

MAG

OdNW_18-Q3-R42-56_BAT3C.134_sub

Skiv_18-Q3-R9-52_BATAC.176

Skiv_18-Q3-R9-52_BATAC.396

Skiv_18-Q3-R9-52_MAXAC.078_sub

Vibo_18-Q3-R45-57_BAT3C.98

bcsABII-A

bcsCII

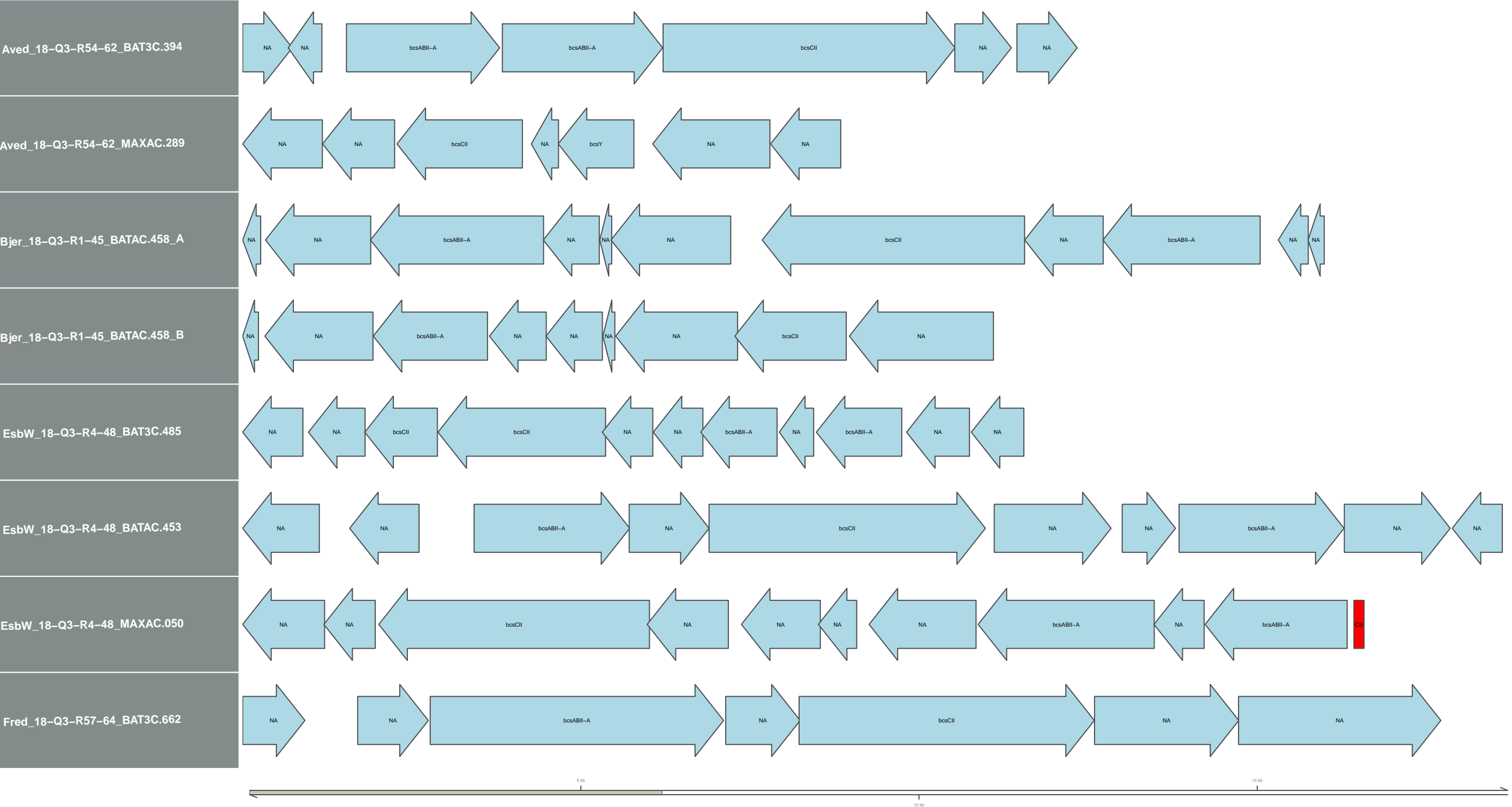
bcsY

Percent Identity

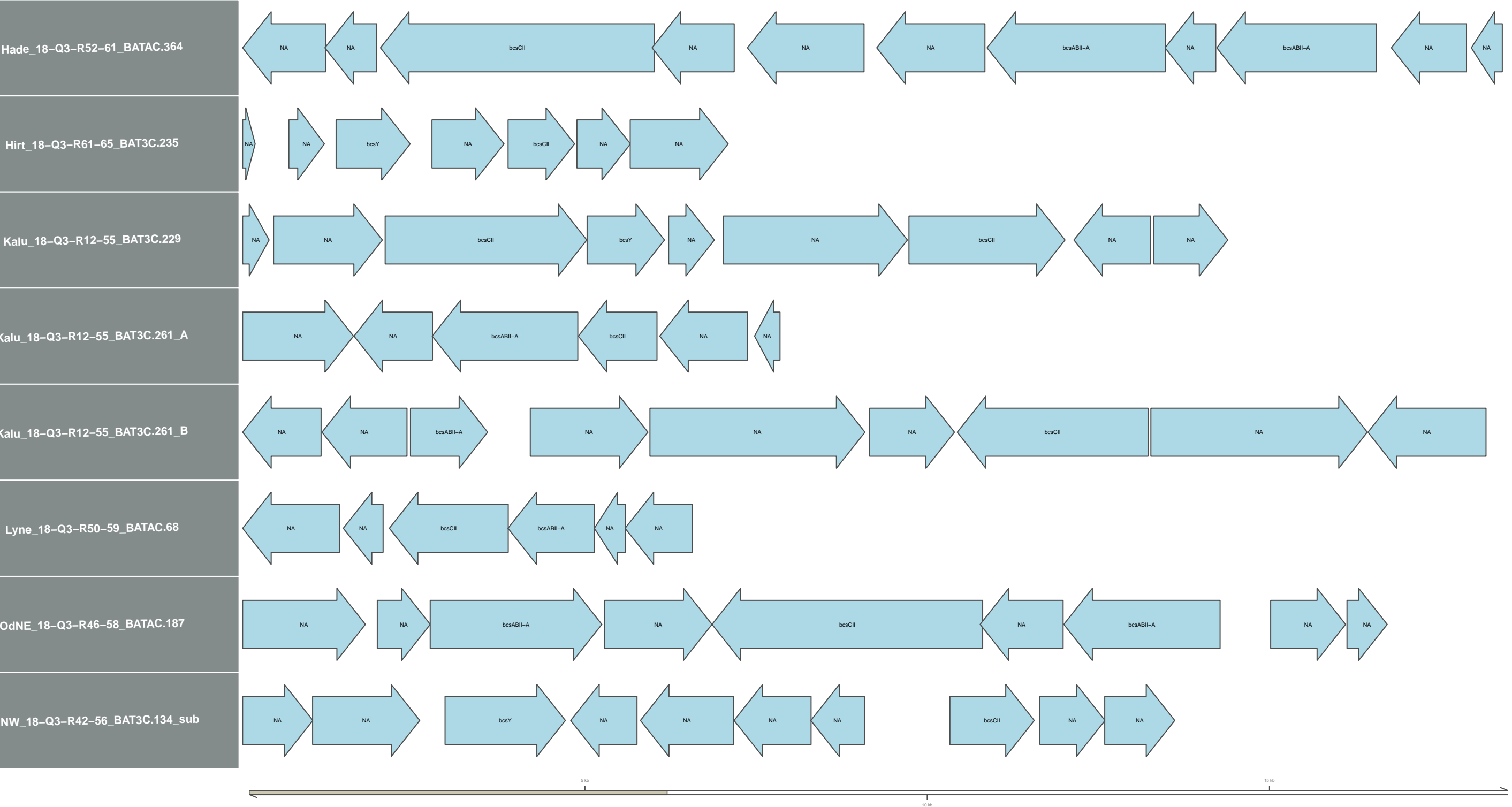
30

25

cellulose2

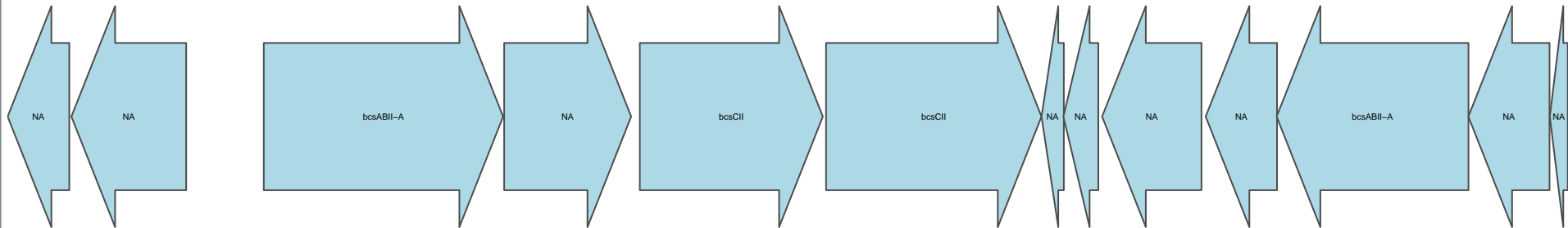


cellulose2

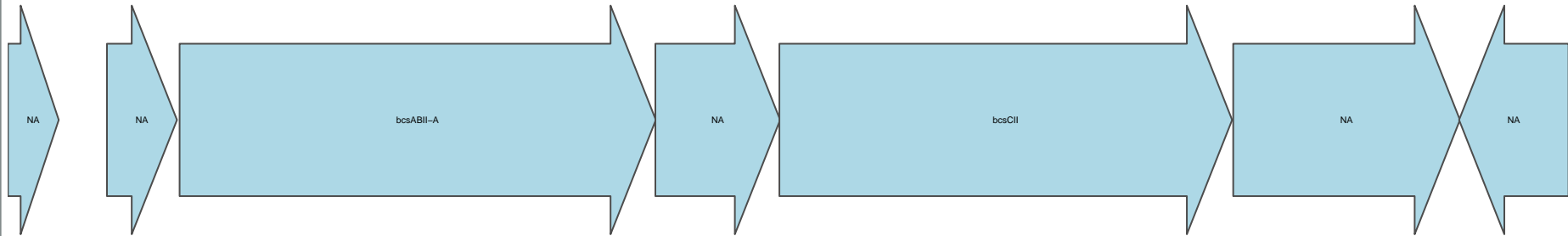


cellulose2

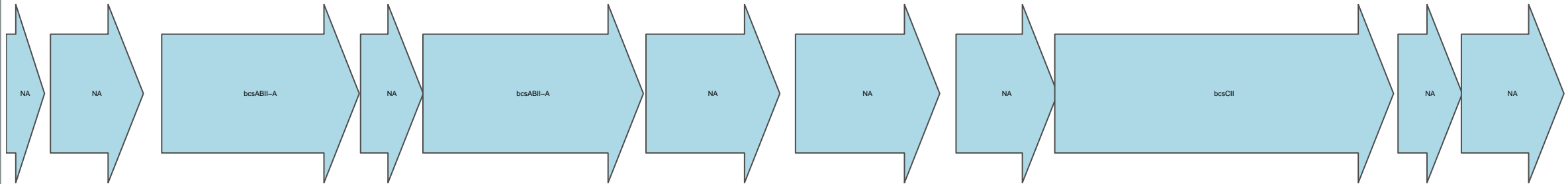
Skiv_18-Q3-R9-52_BATAC.176



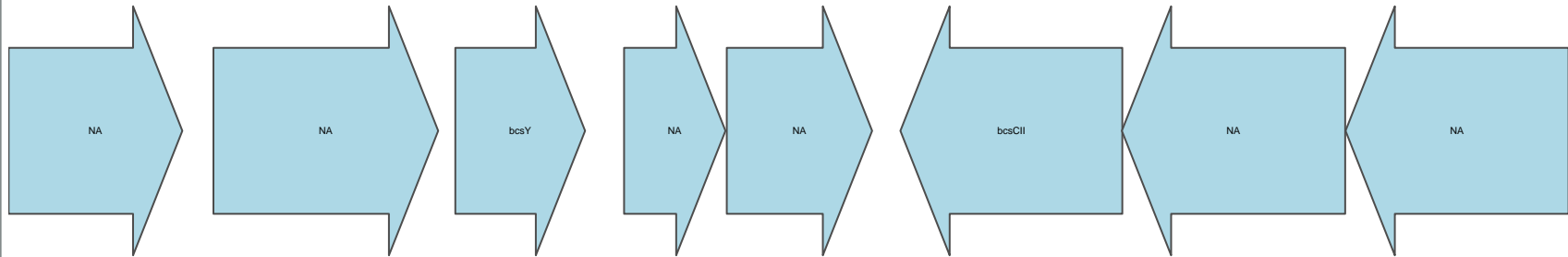
Skiv_18-Q3-R9-52_BATAC.396



Skiv_18-Q3-R9-52_MAXAC.078_sub



Vibo_18-Q3-R45-57_BAT3C.98



	ID	GTDBTax	seqname	start	Query_Label	ProkkaNO
1	Aved_18-Q3-R54-62_BAT3C.394	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Burkholderiaceae;g__PHCI01;s__	tig01608118-10-14295950	486433	bcsABII-A	03920
2	Aved_18-Q3-R54-62_MAXAC.289	d__Bacteria;p__Myxococcota;c__Polyangia;o__GCA-2747355;f__GCA-2747355;g__s__	tig00007292-10-5178180	72477	bcsCII	00090
3	Bjer_18-Q3-R1-45_BATAC.458	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Burkholderiaceae;g__Rhodoferax;s__	tig00001862-10-6955040	315548	bcsABII-A	00263
4	EsbW_18-Q3-R4-48_BAT3C.485	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Burkholderiaceae;g__PHCI01;s__	tig00020141-10-609230	21551	bcsABII-A	03736
5	EsbW_18-Q3-R4-48_BATAC.453	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Burkholderiaceae;g__Simplicispira_A;s__	tig00014047-10-1285900	106609	bcsABII-A	01813
6	EsbW_18-Q3-R4-48_MAXAC.050	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Rhodocyclaceae;g__s__	tig00018082-10-889440	86803	bcsABII-A	03085
7	Fred_18-Q3-R57-64_BAT3C.662	d__Bacteria;p__Acidobacteriota;c__Acidobacteriae;o__Bryobacterales;f__Bryobacteraceae;g__s__	tig00006256-10-5425100	396711	bcsABII-A	01773
8	Hade_18-Q3-R52-61_BATAC.364	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Rhodocyclaceae;g__Dechloromonas;s__	tig00407471-10-3182230	149681	bcsABII-A	03002
9	Hirt_18-Q3-R61-65_BAT3C.235	d__Bacteria;p__Verrucomicrobiota;c__Verrucomicrobiae;o__Opitutales;f__Opitutaceae;g__s__	tig00023131-10-926540	45276	bcsCII	03485
10	Kalu_18-Q3-R12-55_BAT3C.229	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Palsa-1005;g__PALSA-1003;s__	tig00100981-10-4584110	168222	bcsCII	03736
11	Kalu_18-Q3-R12-55_BAT3C.261	d__Bacteria;p__Acidobacteriota;c__Acidobacteriae;o__Bryobacterales;f__Bryobacteraceae;g__s__	tig00004381-10-1940160	135345	bcsABII-A	03277
12	Lyne_18-Q3-R50-59_BATAC.68	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Burkholderiaceae;g__Rubrivivax;s__	tig00007586-10-3042350	69843	bcsABII-A	02354
13	OdNE_18-Q3-R46-58_BATAC.187	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Gallionellaceae;g__UBA7399;s__	tig00013603-10-1108760	21945	bcsABII-A	01453
14	OdNW_18-Q3-R42-56_BAT3C.134_sub	d__Bacteria;p__Myxococcota;c__Myxococcia;o__Myxococcales;f__Anaeromyxobacteraceae;g__Anaeromyxobacter;s__	tig00009616-10-535830	48524	bcsCII	03808
15	Skiv_18-Q3-R9-52_BATAC.176	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Gallionellaceae;g__UBA7399;s__	tig00025195-10-609250	40291	bcsABII-A	01730
16	Skiv_18-Q3-R9-52_BATAC.396	d__Bacteria;p__Acidobacteriota;c__Acidobacteriae;o__Bryobacterales;f__Bryobacteraceae;g__UBA690;s__	tig00002691-10-6419970	11117	bcsABII-A	00856
17	Skiv_18-Q3-R9-52_MAXAC.078_sub	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Rhodocyclaceae;g__Dechloromonas;s__	tig00014165-10-544790	30935	bcsABII-A	00765
18	Vibo_18-Q3-R45-57_BAT3C.98	d__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Rhodanobacteraceae;g__Dokdonella;s__	tig00177979-10-6033000	224892	bcsCII	02186