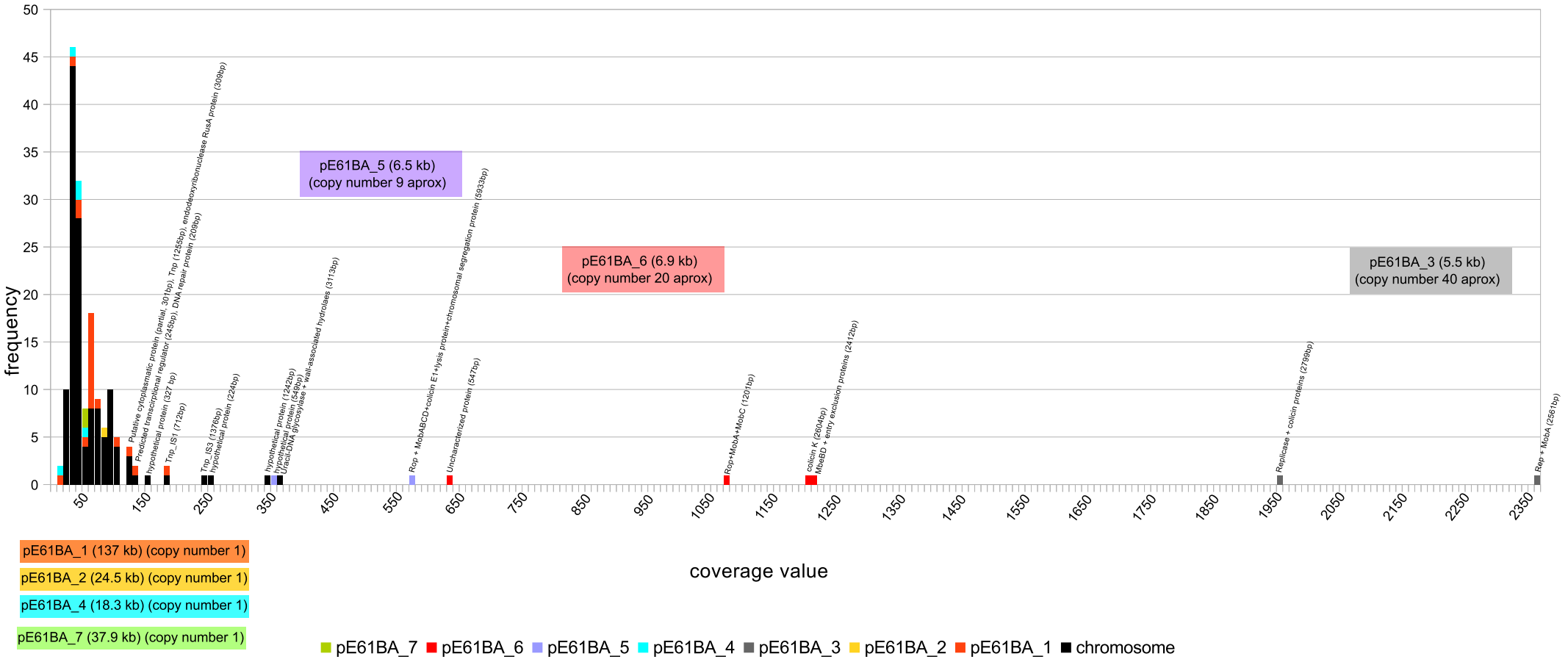
Figure S9


frequency
coverage value
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(copy number 9 aprox)
pE61BA_6 (6.9 kb)
(copy number 20 aprox)
pE61BA_3 (5.5 kb)
(copy number 40 aprox)
Putative cytoplasmatic protein (partial, 301bp), Tnp (1255bp), endodeoxyribonuclease RusA protein (309bp)
Predicted transcriptional regulator (245bp), DNA repair protein (209bp)
hypothetical protein (327 bp)
Tnp_IS1 (712bp)
Tnp_IS3 (1376bp)
hypothetical protein (224bp)
hypothetical protein (1242bp)
hypothetical protein (549bp)
Uracil-DNA glycosylase + wall-associated hydrolaes (3113bp)
Rop + MobABCD+colicin E1+lysis protein+chromosomal segregation protein (5933bp)
Uncharacterized protein (547bp)
Rop+MobA+MobC (1201bp)
colicin K (2604bp)
MbeBD + entry exclusion proteins (2412bp)
Replicase + colicin proteins (2798bp)
Rep + MobA (2561bp)
pE61BA_1 (137 kb) (copy number 1)
pE61BA_2 (24.5 kb) (copy number 1)
pE61BA_4 (18.3 kb) (copy number 1)
pE61BA_7 (37.9 kb) (copy number 1)
pE61BA_7
pE61BA_6
pE61BA_5
pE61BA_4
pE61BA_3
pE61BA_2
pE61BA_1
chromosome