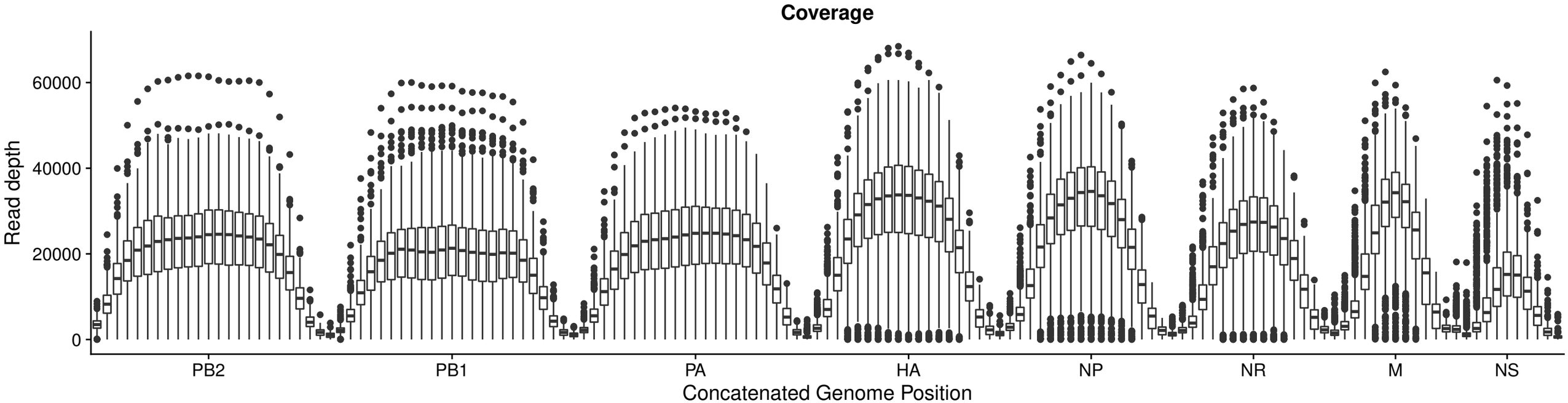
Coverage
Read depth
60000
40000
20000
0
PB2
PB1
PA
HA
NP
NR
M
NS
Concatenated Genome Position