

Table S1. Fit indices for model for Hosmer-Lemshow goodness fit statistics.

Model	χ^2	df	<i>p</i>-value
Double Adult	18.2439	13	0.1485
Hepatitis B	5.7218	13	0.9557
Hepatitis A	11.4485	13	0.5733
HPV	12.0177	13	0.5262
13-valent pneumococcal	6.6939	13	0.9172
23-valent pneumococcal	11.4255	13	0.5752
Meningococcal C	6.7396	13	0.9151
Measles/Mumps/Rubella	10.1338	13	0.6830
Yellow Fever	10.7265	13	0.6337
Adequate Scheme	5.7055	13	0.9562

χ^2 : ratio Chi square; df: degrees of freedom ratio