

Supplementary Table 1: Predicted genomic islands (GI).

Label	Begin	End	Length	Feature	nbr of predicted CDS
GI1	63156	97391	34235	tRNA border	25
GI2	360530	382020	21490	tRNA border+integrase	16
GI3	455772	471292	15520		17
GI4	516395	531476	15081		16
GI5	571179	587135	15956	tRNA border	14
GI6	696813	727942	31129		35
GI7	818344	840045	21701		19
GI8	851752	879285	27533		23
GI9	928330	957648	29318	tRNA border+integrase	31
GI10	1072153	1091983	19830		14
GI11	1105158	1112867	7709	tRNA border	5
GI12	1139167	1158439	19272		20
GI13	1189396	1228741	39345		42
GI14	1511151	1518724	7573		10
GI15	1676634	1698617	21983		15
GI16	1909860	1928847	18987		7
GI17	1950625	1959341	8716		7
GI18	2069502	2100324	30823		26
GI19	2126059	2176889	50831		36
GI20	2216273	2234260	17988		16
GI21	2237972	2247943	9972		8
GI22	2456093	2470601	14509		10
GI23	2582335	2643511	61177		43
GI24	2894754	2965888	71135	tRNA border	55
GI25	3127049	3154670	27622		33