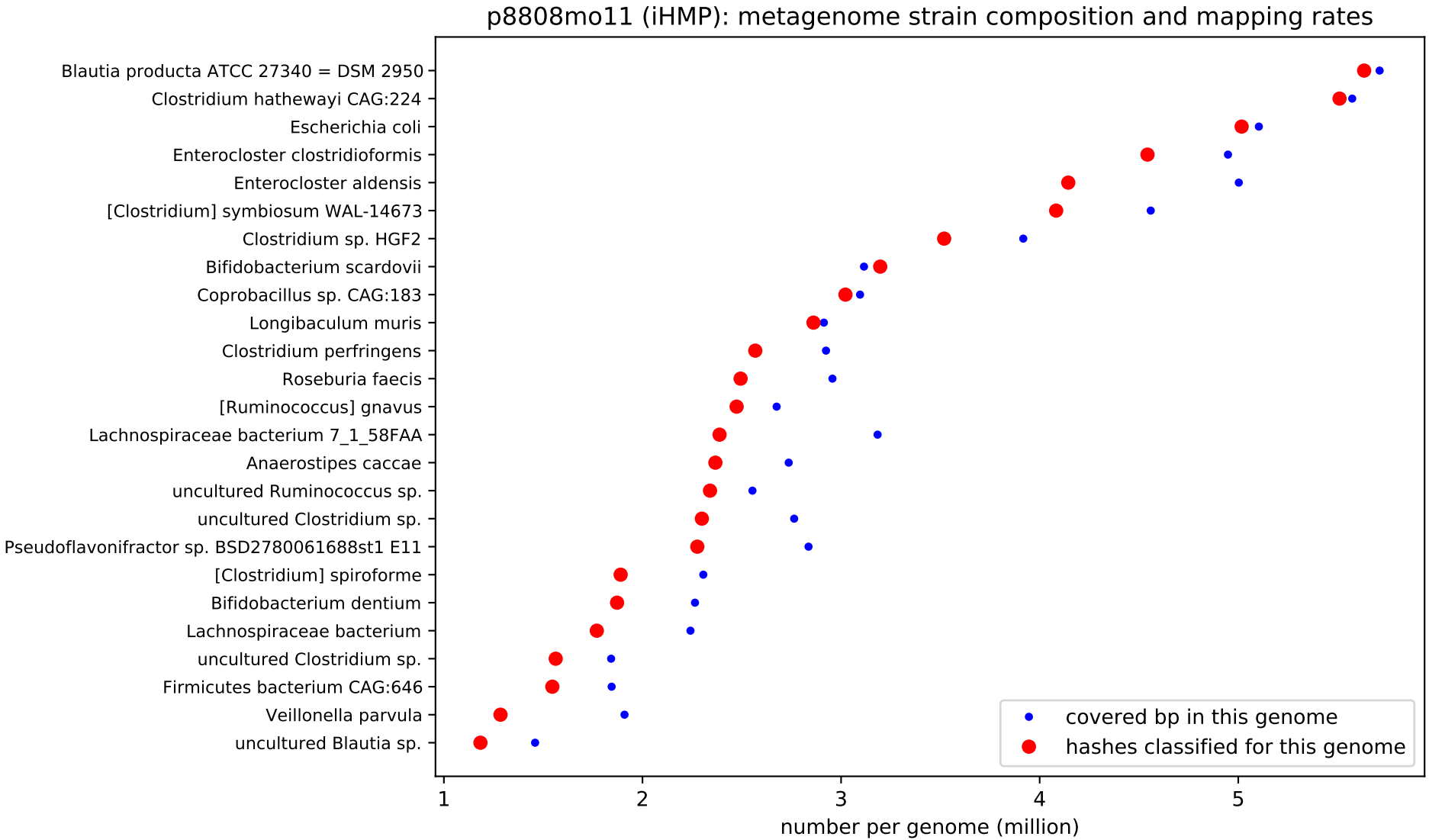
p8808mo11 (iHMP): metagenome strain composition and mapping rates
Blautia producta ATCC 27340 = DSM 2950
Clostridium hathewayi CAG:224
Escherichia coli
Enterocloster clostridioformis
Enterocloster aldensis
[Clostridium] symbiosum WAL-14673
Clostridium sp. HGF2
Bifidobacterium scardovii
Coprobacillus sp. CAG:183
Longibaculum muris
Clostridium perfringens
Roseburia faecis
[Ruminococcus] gnavus
Lachnospiraceae bacterium 7_1_58FAA
Anaerostipes caccae
uncultured Ruminococcus sp.
uncultured Clostridium sp.
Pseudoflavonifractor sp. BSD2780061688st1 E11
[Clostridium] spiroforme
Bifidobacterium dentium
Lachnospiraceae bacterium
uncultured Clostridium sp.
Firmicutes bacterium CAG:646
Veillonella parvula
uncultured Blautia sp.
covered bp in this genome
hashes classified for this genome
1
2
3
4
5
number per genome (million)