

Serotype-associated polymorphisms in a partial *rpoB* gene sequence of *Salmonella enterica*

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Salmonella enterica is a zoonotic bacterium with more than 2500 serotypes, which affect a wide range of hosts and produce diverse clinical outcomes. Strain identification usually involves costly and time-demanding procedures. This paper describes the sequencing of a *rpoB* hypervariable gene segment (847 bp) that allows identification of serotypes in *S. enterica* strains isolated from several hosts. The nucleotide similarity values among *S. enterica* serotypes ranged from 98.23% to 99.88%, with potential usefulness for devising a simple one-step sequencing as a first approach for identification of *S. enterica* strains. In conclusion, the analysis of polymorphisms in the partial *rpoB* sequence can discriminate *S. enterica* strains at the subspecies level.