

Table S1 Global statistics of *Bt* m401

Feature	Value
Total sequence length	6,005,135
Total ungapped length	6,004,585
% GC	34.72
Gaps between scaffolds	0
Number of scaffolds	45
Scaffold N50	391,966
Scaffold L50	4
Number of contigs	64
Contig N50	209,993
Contig L50	9
Total number of plasmids	4
Number of component sequences (WGS or clone)	45