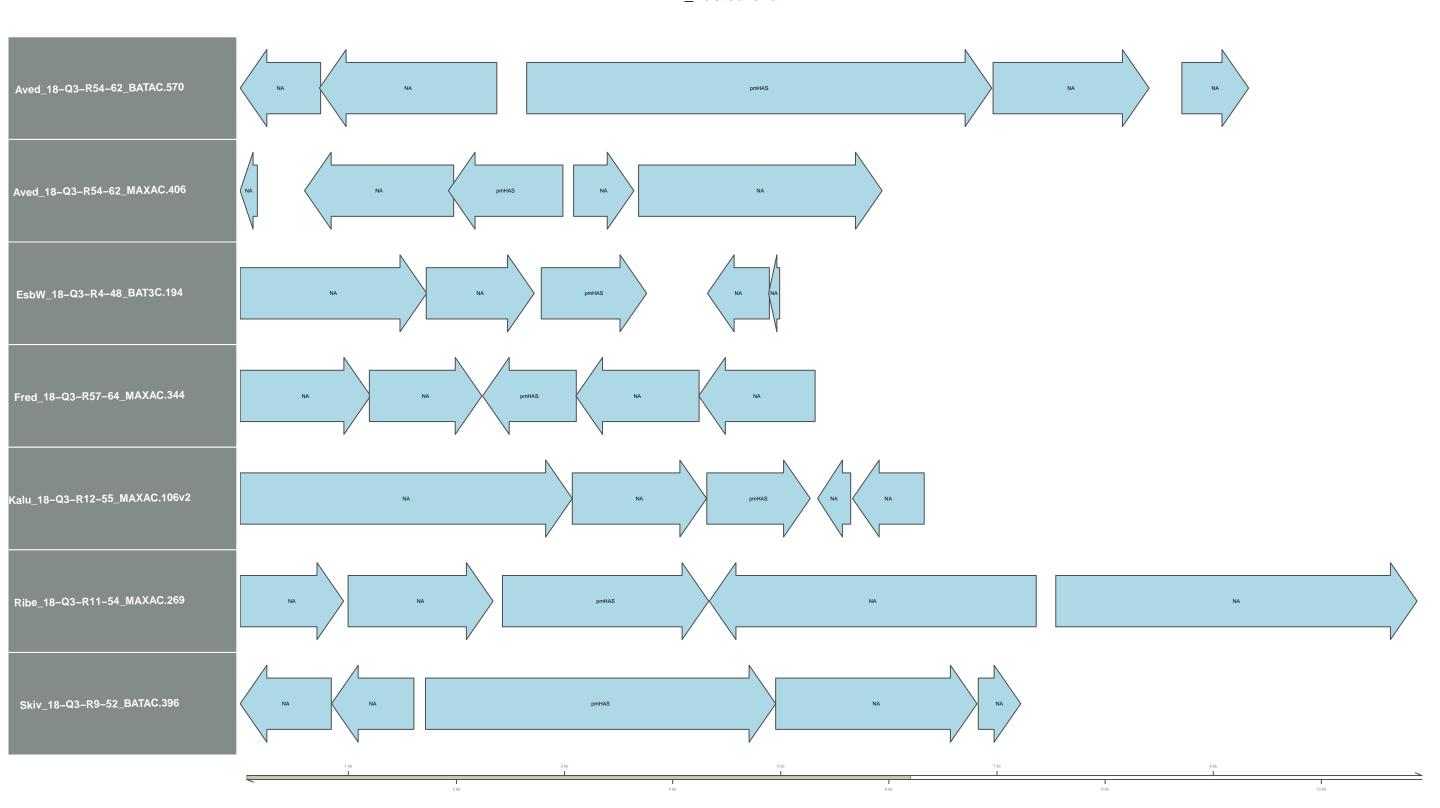


HA_Pasteurella



	ID	GTDBTax	seqname	start	Query_label	ProkkaNO
1	Aved_18-Q3-R54-62_BATAC.570	d_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Burkholderiales;f_Hydrogenophilaceae;g_UBA3361;s_	tig00020161-10-1851290	105983	pmHAS	03279
2	Aved_18-Q3-R54-62_MAXAC.406	d_Bacteria;p_Krumholzibacteriota;c_Krumholzibacteria;o_LZORAL124-64-63;f_LZORAL124-64-63;g_;s_	tig00007022-10-3217080	5661	pmHAS	01084
3	EsbW_18-Q3-R4-48_BAT3C.194	d_Bacteria;p_Acidobacteriota;c_Blastocatellia;o_Pyrinomonadales;f_;g_;s_	tig00003338-10-6811220	646617	pmHAS	00537
4	Fred_18-Q3-R57-64_MAXAC.344	d_Bacteria;p_Bacteroidota;c_Bacteroidia;o_Flavobacteriales;f_PHOS-HE28;g_PHOS-HE28;s_	tig00000591-10-23971360	1311899	pmHAS	01083
5	Kalu_18-Q3-R12-55_MAXAC.106v2	d_Bacteria;p_Chloroflexota;c_Anaerolineae;o_Promineofilales;f_Promineofilaceae;g_GCA-2699125;s_	tig00000173-10-21988890	2024772	pmHAS	01733
6	Ribe_18-Q3-R11-54_MAXAC.269	d_Bacteria;p_Planctornycetota;c_UBA8742;o_UBA2392;f_UBA2392;g_;s_	tig00000386-10-27093100	1588438	pmHAS	01313
7	Skiv_18-Q3-R9-52_BATAC.396	$\verb d_Bacteria:p_Acidobacteriota;c_Acidobacteriae;o_Bryobacterales;f_Bryobacteraceae:g_UBA690;s_DRA690;a_Bryobacteraceae:g_UBA690$	tig01148979-10-1755960	165172	pmHAS	06879