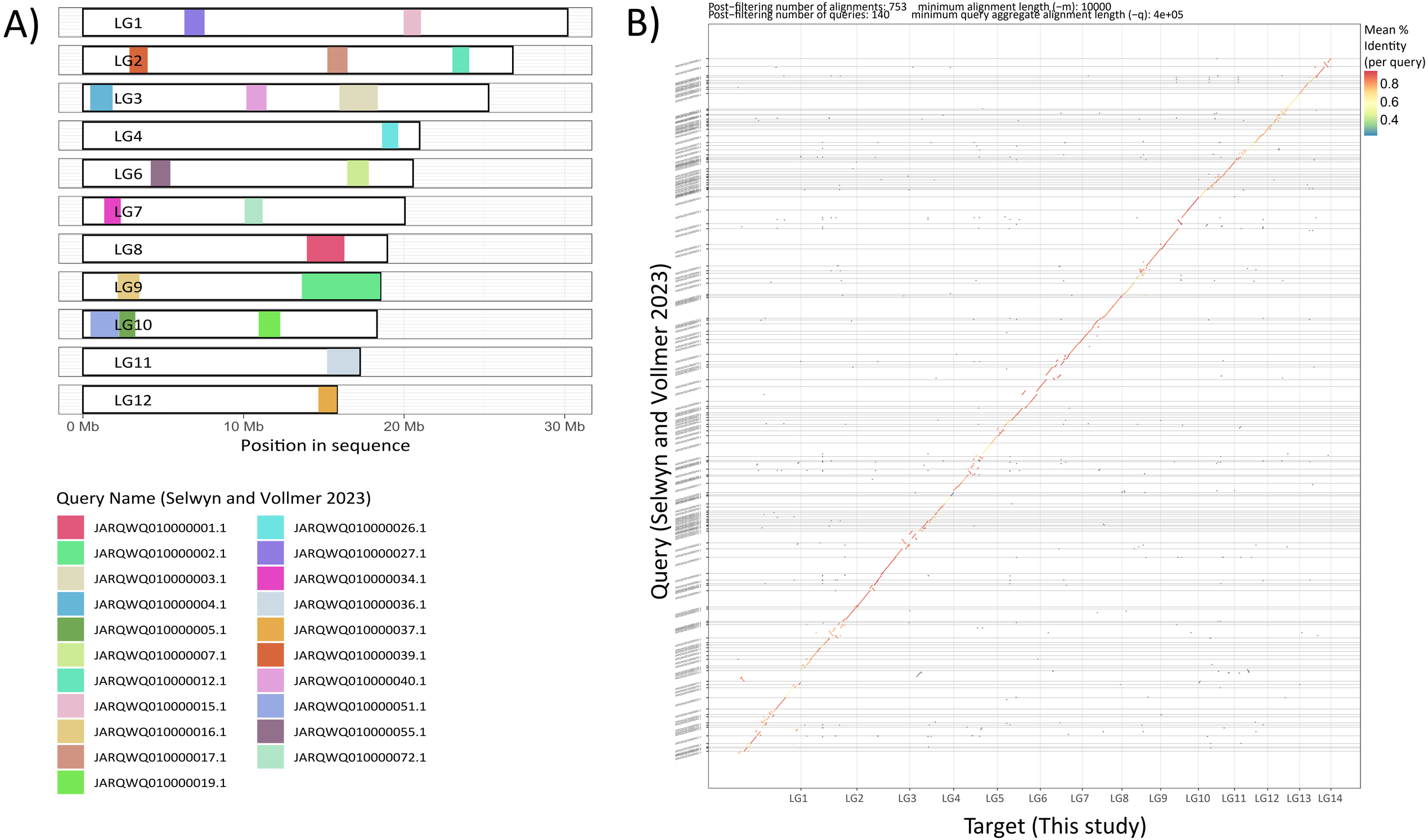
A)
LG1
LG2
LG3
LG4
LG6
LG7
LG8
LG9
LG10
LG11
LG12
0 Mb
10 Mb
20 Mb
30 Mb
Position in sequence
Query Name (Selwyn and Vollmer 2023)
JARQWQ010000001.1
JARQWQ010000002.1
JARQWQ010000003.1
JARQWQ010000004.1
JARQWQ010000005.1
JARQWQ010000007.1
JARQWQ010000012.1
JARQWQ010000015.1
JARQWQ010000016.1
JARQWQ010000017.1
JARQWQ010000019.1
JARQWQ010000026.1
JARQWQ010000027.1
JARQWQ010000034.1
JARQWQ010000036.1
JARQWQ010000037.1
JARQWQ010000039.1
JARQWQ010000040.1
JARQWQ010000051.1
JARQWQ010000055.1
JARQWQ010000072.1
B)
Post−filtering number of alignments: 753 minimum alignment length (−m): 10000
Post−filtering number of queries: 140 minimum query aggregate alignment length (−q): 4e+05
Mean % Identity (per query)
0.8
0.6
0.4
Query (Selwyn and Vollmer 2023)
LG1
LG2
LG3
LG4
LG5
LG6
LG7
LG8
LG9
LG10
LG11
LG12
LG13
LG14
Target (This study)