

Table S1. General genome^a features of strain HdN1.

Size (bp)	4,587,455
G + C content (mol%)	53.26
Stable RNAs	
rRNAs	9 (3 operons)
tRNAs	47
Protein-coding sequences (CDS) ^b	3,763
Coding (%)	89.2
Average length (bp)	1,088
Pseudo genes	45

^a Accession no. FP929140

^b Without "pseudo"-qualifier