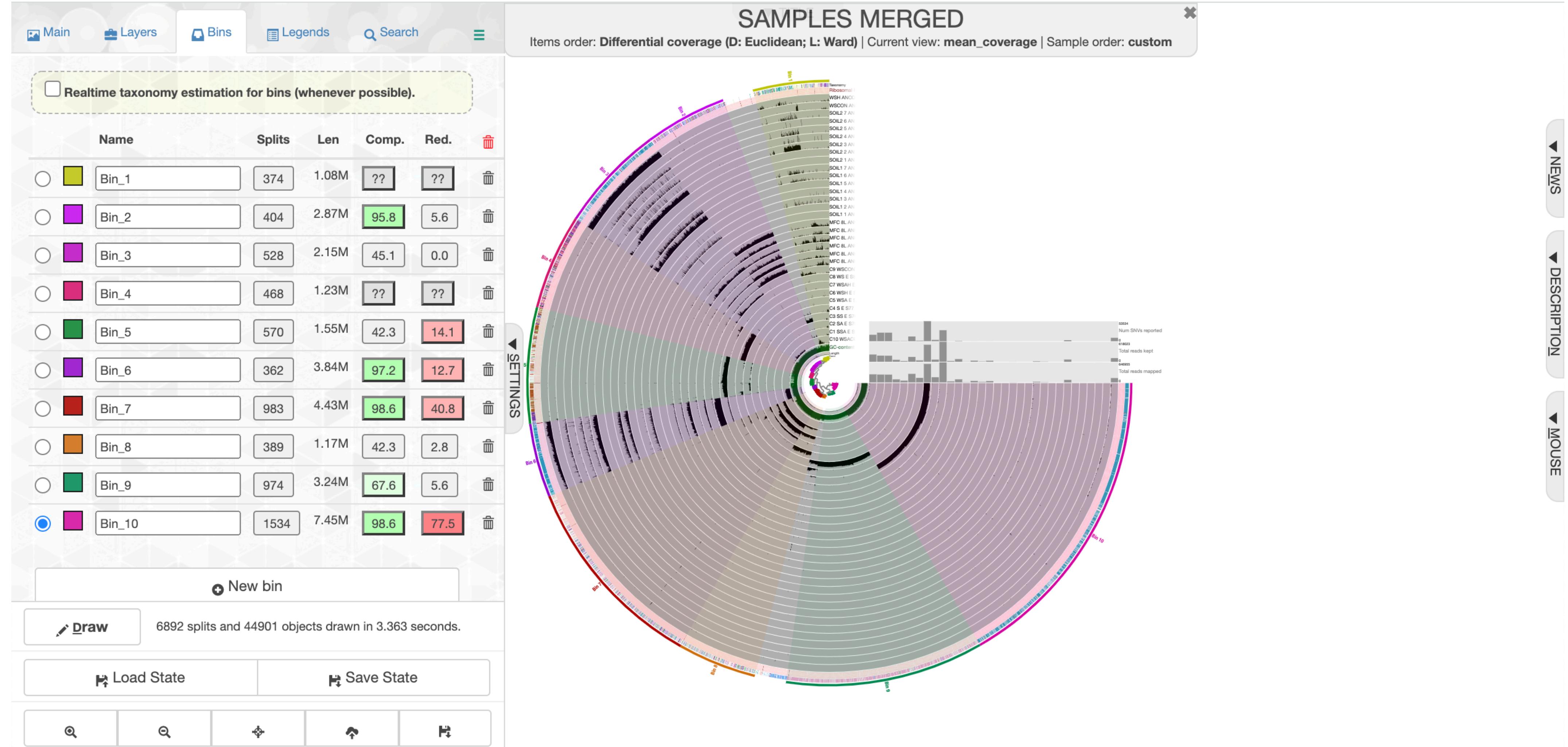
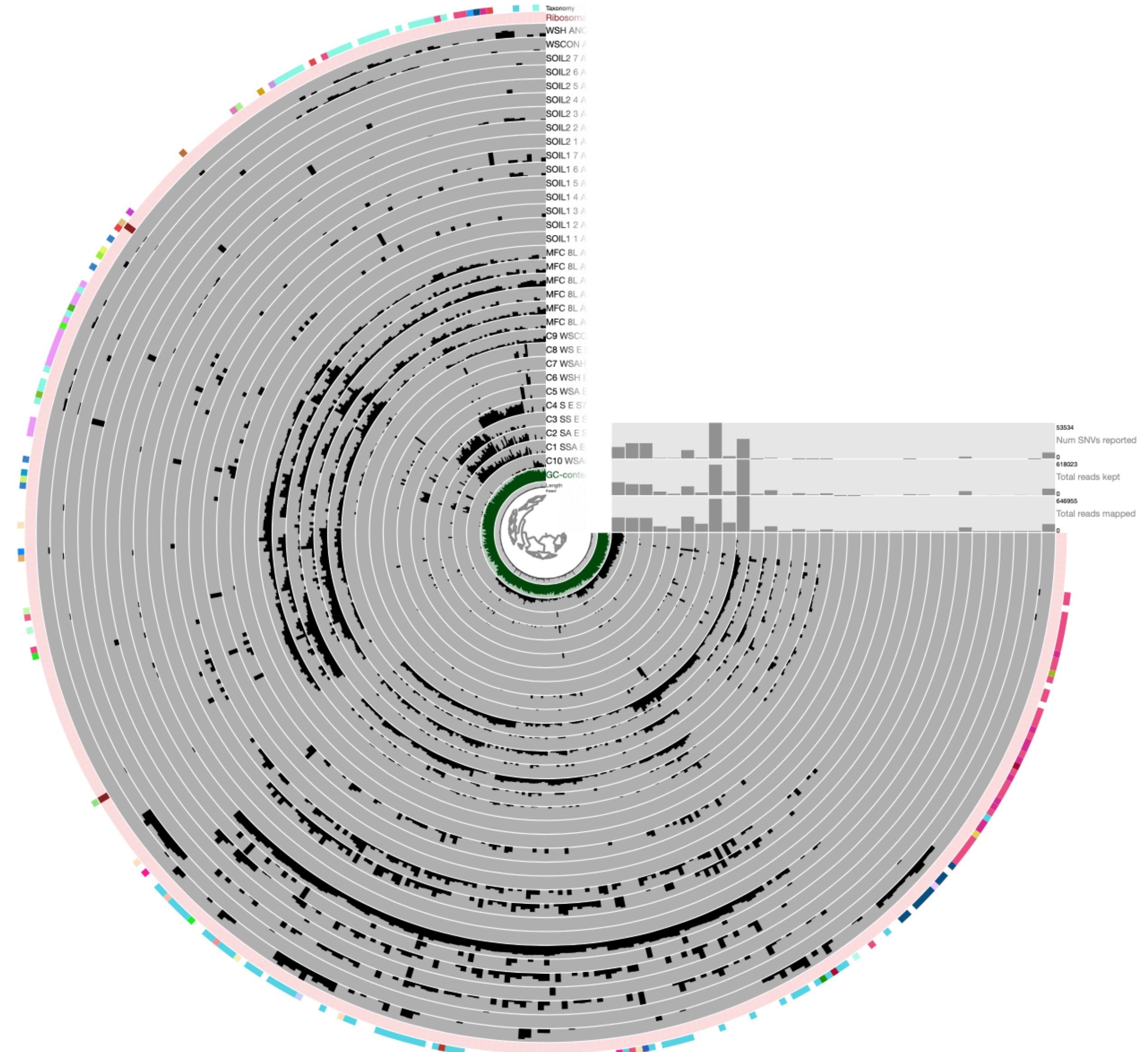


My bins 3



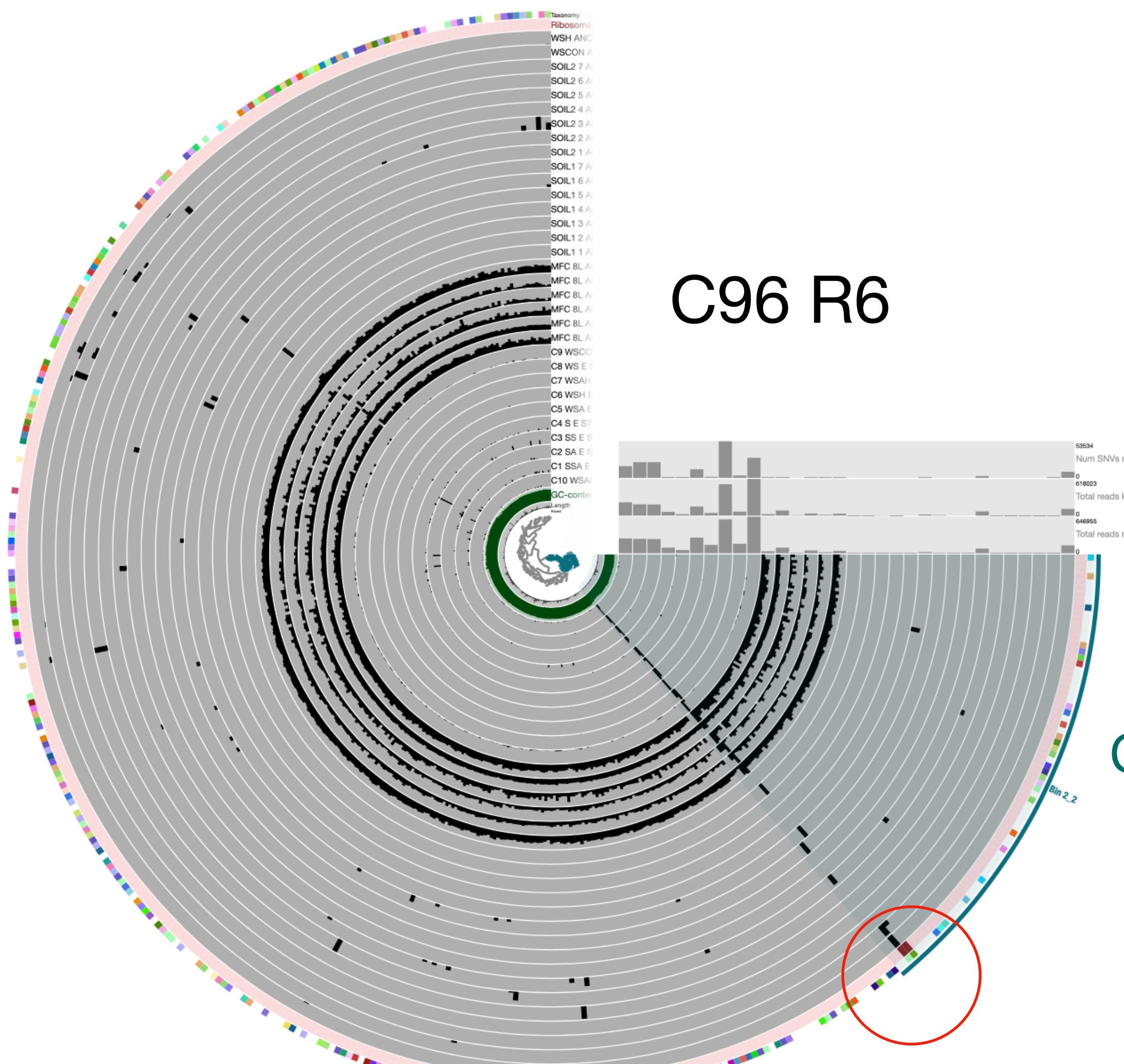
Refining Bin_1 from "collection4"

Items order: Seq. Composition + Diff. Coverage (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom



Refining Bin_2 from "collection4"

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



Sequences producing significant alignments:

Select: All None Selected:0

Alignments Download GenBank Graphics Distance tree of results

Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
Azospira sp. I09 DNA, complete genome	4438	17754	100%	0.0	92.98%	AP021844.1
Dechlorosoma suillum PS, complete genome	4433	8866	100%	0.0	92.95%	CP003153.1
Uncultured bacterium clone fosmid K823 genomic sequence	3960	4240	84%	0.0	96.99%	KX576361.1
Denitratisoma sp. DHT3 genome	3243	11184	90%	0.0	91.63%	CP020914.1
Sterolibacterium denitrificans strain Chol genome assembly, chromosome: SDENCHOL	3195	3556	87%	0.0	91.15%	LT837803.1
Azospira oryzae strain PS 23S ribosomal RNA gene, complete sequence	3169	3169	76%	0.0	91.16%	NR_076936.1

Sequences producing significant alignments:

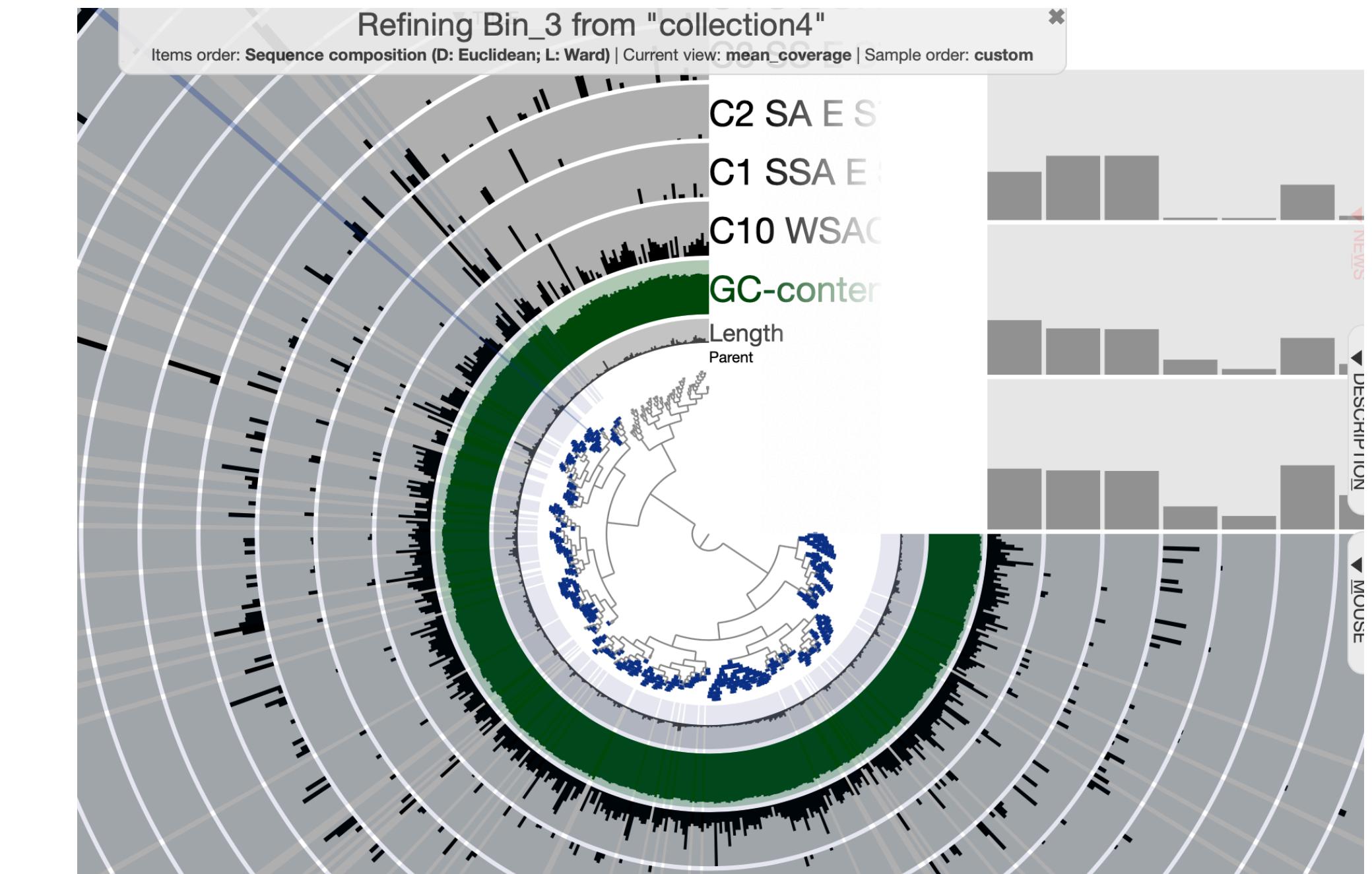
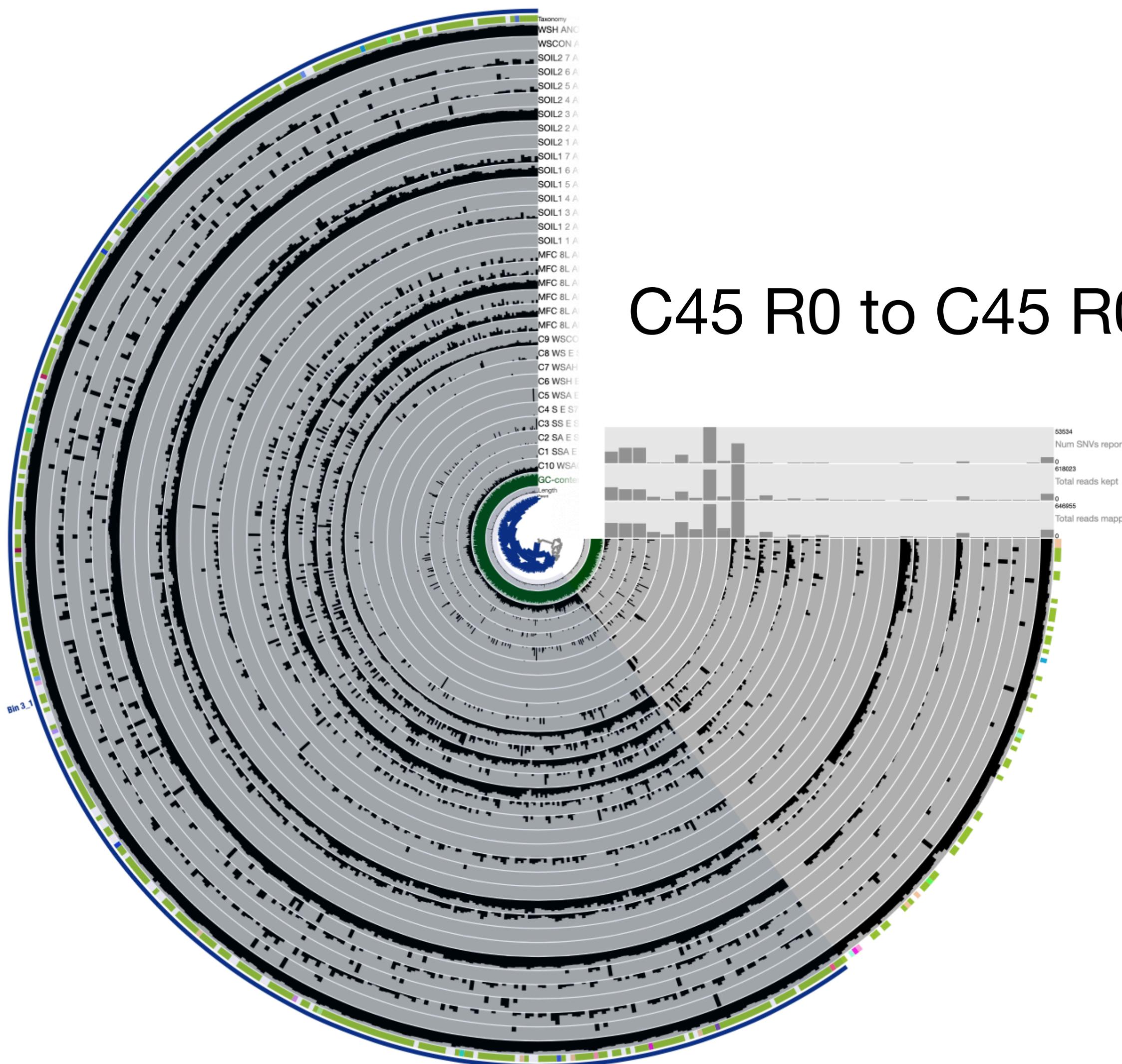
Select: All None Selected:0

Alignments Download GenBank Graphics Distance tree of results

Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
Rugobacter aromaticivorans strain Ca6, complete genome	2918	8878	91%	0.0	93.25%	CP010554.1
Uncultured bacterium clone 9_shallow 16S ribosomal RNA gene, partial sequence	2782	2782	46%	0.0	97.44%	MF942658.1
Uncultured bacterium clone UA_23 16S ribosomal RNA gene, partial sequence	2780	2780	42%	0.0	99.87%	JX120393.1
Uncultured bacterium clone LYC178 16S ribosomal RNA gene, partial sequence	2780	2780	42%	0.0	99.74%	DQ984610.1
Uncultured bacterium clone M17-10-B10 16S ribosomal RNA gene, partial sequence	2774	2774	42%	0.0	99.80%	JQ088386.1
Uncultured bacterium clone P2C10 16S ribosomal RNA gene, partial sequence	2702	2702	42%	0.0	99.07%	KP636125.1
Uncultured bacterium clone P2B5 16S ribosomal RNA gene, partial sequence	2682	2682	42%	0.0	99.00%	KP636120.1
Rugobacter aromaticivorans strain Ca6 16S ribosomal RNA, complete sequence	2627	2627	43%	0.0	97.07%	NR_156019.1

Refining Bin_3 from "collection4"

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

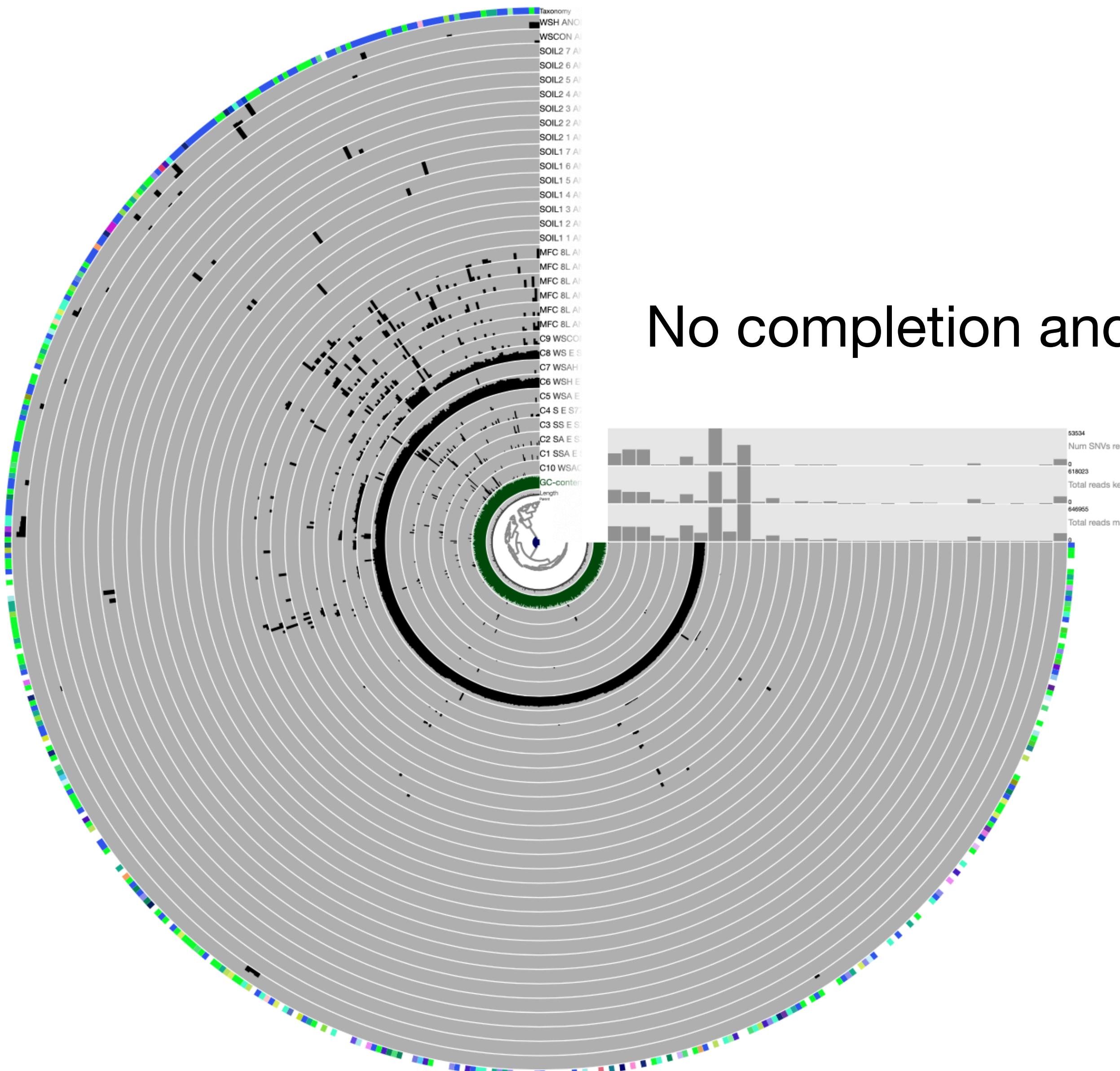


To C31 R0

Geobacter

Refining Bin_4 from "collection4"

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



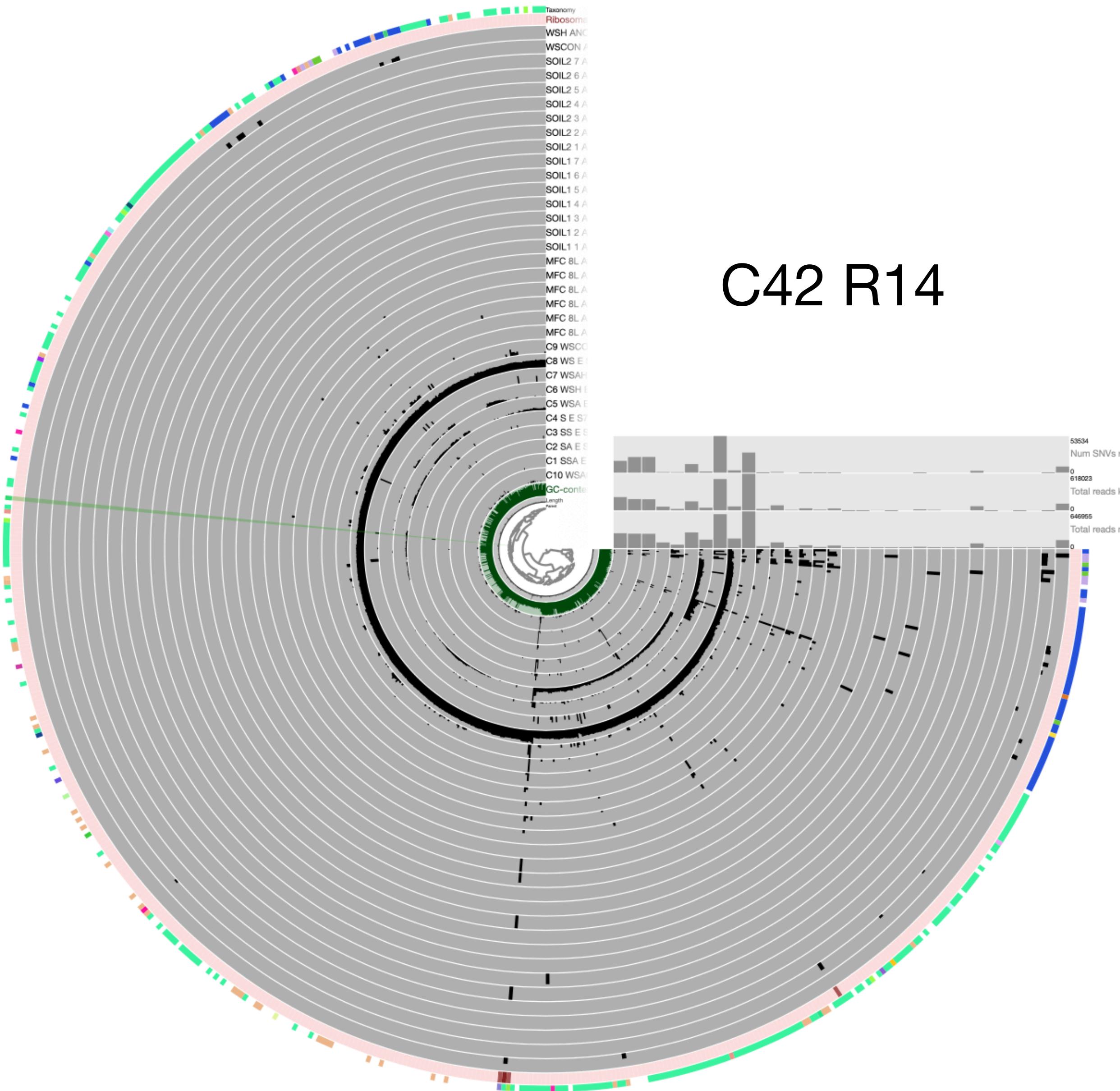
No completion and redundancy estimation



Refining Bin_5 from "collection4"

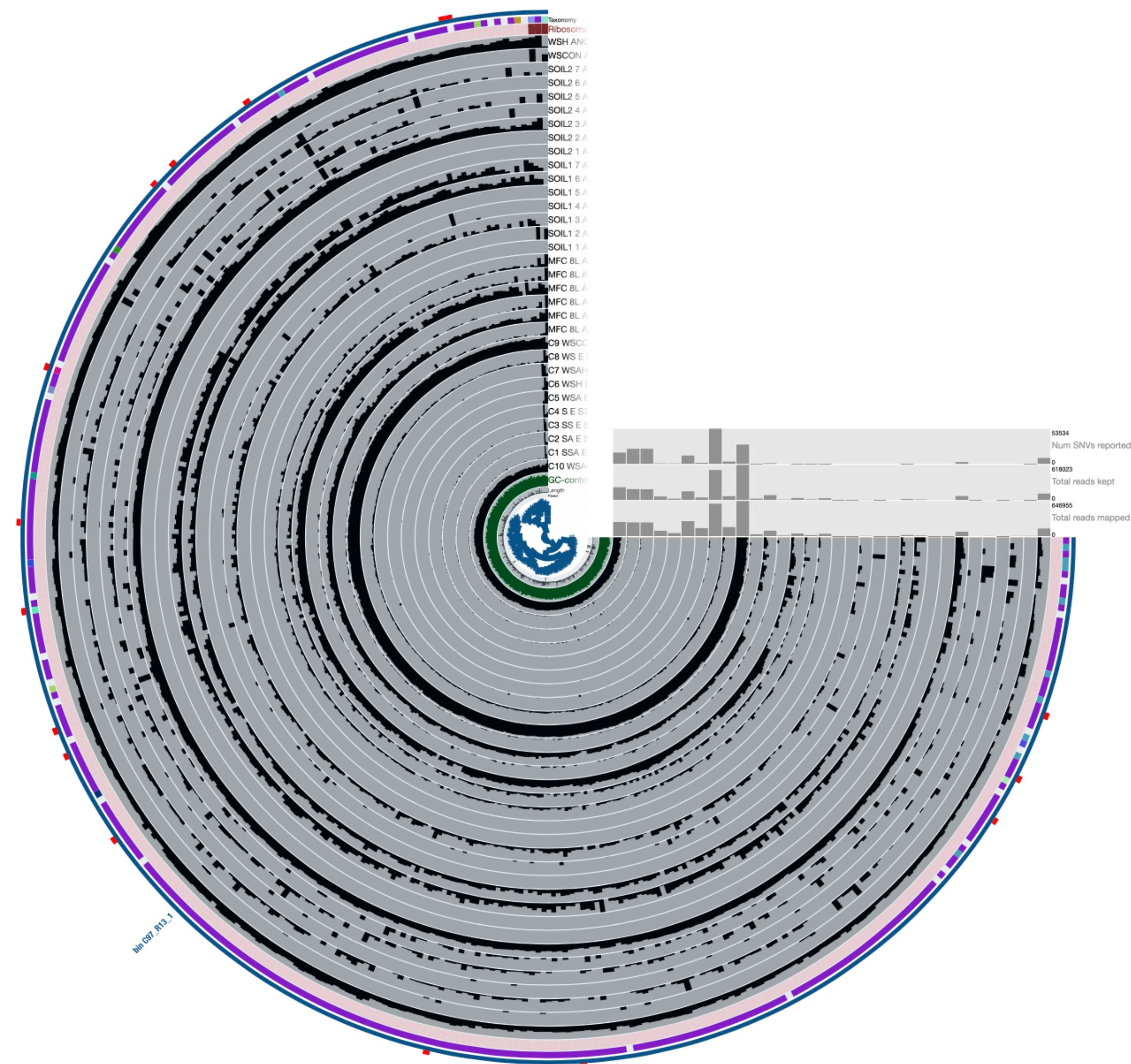
Items order: Differential coverage (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom

C42 R14



Refining bin_C97_R13 from "collection4"

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

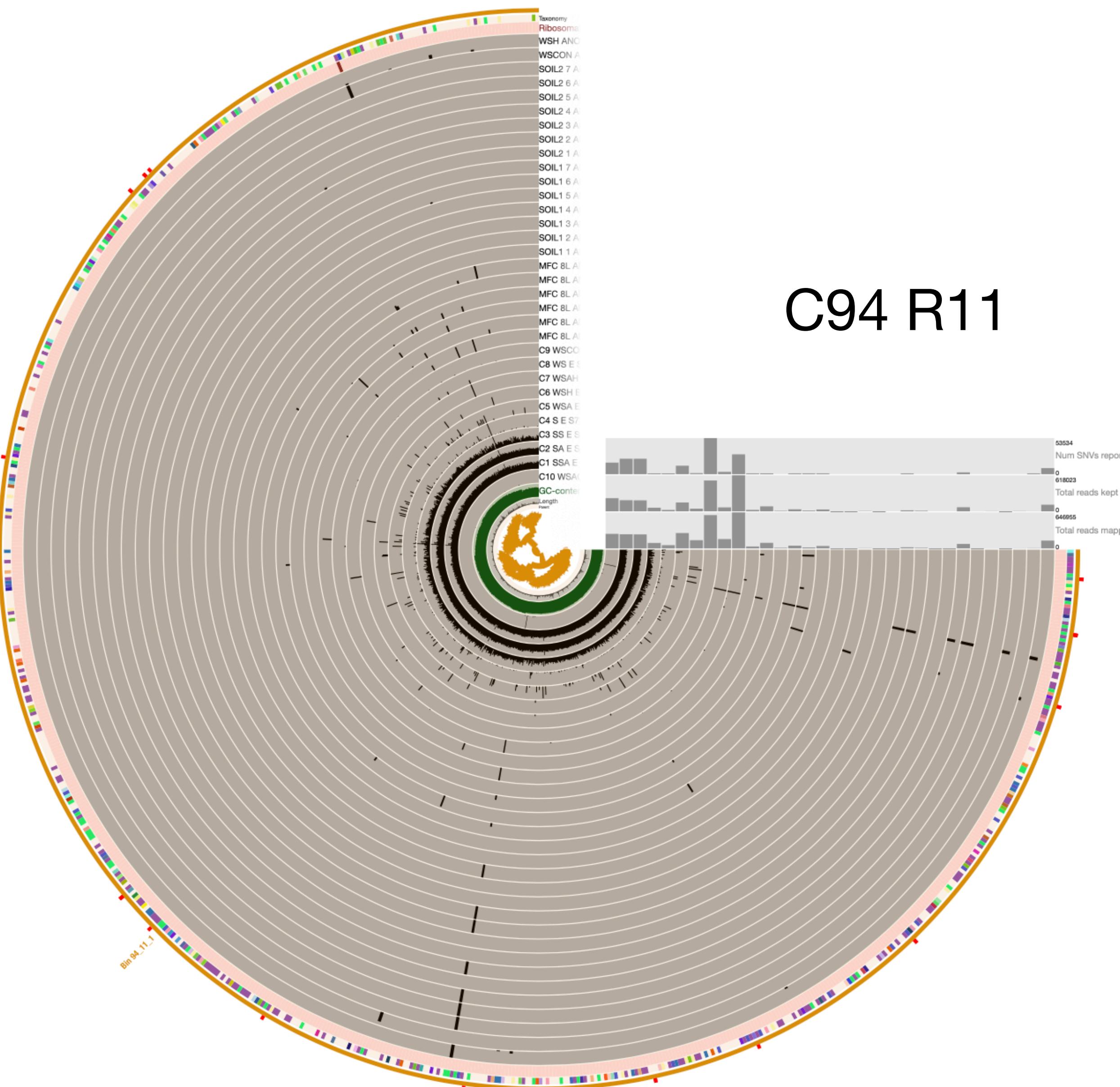


Geobacter

Refining Bin_94_11 from "collection4"

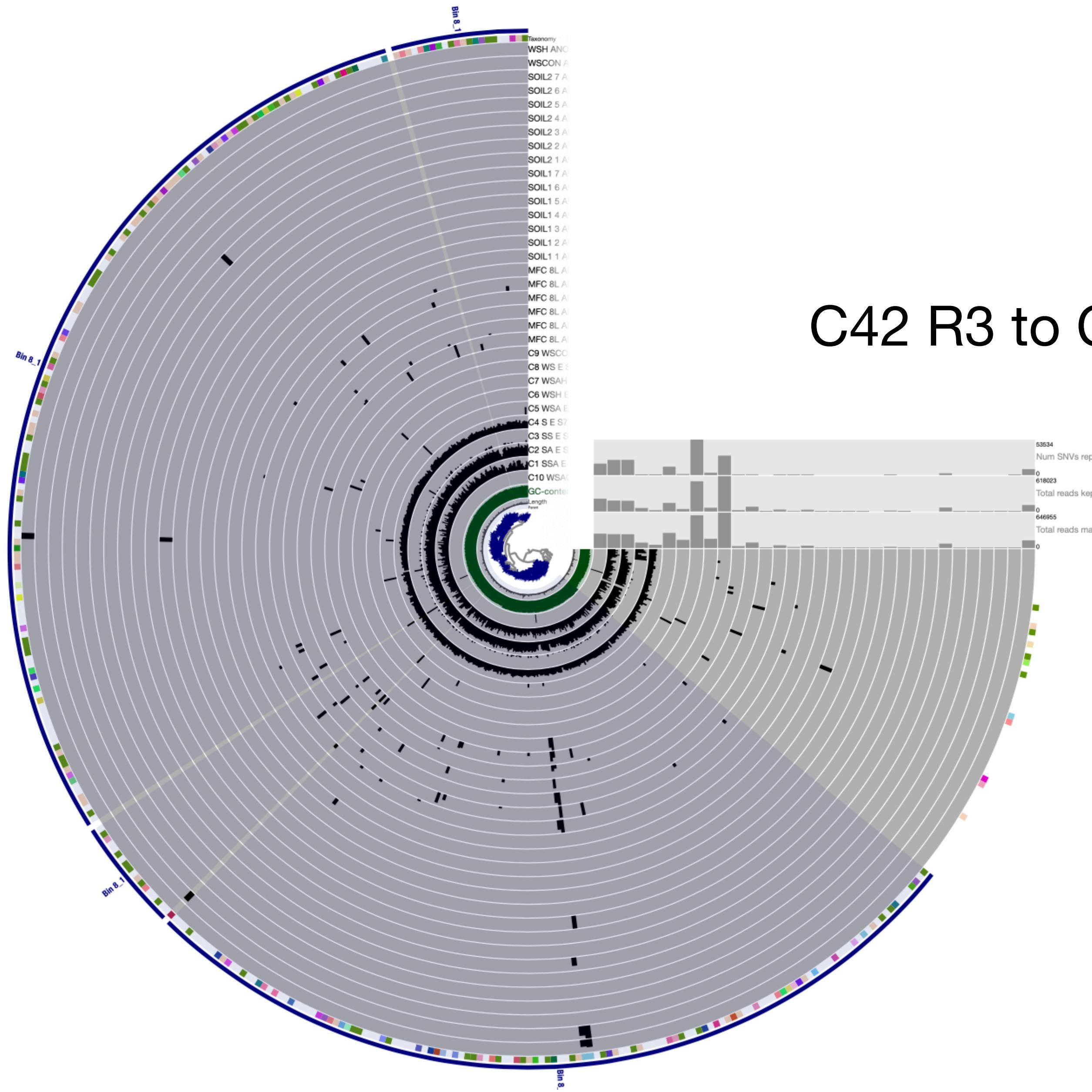
Items order: Sequence composition (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom

C94 R11



Refining Bin_8 from "collection4"

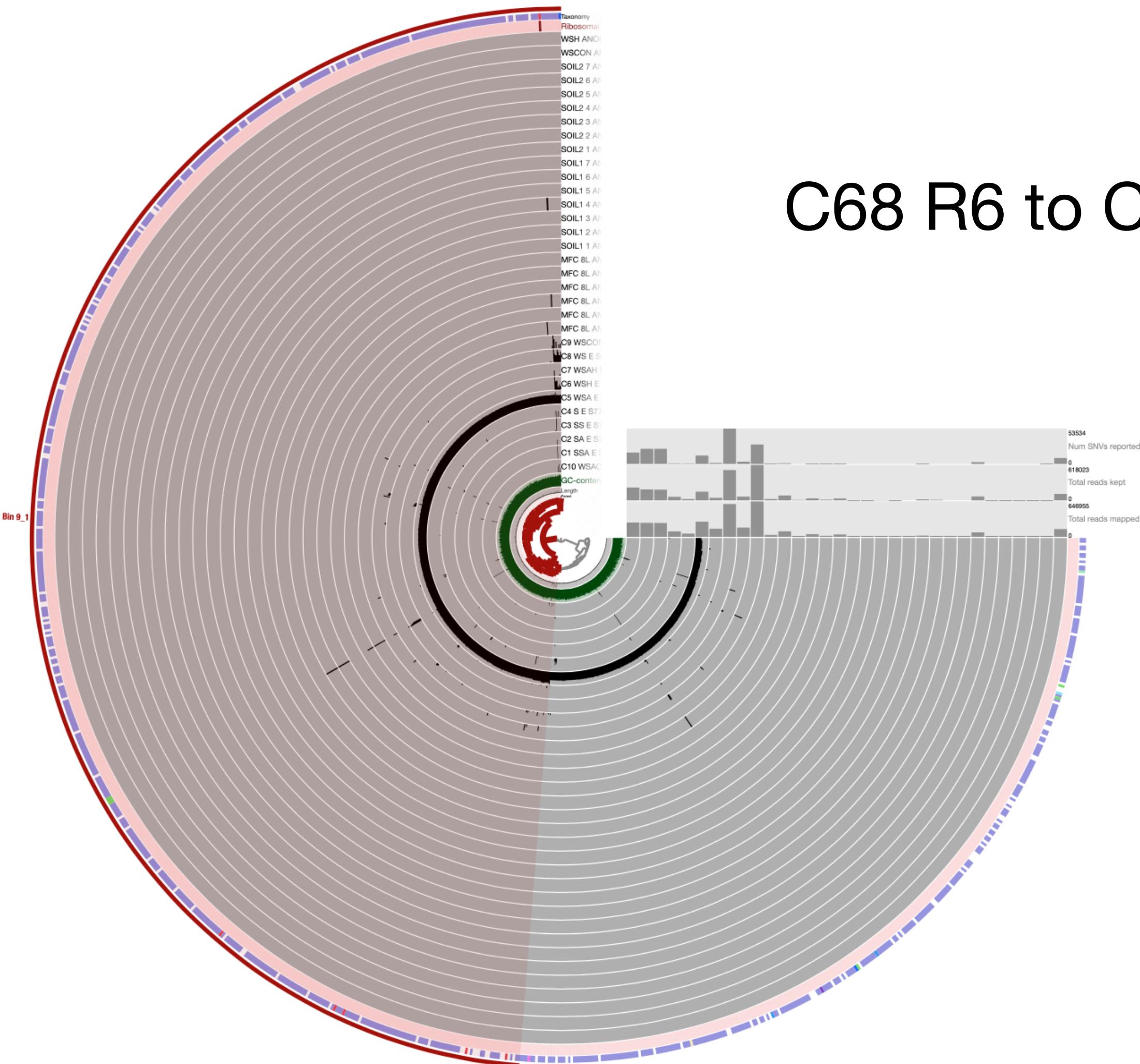
Items order: Sequence composition (D: Euclidean; L: Ward) | Current view: mean_coverage | Sample order: custom



C42 R3 to C42 R1

Refining Bin_9 from "collection4"

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

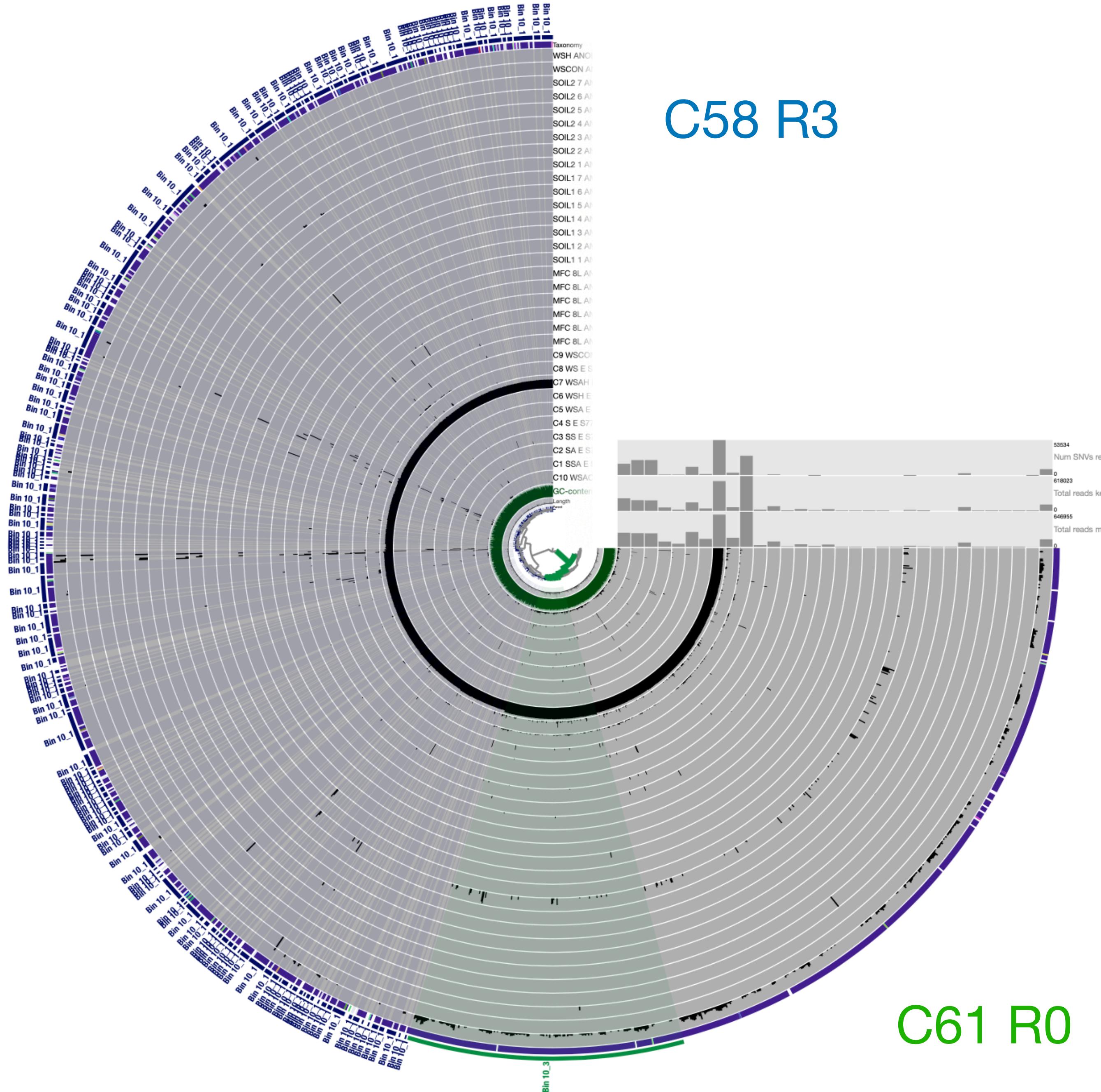


C68 R6 to C58 R1

Desulfobacterium

Refining Bin_10 from "collection4"

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



C61 R0

Geobacter

