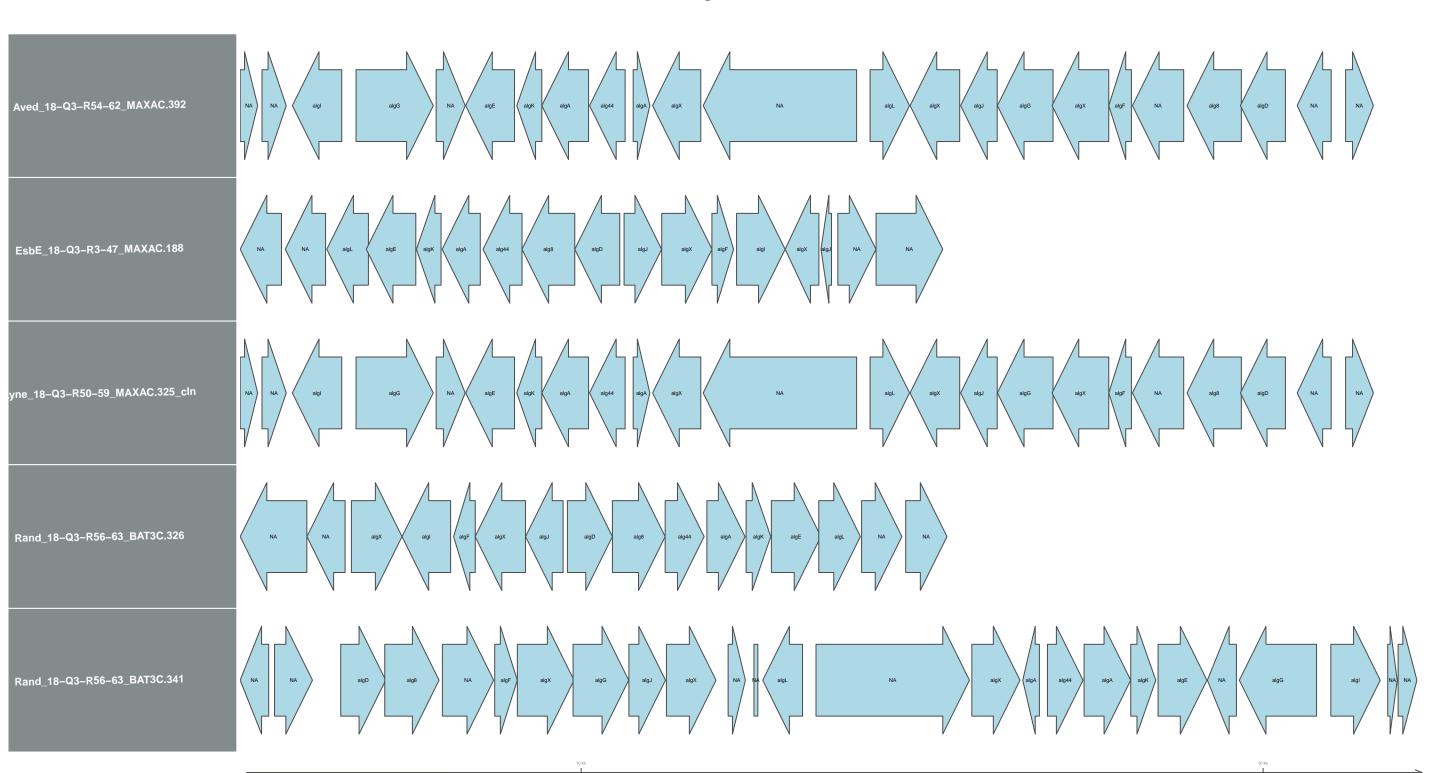


alginate



	ID	GTDBTax	seqname	start	Query_label	ProkkaNO
1	Aved_18-Q3-R54-62_MAXAC.392	d_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Micavibrionales;f_Micavibrionaceae;g_;s_	tig00000035-10-27699490	660128	alg44	00678
2	EsbE_18-Q3-R3-47_MAXAC.188	d_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Micavibrionales;f_Micavibrionaceae;g_UBA7657;s_UBA7657 sp002483465	tig00000486-10-13516940	61819	alg44	00067
3	Lyne_18-Q3-R50-59_MAXAC.325_cln	d_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Micavibrionales;f_Micavibrionaceae;g_;s_	tig00000079-10-27418810	2546594	alg44	02512
4	Rand_18-Q3-R56-63_BAT3C.326	d_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Micavibrionales;f_Micavibrionaceae;g_UBA7657;s_UBA7657 sp002483465	tig00000230-10-11682560	447802	alg44	00444
5	Rand_18-Q3-R56-63_BAT3C.341	$\label{lem:d_Bacteria} \begin{tabular}{ll} d_Bacteria:p_Proteobacteria:c_Alphaproteobacteria:o_Micavibrionales:f_Micavibrionaceae:g_:s_\\ \end{tabular}$	tig00811517-10-16736710	1237518	alg44	02408