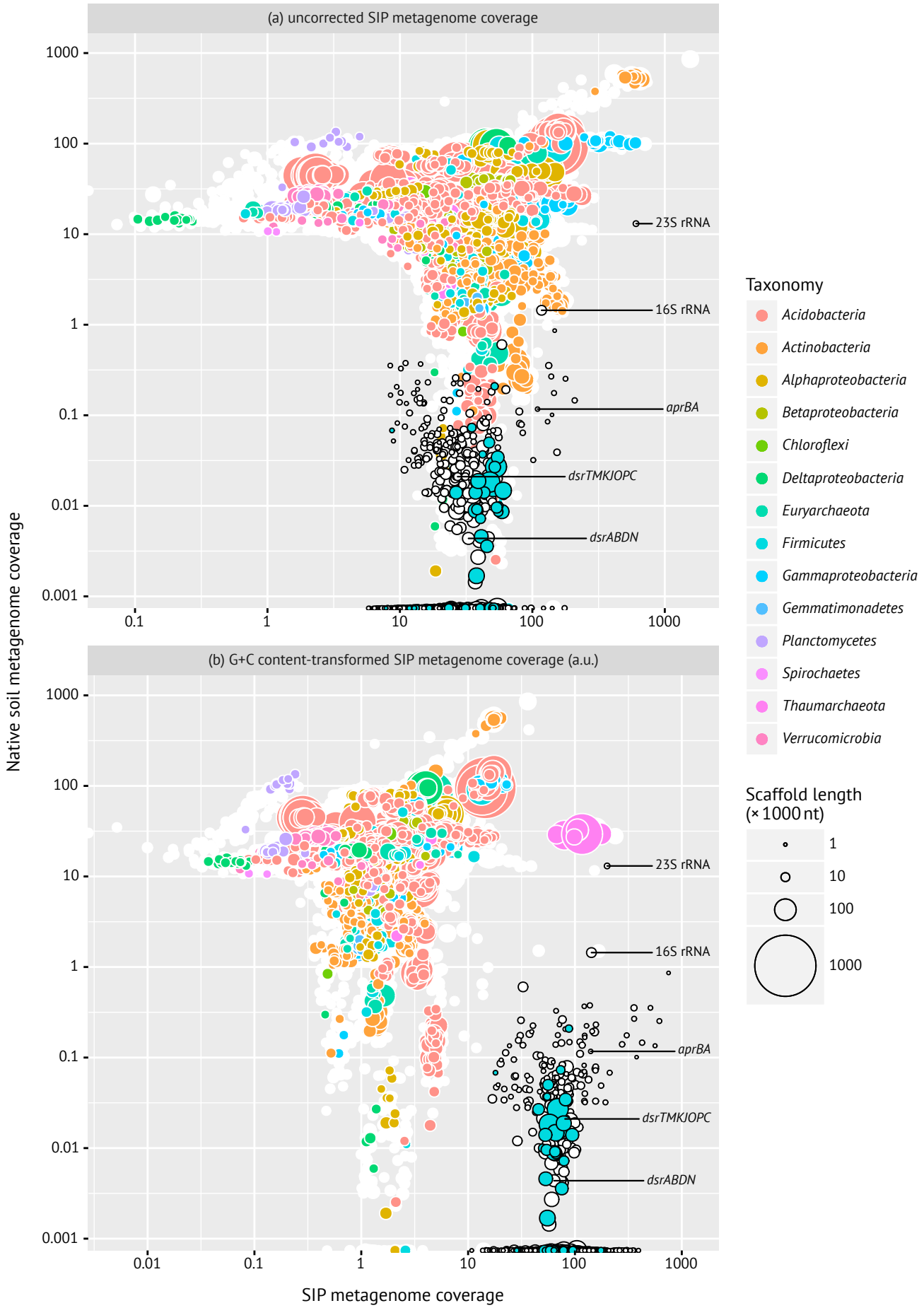
Fig. S1.


(a) uncorrected SIP metagenome coverage
(b) G+C content-transformed SIP metagenome coverage (a.u.)
Native soil metagenome coverage
SIP metagenome coverage
23S rRNA
16S rRNA
aprBA
dsrTMKJOPC
dsrABDN
Taxonomy
Acidobacteria
Actinobacteria
Alphaproteobacteria
Betaproteobacteria
Chloroflexi
Deltaproteobacteria
Euryarchaeota
Firmicutes
Gammaproteobacteria
Gemmatimonadetes
Planctomycetes
Spirochaetes
Thaumarchaeota
Verrucomicrobia
Scaffold length
(× 1000 nt)
1
10
100
1000