**Figure S1. Nm *sodC* compared to Hi *sodC* and primer and probe location.** *sodC* was sequenced from 14 groupable and NG meningococci and a consensus nucleotide sequence was generated. An *H. influenzae* *sodC* consensus was built using two sequences from GenBank (accession numbers M84012 and AF549211). Nucleotide differences from the meningococcal sequence are white letters highlighted in black in the *Haemophilus* sequence. The primers and probe (highlighted in gray) were chosen at positions not only containing at least three nucleotide differences between Nm and Hi*,* but also where *sodC* nucleotide sequence was conserved among the meningococcal isolates that were sequenced. The meningococcal *sodC* open reading frame is 560 nucleotides long; the portion of the gene depicted corresponds to nucleotides 255-513.

Supplementary Data Figure S1

