

Table S5. Copy number of *pilE*, *pilS* and *opa* genes in *Neisseria* species. See supplemental references (Text S1) for additional information on *N. meningitidis* MC58 [3,20] and *N. gonorrhoeae* FA1090 [20,21].

Genome	<i>pilE</i>	<i>pilS</i>	<i>Opa</i>
<i>N. elongata</i>	1	5*	0
<i>N. sicca</i>	1	5	0
<i>N. mucosa</i>	1	5	0
<i>N. subflava</i>	2	2	0
<i>N. flavescens</i>	1	4	2
<i>N. cinerea</i>	2	3	0
<i>N. polysaccharea</i>	1	3	1
<i>N. lactamica</i> 23970	1	3	3
<i>N. meningitidis</i> MC58	1	9	4
<i>N. gonorrhoeae</i> FA1090	1	19	11

*The *pilS* copy upstream of *pilE* in *N. elongata* contains a complete open reading frame, but appears to be missing the promoter region.