

circular	linear	pAtC58	pTiC58	Regulatory motif
2	1	0	0	Adenylate and Guanylate cyclase catalytic domain
0	0	0	0	Adenylate cyclase, class-I
4	10	0	0	AsnC (Irp) family
0	0	0	1	Autoinducer synthetase
11	19	6	0	Bacterial regulatory helix-turn-helix proteins, araC family
20	34	16	5	Bacterial regulatory helix-turn-helix proteins, lysR family
10	3	1	0	Bacterial regulatory proteins, arsR family
7	6	0	0	Bacterial regulatory proteins, crp family
2	3	1	1	Bacterial regulatory proteins, deoR family
5	3	0	0	Bacterial regulatory proteins, Fis family
17	25	5	1	Bacterial regulatory proteins, gntR family
3	13	6	0	Bacterial regulatory proteins, lacl family
12	9	1	1	Bacterial regulatory proteins, luxR family
4	2	1	0	Bacterial regulatory proteins, merR family
13	12	7	0	Bacterial regulatory proteins, tetR family
1	5	2	0	Bacterial transcriptional regulator
1	0	0	0	CarD-like transcriptional regulator
3	2	1	0	Cold-shock DNA-binding domain
3	0	0	0	Ferric uptake regulator family
18	10	1	0	GGDEF domain
2	2	1	0	Helix-turn-helix domain, rpiR family
26	10	3	1	His Kinase A (phosphoacceptor) domain
31	13	5	1	Histidine kinase-, DNA gyrase B-, phytochrome-like ATPase
0	0	0	0	LuxS protein
15	8	4	0	MarR family
2	0	0	0	Nitrogen regulatory protein P-II
37	21	5	1	Response regulator receiver domain
2	8	0	0	ROK family
1	0	0	0	Sigma-54 factors family
4	5	0	0	Sigma-54 interaction domain
2	0	0	0	Sigma-70 factor
5	3	3	0	Sigma-70 factor (ECF subfamily)
3	2	1	0	SIS domain
11	6	4	1	Transcriptional regulatory protein, C terminal
2790	1882	550	198	Total genes in replicon
2.8	2.1	0.5	0.2	Genome size (Mb)
277	235	74	13	Total regulatory motifs
215	196	65	10	Distinct regulatory genes
7.71	10.41	11.82	5.05	% regulatory genes