

Table S8: Location of predicted genomic islands in the genome of *Y. entomophaga* MH96 based on Island Viewer 4 analysis.

Island	Start	Stop
Is.1	4,245,874	43,205
Is.2	63,587	68,974
Is.3	190,789	198,981
Is.4	642,845	691,291
Is.5	749,459	763,698
Is.6	840,932	892,566
Is.7	911,590	931,192
Is.8	1,042,716	1,047,262
Is.9	1,161,921	1,166,559
Is.10	1,323,793	1,330,066
Is.11	1,5726,53	1,602,200
Is.12	1,693,457	1,718,980
Is.13	2,171,815	2,180,723
Is.14	2,633,449	2,651,765
Is.15	3,084,579	3,102,232
Is.16	3,204,015	3,214,229
Is.17	3,263,328	3,292,504
Is.18	3,326,808	3,340,556
Is.19	3,516,027	3,562,485
Is.20	3,655,950	3,662,717
Is.21	3,729,760	3,736,838