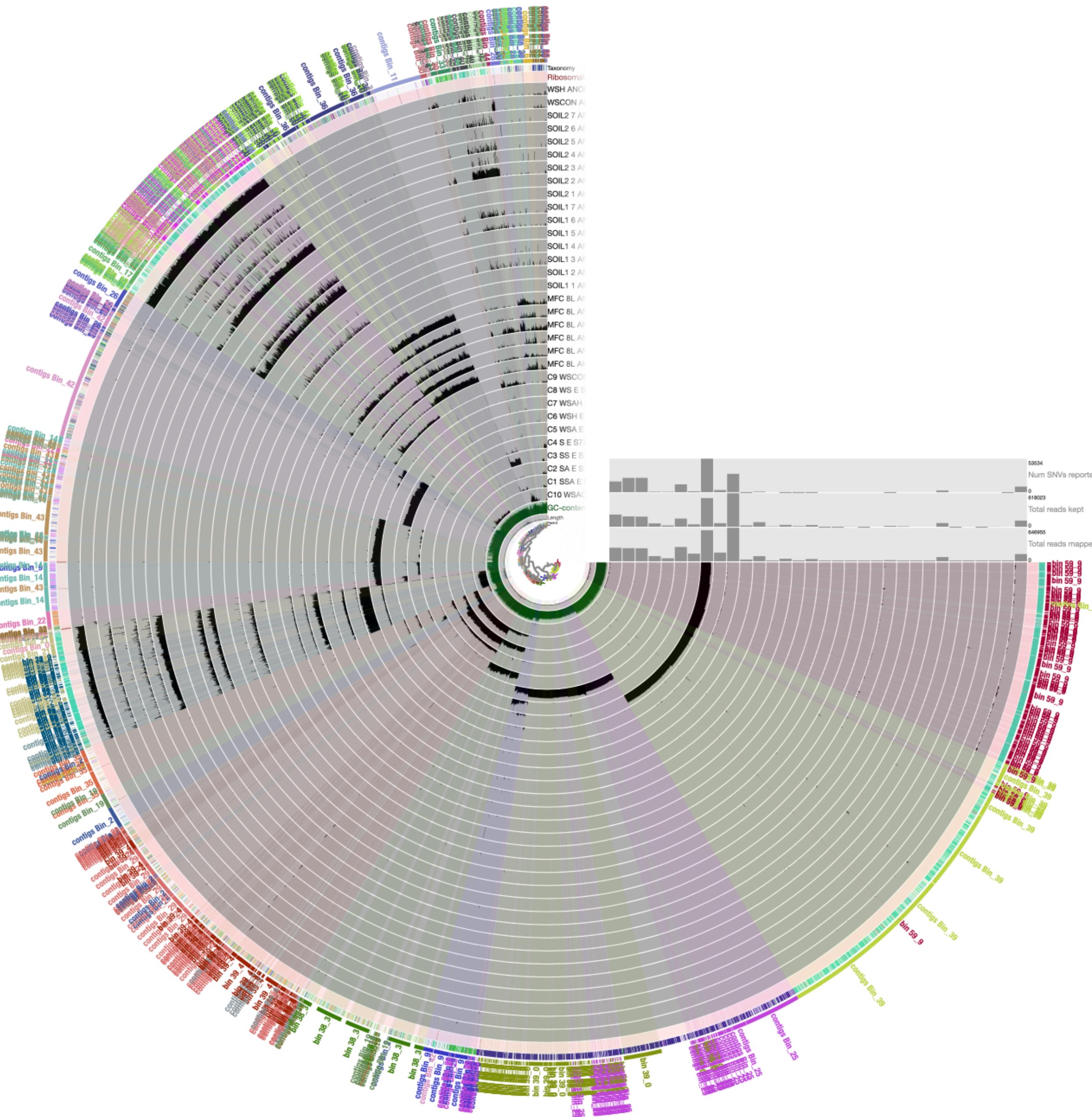


My bins, second attempt

SAMPLES MERGED

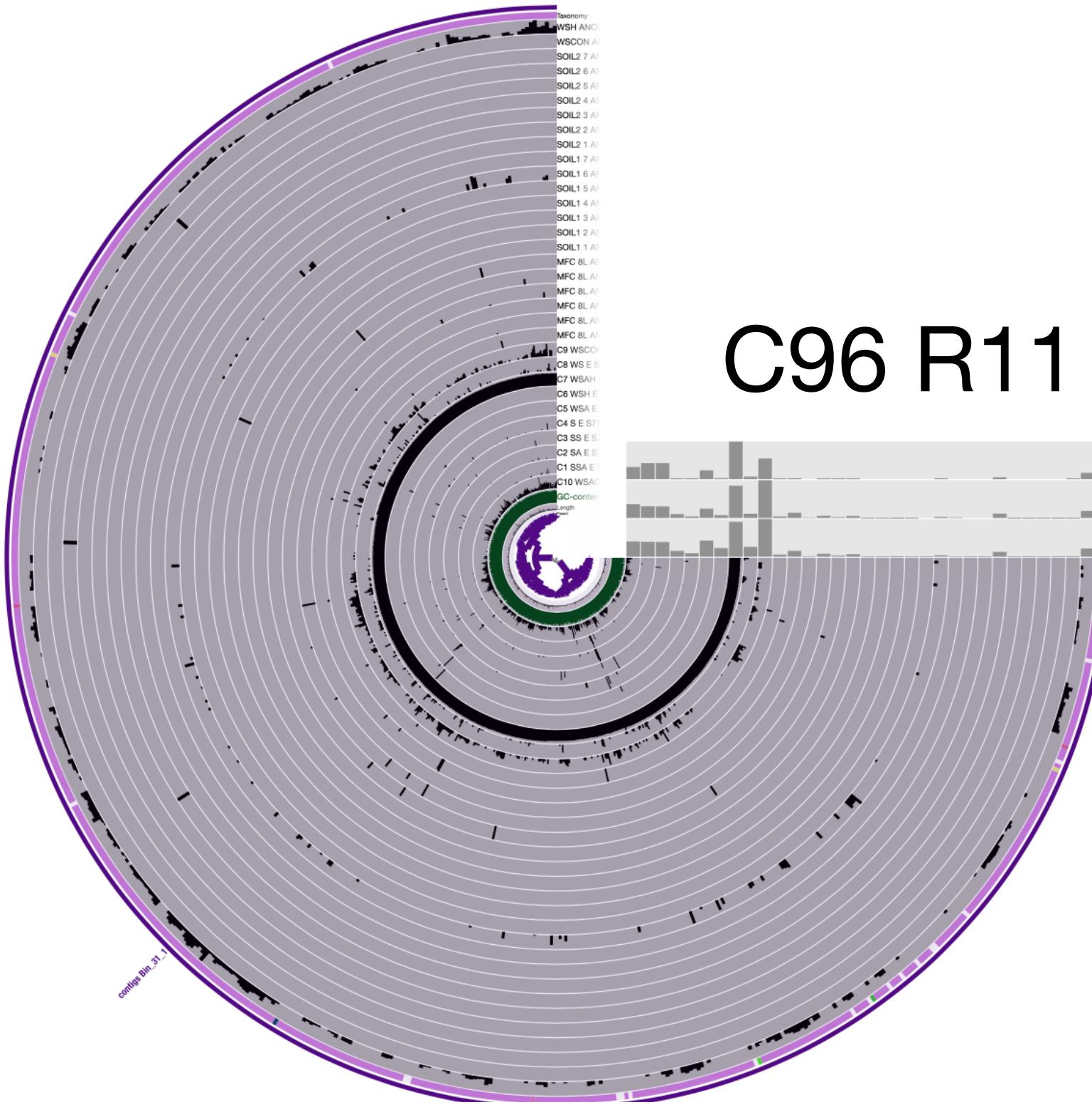
X

Items order: Differential coverage (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom



Refining contigs_Bin_31 from "collection1"

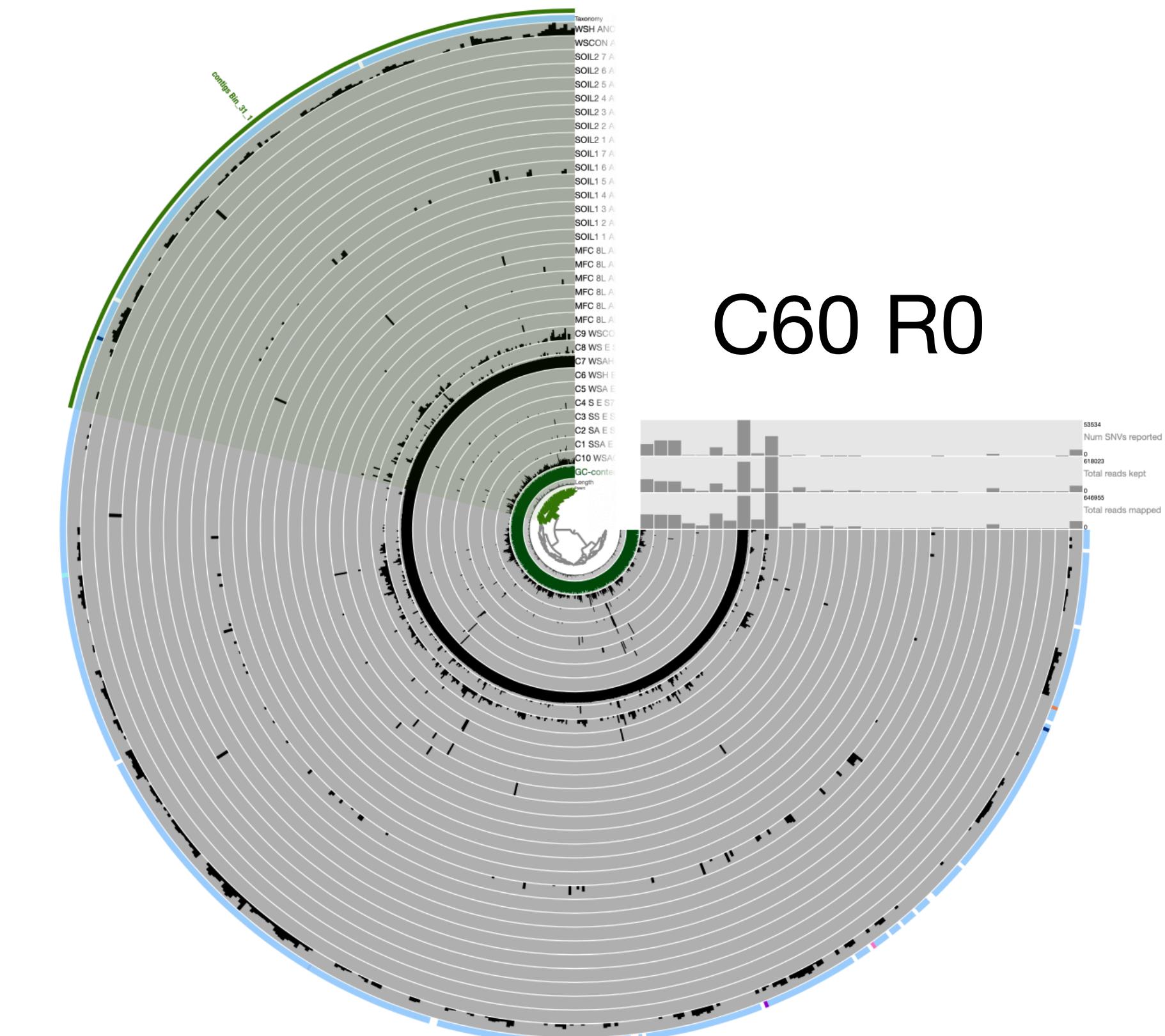
Items order: Differential coverage (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom



C96 R11

Refining contigs_Bin_31 from "collection1"

Items order: Differential coverage (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom



C60 R0

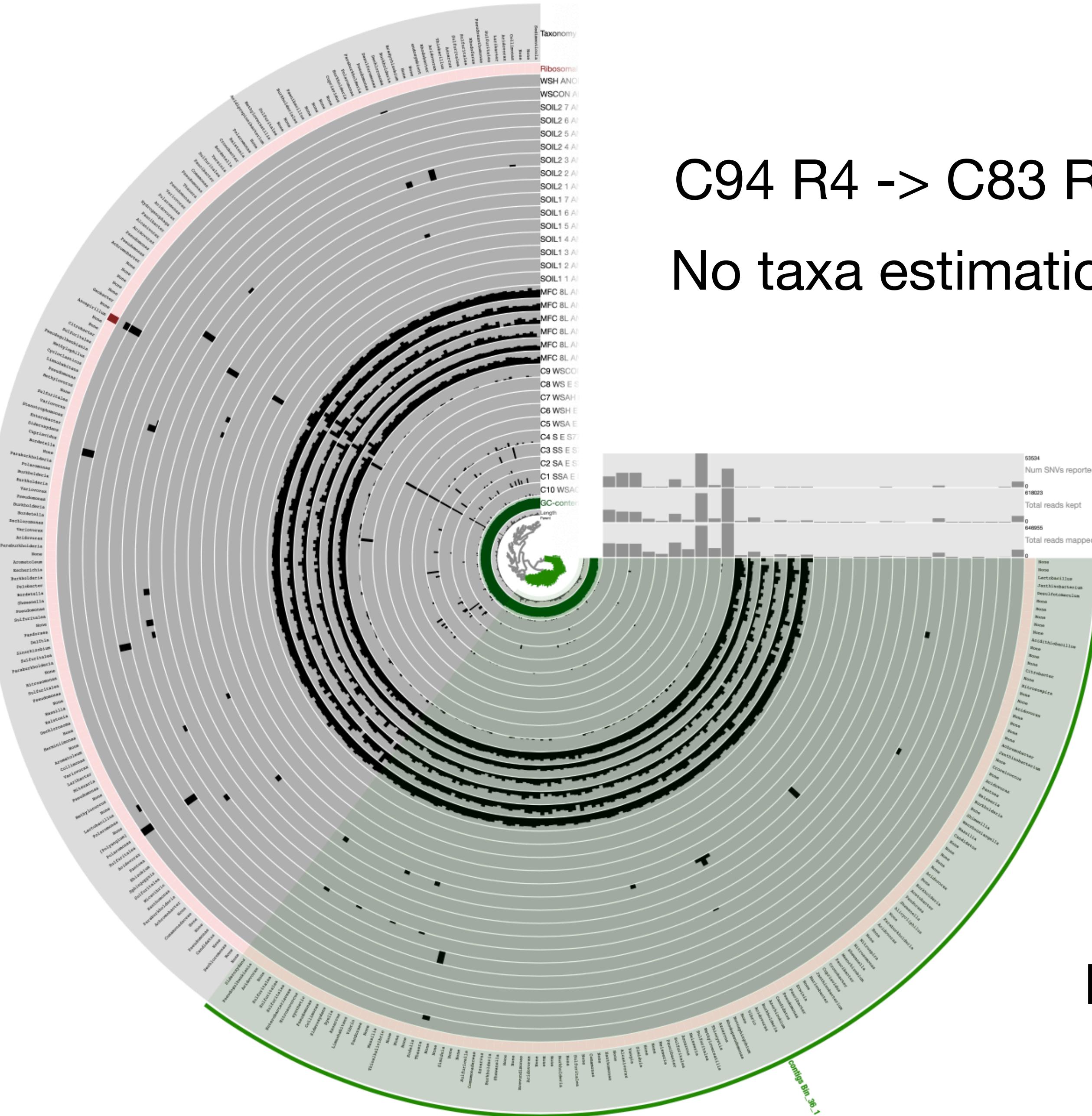
Geobacter

bin_60_0

+ bin_41_6

Refining contigs_Bin_36 from "collection1"

Items order: Sequence composition (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom



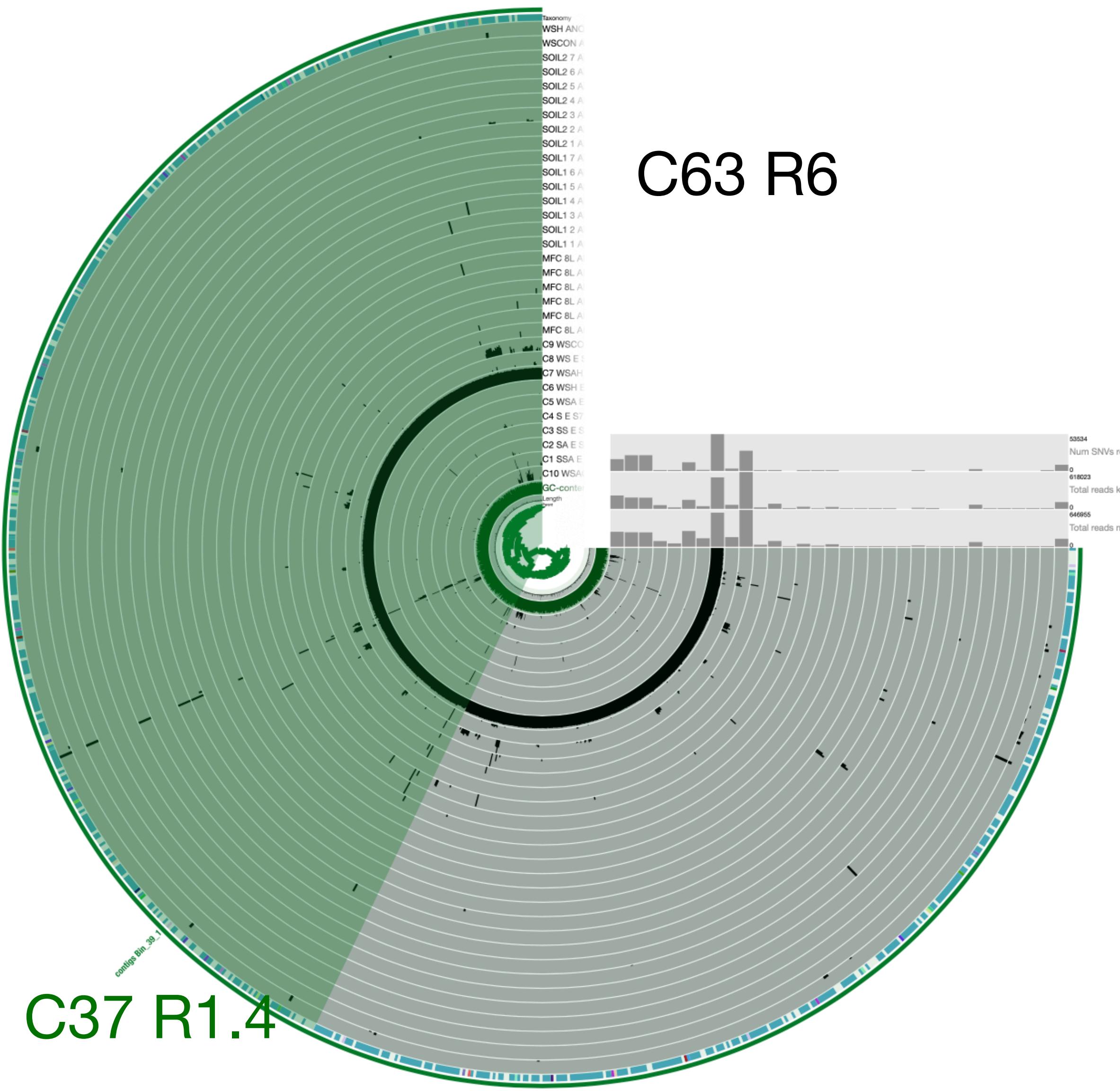
C94 R4 -> C83 R3

No taxa estimation

bin_83_2

Refining contigs_Bin_39 from "collection1"

Items order: **Differential coverage (D: Euclidean; L: Ward)** | Current view: **mean_coverage** | Sample order: **custom**

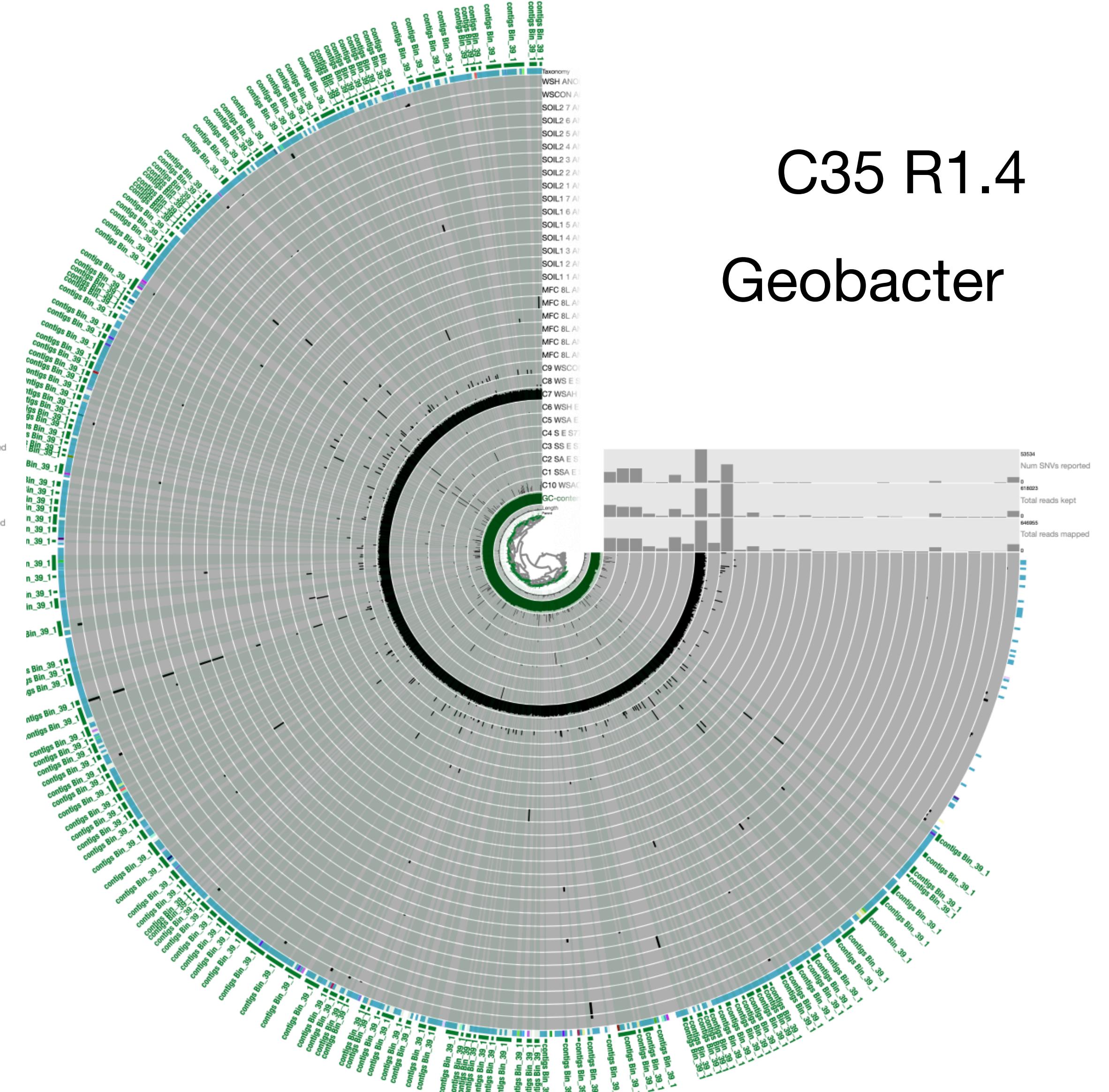


C63 R6

C37 R1.4

Refining contigs_Bin_39 from "collection1"

Items order: **Sequence composition (D: Euclidean; L: Ward)** | Current view: **mean_coverage** | Sample order: **custom**



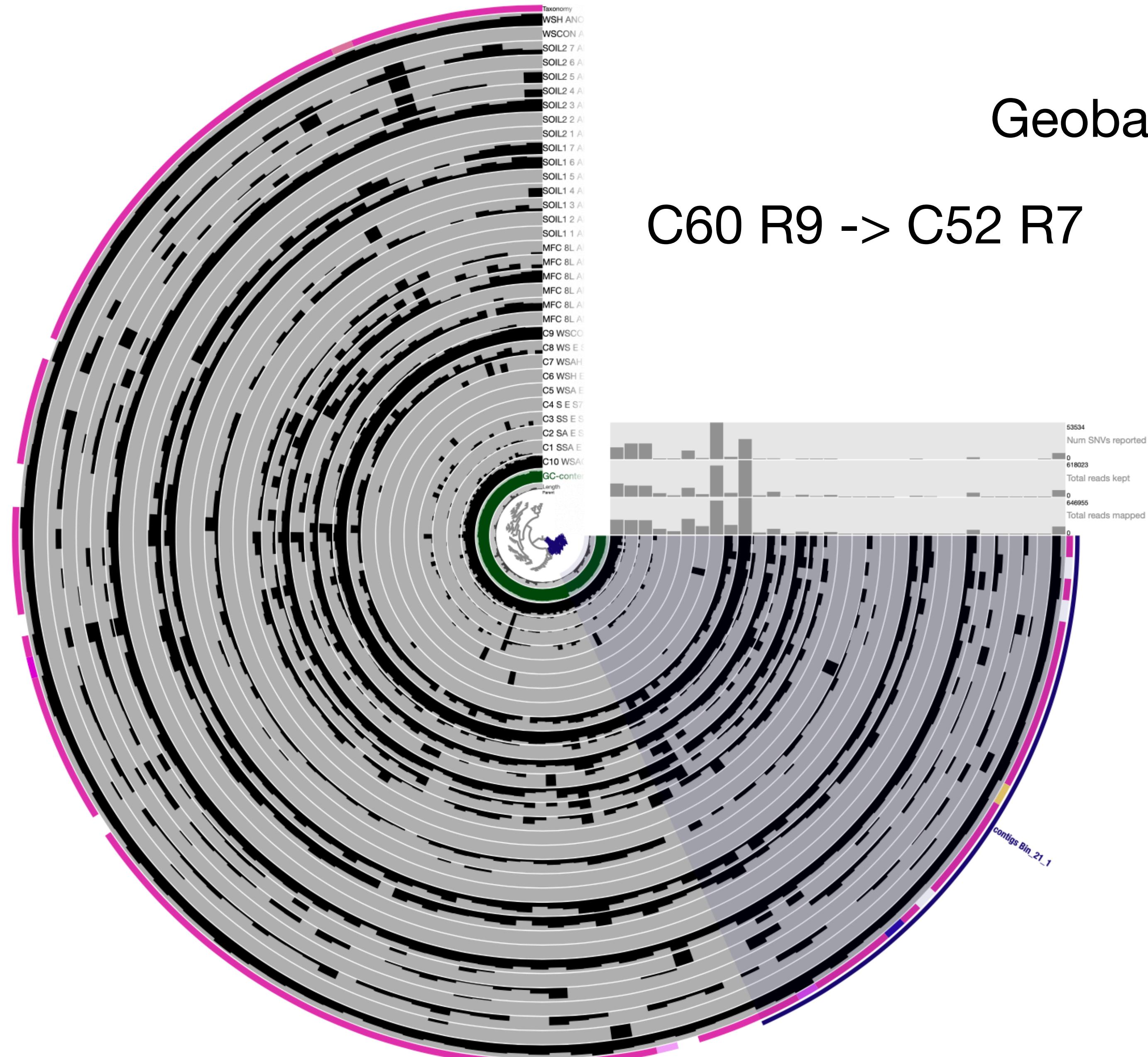
C35 R1.4

Geobacter

bin_35_1

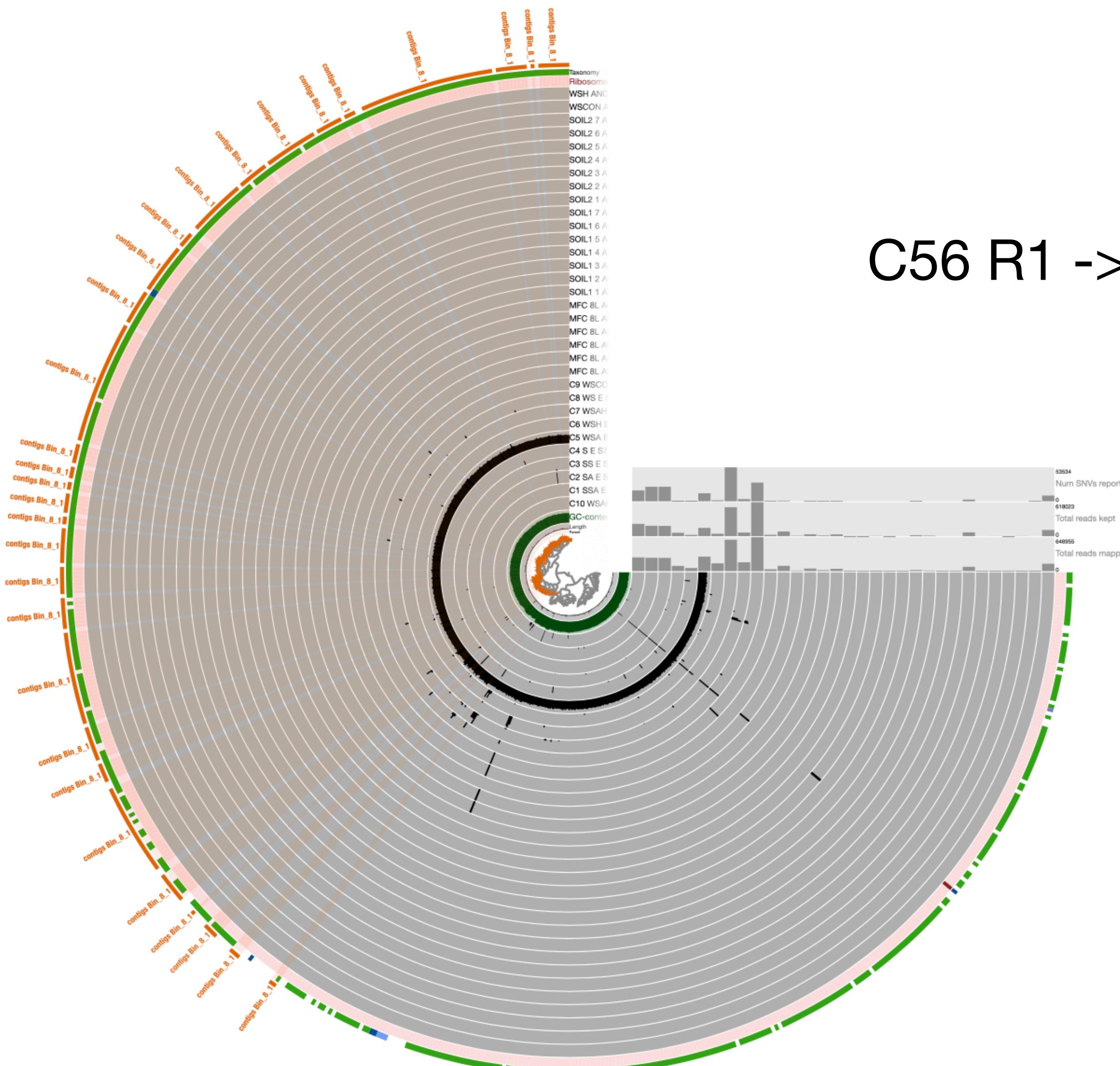
Refining contigs_Bin_21 from "collection1"

Items order: Sequence composition (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom



Refining contigs_Bin_8 from "collection1"

Items order: **Sequence composition (D: Euclidean; L: Ward)** | Current view: **mean_coverage** | Sample order: **custom**



C56 R1 → C31 R0

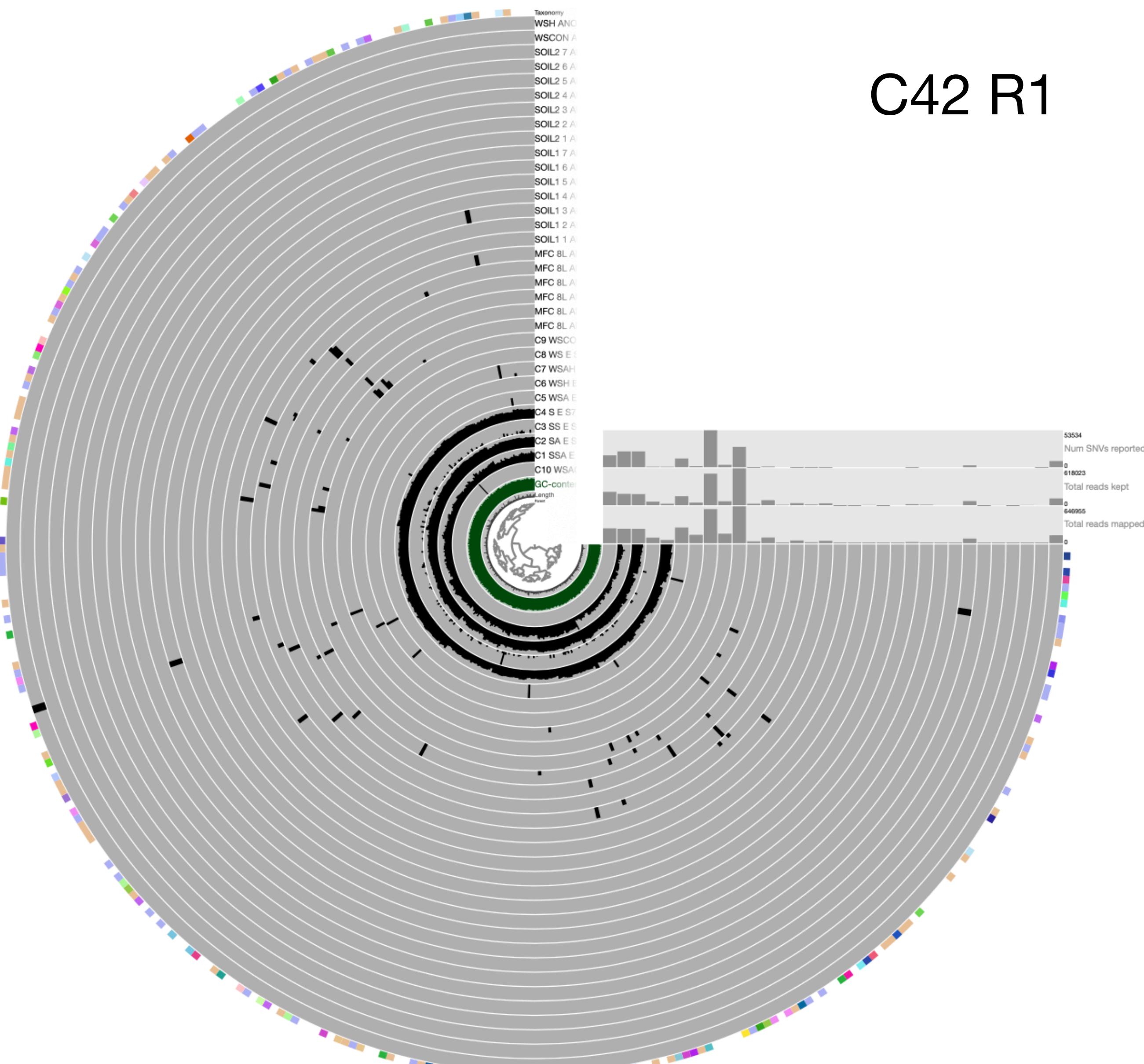
Desulfobacterium

bin_31_0 ?

Refining contigs_Bin_5_1 from "collection1"

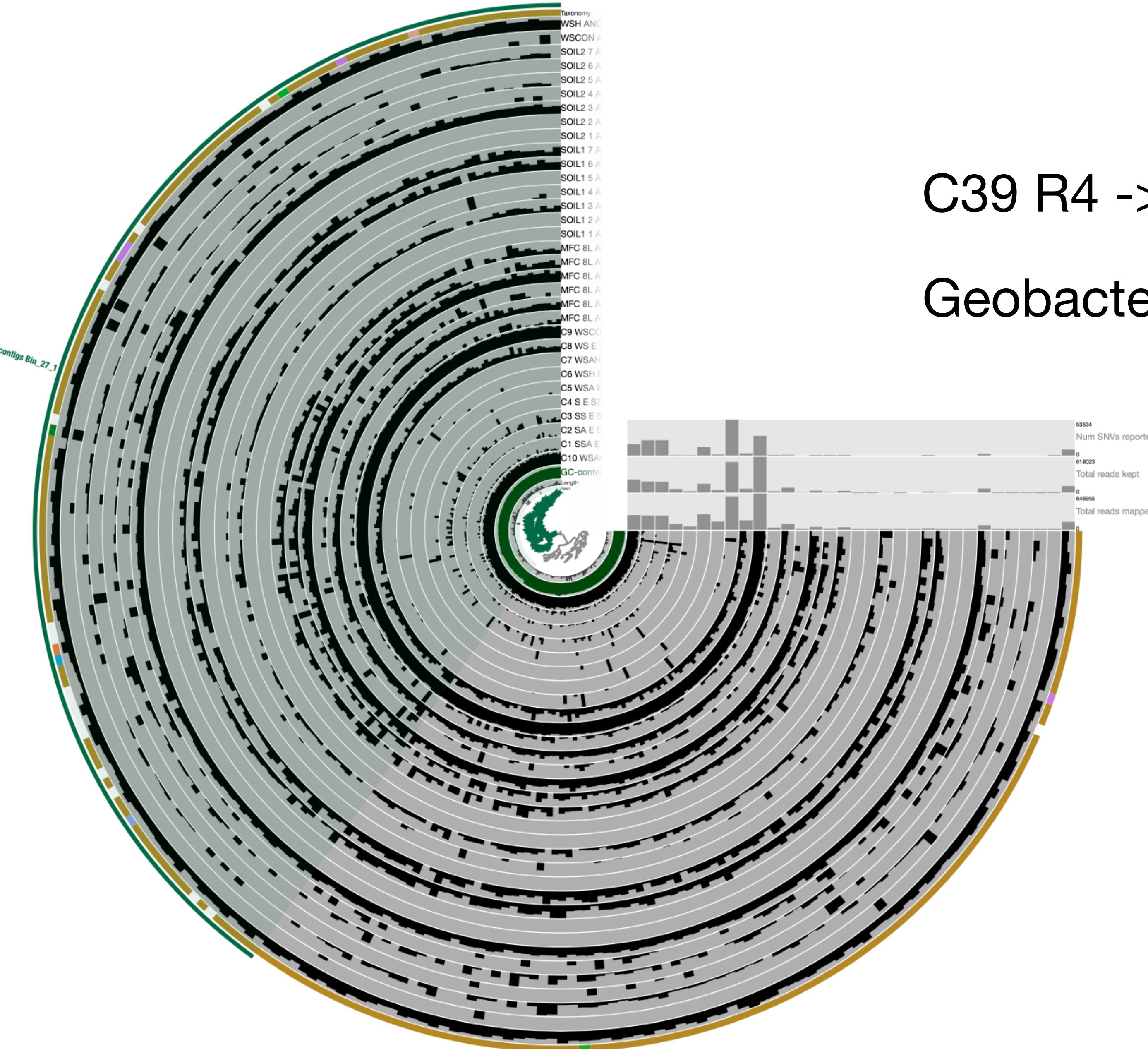
Items order: Differential coverage (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom

C42 R1



Refining contigs_Bin_27 from "collection1"

Items order: Sequence composition (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom



C39 R4 -> C32 R3

Geobacter

53534
Num SNVs reported
0
618023
Total reads kept
0
646955
Total reads mapped
0

Refining contigs_Bin_13 from "collection1"

Items order: Sequence composition (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom

C80 R7 -> C39 R4

Desulfoboromas and geobacter...

