



Public Health
England



England (UK)

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General information - England



Population size	Category	54.8 million (= 84% of UK's population)
Structure of surveillance of meningococcal disease	Statutory notification of disease Enhanced surveillance Centralized Laboratory Surveillance	Yes Yes Yes
NRL: repertoire of typing targets	Serogroup / genogroup PorA antigen sequence typing FetA antigen sequence typing Vaccine Antigen sequence typing (e.g. fhbp) Antibiotic resistance gene typing MLST (7 loci) Genome sequencing	Yes culture & non-culture Yes culture & non-culture Yes culture only Yes culture & non-culture Yes culture only Yes culture only Yes culture only
Meningococcal vaccine recommendation	Polysaccharide vaccines (not conjugated) MenC conjugate MenACWY conjugate Protein vaccines targeting MenB	No YES UK imm schedule at 12M only YES teenage booster (13/14Y) & catch-up YES Imm schedule (2,4 and 12M) & risk groups
Uptake of MenWY vaccines	estimated % per target group size	ACWY 36% for 1st targeted group*

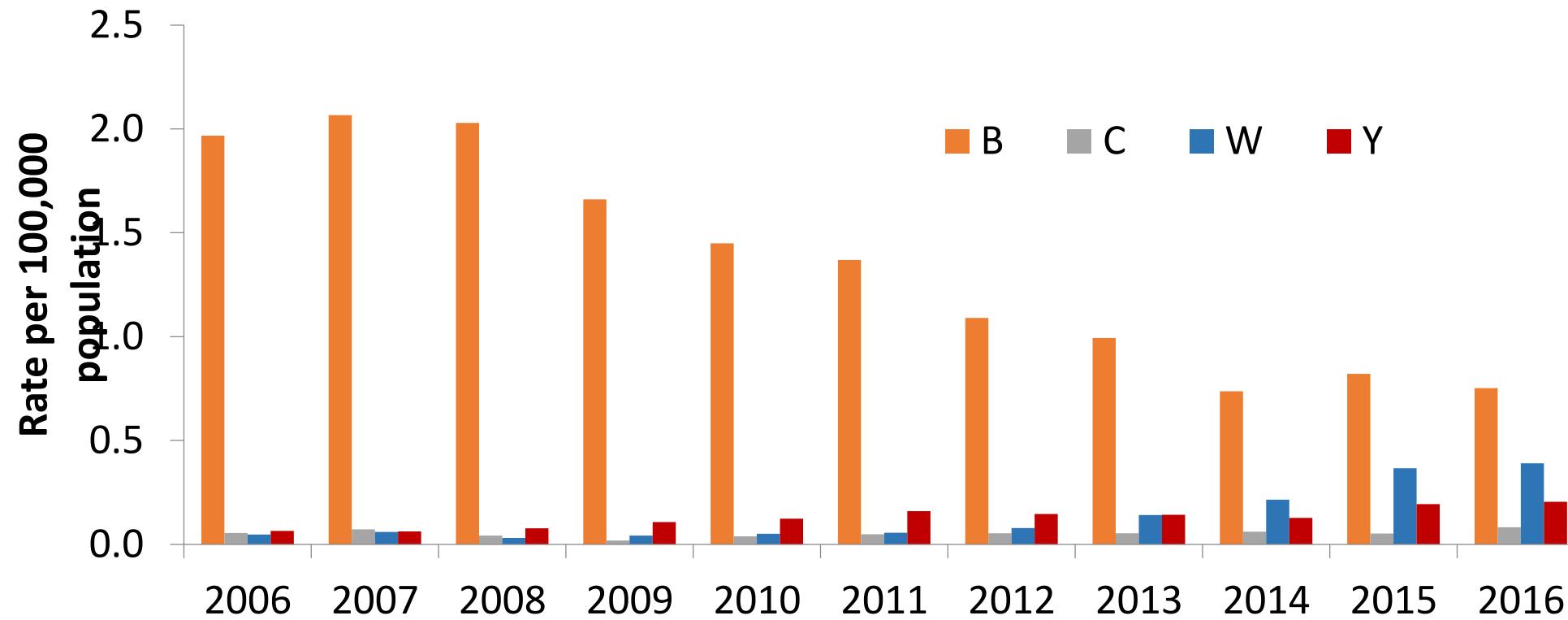


Basic numbers for 2015

No of reported cases of invasive meningococcal disease * (incidence rate)	N=796 (1.5 /100,000)
Proportion of reported cases of invasive meningococcal disease, whose strains or samples were processed at NRL (<i>=laboratory surveillance coverage of reported cases</i>)	100 %
MenW <ul style="list-style-type: none">• no of cases,• incidence rate,• % of all cases	N=201 0.4 /100,000 25 %
MenY <ul style="list-style-type: none">• no of cases,• incidence rate,• % of all cases	N=106 0.2 /100,000 13%

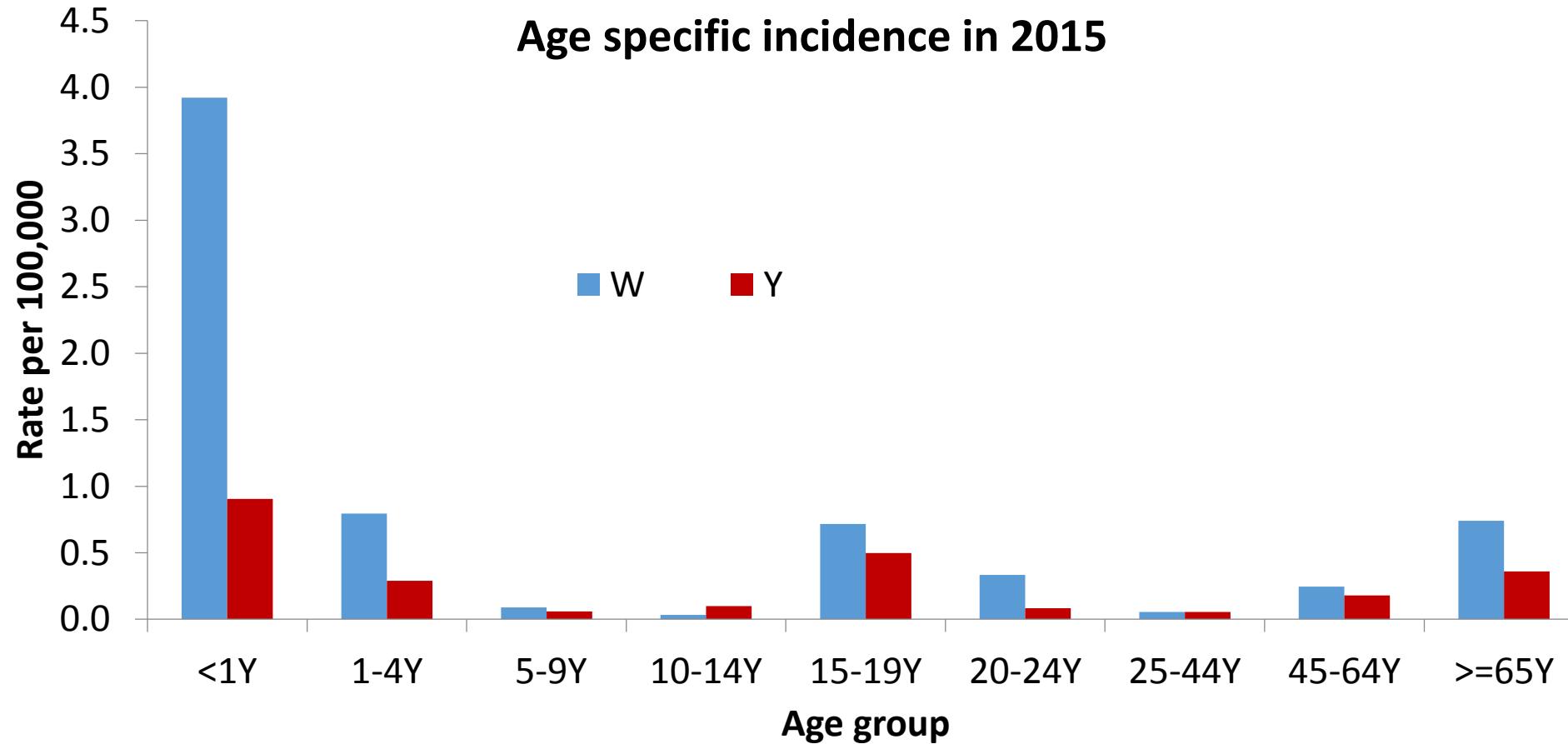
* please use [ECDC case definition](#)

England - Incidence rates (N/100,000/a) for serogroups B, C, W, Y from 2006-2016*

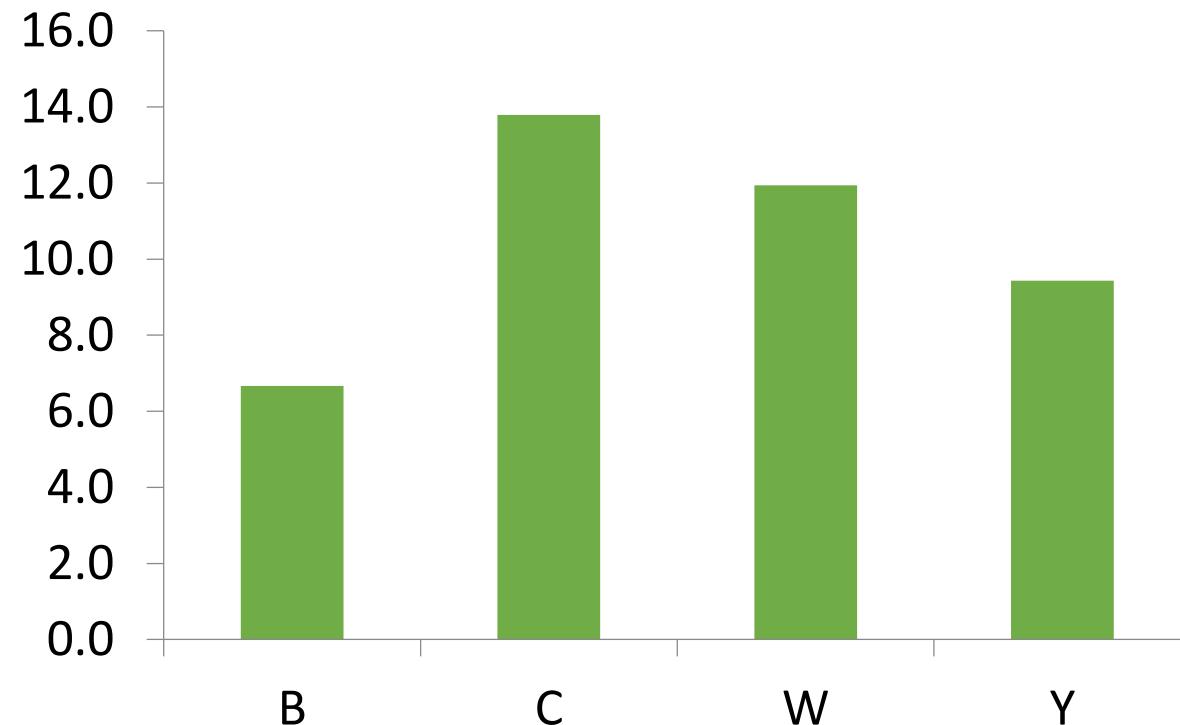


*calculated annual incidence based on 37 weeks of data in 2016

England -Age specific incidence rates 2015 and/or 2016 (N/100,000 of age group/a) for serogroups W and Y



Case fatality rates* 2015 (in %) for serogroups B, C, W, Y



*Based on those with IMD certified as an underlying cause of death in ONS records

Sequence typing of MenW in 2015 and / or 2016



Year	2015 & 2016
Dominant variant according to the typing method used	P1.5,2: F1-1: ST-11 (cc11)
Dominant variant according to the typing method used, in % of all MenW cases	80% (220/276; cultures only; 5 not determined)
cc11 in % of all serogroup W cases	94.3% (265/281 cultures only)

Outbreaks caused by serogroups W and Y 2012 – 2016, England



	MenW	MenY
Number of outbreaks or clusters	Probable 5 (checked by WGS), <i>Possible 12¹</i>	2
Total number of patients involved	Probable Total = 10 Median 2, Range 2-7.	Total = 6, Range 2-4
Settings	Probable² Nursing home (2), Ambulance (1), Local Geog. (1), Wider Geog. (1)	Social group (1), Geog. (1)

¹PossibleIncidents additionally includes family (2) , nursing home (1) schools (1), university (2) local geography (1); where cases range <1 to 186 days, Patient ages 26d to 95 yrs.

²Probable cases <1 to 8 days apart .