Annual Report of the Australian Meningococcal Surveillance Programme, 2014

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Abstract

In 2014 there were 165 laboratory-confirmed cases of invasive meningococcal disease analysed by the Australian National Neisseria Network. This number was higher than the number reported in 2013, but was the second-lowest reported since inception of the Australian Meningococcal Surveillance Programme in 1994. Probable and laboratory confirmed invasive meningococcal disease (IMD) is notifiable in Australia, and there were 170 IMD cases notified to the National Notifiable Diseases Surveillance System in 2014. This was also higher than in 2013, but was the second-lowest number reported IMD cases recorded by this Program. The meningococcal serogroup was determined for 161/165 (98%) of laboratory confirmed IMD cases. Of these: 80.1% (129 cases) were serogroup B infections; 1.9% (3 cases) were serogroup C infections; 9.9% (16 cases) were serogroup W135; and 8.1% (13 cases) were serogroup Y. Primary and secondary disease peaks were observed in those aged 4 years or less, and in adolescents (15-19 years) respectively. Serogroup B cases predominated in all jurisdictions and age groups, except for those aged 65 years or over, where serogroups Y and W135 combined predominated. The overall proportion and number of IMD caused by serogroup B was higher than in 2013, but has decreased from previous years. The number of cases of IMD caused by serogroup C is the lowest reported to date. The number of IMD cases caused by serogroup Y was similar to previous years, but the number of IMD cases caused serogroup W135 was higher than in 2013. The proportion of IMD cases caused by serogroups Y and W135 has increased in recent years, whilst the overall number of cases of IMD has decreased. Molecular typing was able to be performed on 106 of the 165 IMD cases. In 2014, the most common *porA* genotypes circulating in Australia were P1.7-2,4 and P1.22,14. All IMD isolates tested were susceptible to ceftriaxone and

ciprofloxacin. There were two isolates that were resistant to rifampicin. Decreased susceptibility to penicillin was observed in 88% of isolates.

Keywords: antibiotic resistance; disease surveillance; meningococcal disease; Neisseria meningitidis Introduction

The Australian National Neisseria Network (NNN) is a collaborative network of reference laboratories in each state and territory that contribute to the laboratory surveillance system of the pathogenic *Neisseria* species (*N. meningitidis* and *N. gonorrhoeae*). Since 1994 the NNN has coordinated laboratory data from the examination of *N. meningitidis* cases of invasive meningococcal disease (IMD) for the Australian Meningococcal Surveillance Programm2e (AMSP). (1) The AMSP is funded by the Australian Government Department of Health. Participating NNN laboratories supply phenotypic and genotypic data on invasive meningococci for the AMSP. These data supplement the notification data from the National Notifiable Diseases Surveillance System (NNDSS), which includes cases of probable IMD as well as laboratory confirmed IMD. The characteristics of meningococci responsible for IMD and the associated demographic information are important considerations for management of individual patients and their contacts. These data also inform public health responses for outbreaks or case clusters, locally and nationally. The introduction of the publicly funded conjugate serogroup C meningococcal vaccine onto the National Immunisation Program in 2003 has seen a significant and sustained reduction in the number of cases of serogroup C IMD after 2003.(2) However, IMD remains an issue of public health concern in Australia.

Methods

Case confirmation of invasive meningococcal disease

Case confirmation is based on isolation of *N. meningitidis*, or a positive nucleic acid amplification testing (NAAT) from a normally sterile site, defined as laboratory definitive evidence of IMD by the Communicable Diseases Network Australia criteria.(3) Information regarding the site of infection, age and sex of the patients is collated by the NNN for the AMSP.

IMD cases are categorised on the basis of the site from which *N. meningitidis* was isolated, or from which meningococcal DNA was detected. When *N. meningitidis* is grown from blood only, the IMD case is classified as septicaemia; cerebrospinal fluid (CSF) only cultures are classified as meningitis.

When *N. meningitidis* is grown from both blood and cerebrospinal fluid (CSF) cultures from the same patient, the case is classified as one of meningitis.

Phenotyping and genotyping of Neisseria meningitidis

Phenotyping is limited to the determination of the serogroup by detection of soluble polysaccharide antigens. Genotyping of both isolates and DNA extracts is performed by sequencing of products derived from amplification of the porin genes *porA*, *porB* and *FetA*.

Antibiotic susceptibility testing

Isolates were tested to determine their minimum inhibitory concentration (MIC) values to antibiotics used for therapeutic and prophylactic purposes: ceftriaxone, ciprofloxacin; rifampicin. This program uses the following parameters to define the various levels of penicillin susceptibility or resistance when determined by a standardised agar plate dilution technique: (4) These are: Sensitive (MIC \leq 0.03 mg/L); Less Sensitive (MIC 0.06–0.5 mg/L) and Resistant (MIC \geq 1 mg/L).

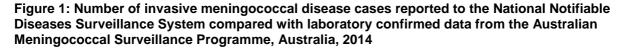
Results

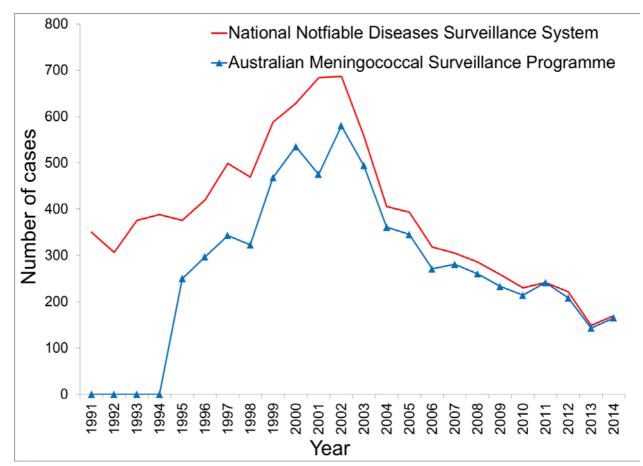
In 2014, there were 165 laboratory-confirmed cases of IMD analysed by the NNN, and 170 cases notified to the NNDSS. Thus, laboratory data were available for 97% of notified cases of IMD in Australia in 2014 (Figure 1). This is the second-lowest annual number of IMD cases recorded by the NNDSS, and by the AMSP (In 2013, there were 149 IMD cases recorded by NNDSS, 143 laboratory confirmed IMD cases reported by the AMSP). As in previous years, the peak incidence for IMD was in late winter and early spring (1 July to 30 September) (Table 1).

Table 1: Laboratory confirmed cases of invasive meningococcal disease, Australia, 2014, by quarter

Serogroup	1 January– 31 March	1 April– 30 June	1 July– 30 September	1 October– 31 December	2014 Total
В	20	29	45	35	129
С	2	0	1	0	3
Y	0	7	3	3	13
W135	4	3	4	5	16
NG	0	0	0	0	0
ND	0	2	0	2	4
Total	26	41	53	45	165

(NG: non groupable; ND: not determined)





In 2014, the highest number of laboratory confirmed cases was from Queensland (39 cases), which was higher than that reported in this state in 2013 (32 cases). Other states that recorded a rise in IMD cases in 2014 compared with 2013 were: Victoria (33 cases in 2014, compared with 23 in 2013), and South Australia (31 cases in 2014, compared with 21 in 2013). By contrast, New South Wales recorded a fall in the number of IMD cases in 2014 (36 cases) compared with 2013 (43 cases). Numbers for the other states were similar to 2013 (Table 2).

			Sero	group			
State or territory	В	С	Y	W135	NG	ND	Total
ACT	1	0	1	0	0	0	2
NSW	21	0	8	6	0	1	36
NT	4	0	0	0	0	0	4
Qld	31	1	2	3	0	2	39
SA	31	0	0	0	0	0	31
Tas.	1	0	0	1	0	0	2
Vic.	27	0	1	4	0	1	33
WA	13	2	1	2	0	0	18
Australia	129	3	13	16	0	4	165
	78.2	1.8	7.9	9.7	0	2.4	%

 Table 2: Number of laboratory confirmed cases of invasive meningococcal disease, Australia,

 2014, by state or territory and serogroup

(NG: non groupable; ND: not determined).

Age distribution

Nationally, the peak number of IMD cases was in children less than 5 years of age, similar to previous years. Between 2007 and 2013, 28% to 36% of cases were in this age group. In 2014, 46/165 (28%) IMD cases occurred in this age group (Table 3). A secondary disease peak has also been observed in previous years amongst adolescents aged 15 to 19 years. Of the total cases of IMD, 30/165 (18%) were in those aged 15 to 19 years in 2014, which was the same as the proportion reported for 2013; and similar to the proportion reported in this age group during the period 2007 to 2011 (17% to 20%). The proportion of IMD cases in those aged 25 to 44 (14.5%, 24 cases) was almost double than that in 2013 (7.7%, 11 cases).

		Age group									
Serogroup	<1	1–4	5–9	10–14	15–19	20–24	25–44	45–64	65+	NS	Total
В	24	16	8	5	22	20	13	12	7	2	129
С	1	1	0	0	0	1	0	0	0	0	3
Y	1	0	0	0	2	1	1	1	7	0	13
W135	2	0	0	1	5	1	0	2	5	0	16
NG	0	0	0	0	0	0	0	0	0	0	0
ND	1	0	0	0	1	1	1	0	0	0	4
Total	29	17	8	6	30	24	15	15	19	2	165
% B of within age group	82.8	94.1	100.0	83.3	73.3	83.3	86.7	80.0	36.8		

Table 3: Laboratory confirmed cases of invasive meningococcal disease, Australia, 2014, by age and serogroup

(NS: age not stated; NG: non groupable: ND not determined)

Anatomical site of samples and method of confirmation

In 2014, diagnosis was made by a positive culture in 95/165 (58%) cases and, 70/165 (42%) cases were confirmed by NAAT testing (Table 4).

There were 58 diagnoses of meningitis based on cultures or NAAT examination of CSF either alone or with a positive blood sample. There were 103 diagnoses of septicaemia based on cultures or NAAT examination from blood samples alone (Table 4). There were 4 IMD diagnoses by positive joint fluid culture (n=2) and NAAT (n=2).

Table 4: Number of laboratory confirmed cases of invasive meningococcal disease, Australia,
2014, by anatomical source and method of confirmation

Specimen type	Bacterial culture	NAAT	Total
Blood	69	34	103
CSF +/- blood	24	34	58
Other [‡]	2	2	4
Total	95	70	165

NAAT: nucleic acid amplification testing; CSF = cerebrospinal fluid.

Serogroup data

Number of cases of invasive meningococcal disease by serogroup B, C, Y, W135

The serogroup was determined for 161 of 165 laboratory confirmed cases of IMD in 2014 (Tables 2 and 3). There has been an overall decrease in the number of cases of IMD in Australia in recent years, which was initially predominantly due to a reduction in the number of cases of IMD caused by serogroup C from 2003 to 2007. This was followed by a decline in the numbers IMD cases caused by serogroup B from 194 cases in 2009, to 104 cases in 2013. In 2014, there was an increase in the numbers of IMD caused by serogroup Y has remained stable since 2011, whereas, the number of cases of serogroup W135 IMD has increased in recent years (7 to 16 cases in 2011 to 2014, compared with 4 to 9 cases in 2007 to 2010). In 2014 there were 16 cases, the highest number ever reported by the AMSP.

Proportions of serogroup B, C, Y, W135 invasive meningococcal disease

Of the 161 IMD strains for which the serogroup was determined, 80.1% were serogroup B, which was higher than in 2013 (74.8%), but lower than that reported in the years 2006 to 2012 (84% to 88%). The proportion of cases of IMD caused by serogroup B in children less than 5 years in 2014 was lower than in previous years (2008 to 2013) (Table 3, Figure 2). However, in young adults 20–24 years, the number of cases of serogroup B IMD in 2014 was higher than in 2007 to 2011 and 2013 (61% to 67%), and similar to 2012. The proportion of cases of IMD caused by serogroup B in those aged 15–19 has remained relatively stable since 2008, but was lower in 2014. Serogroup B IMD was prominent in IMD in all age groups excepting 65 years or more where, serogroup Y was equally prevalent, and serogroup W135 slightly less so,.

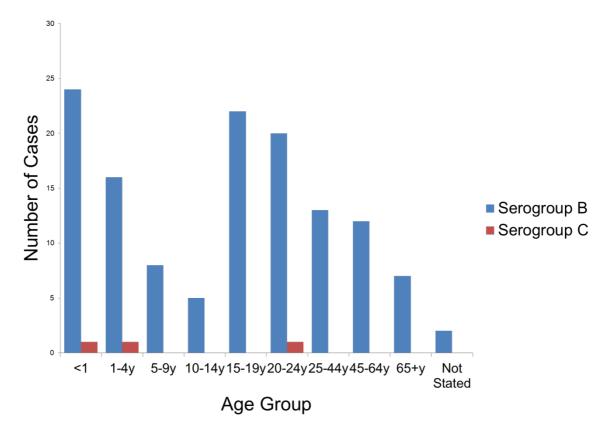


Figure 2: Number of serogroup B and C cases of confirmed invasive meningococcal disease, Australia, 2014, by age group

The number and proportion of IMD caused by serogroup C in 2014 was lowest since the inception of the Australian Meningococcal Surveillance Programme (1.9% and the number was?). Two of the three cases of IMD caused by serogroup C in 2014 were in those aged less than 20 years in 2014, compared with 1 case in 2013, 2 cases in 2012 and no cases in 2011 in this age group.

Of note, coincident with the decline in serogroup C IMD, the proportion of IMD caused by serogroups Y and W135 has been increasing in recent years. In 2012 to 2014 serogroup Y accounted for 7.7% to 10.8% of IMD, higher than the proportion reported in the period 2007 to 2011: 3.5% to 5.0%. Similarly the proportion by serogroup W135 IMD was 8.6% to 9.9% of IMD in 2013 to 2014, higher than the 1.8% to 4.5% reported in the period 2007 to 2011. The number and proportion of IMD cases caused by serogroup Y was highest in people aged 65 years or over in 2014. The number and proportion of IMD cases caused by serogroup W135 was highest in people aged 65 years or over, and also in people aged 15–19 years.

Genotyping

In 2014, genotyping was performed for 106/165 (64%) IMD cases (Tables 5 and 6). The predominant *porA* genotypes for serogroup B IMD cases were again P1.7-2,4 (14 cases) and P1.22,14 (14 cases). Other *porA* genotypes for serogroup B IMD cases more frequently seen in 2014 were P1.7,16-26 (7 cases); and P1.18-1,34 and P1.22,9 (6 cases each) The AMSP was not aware of any epidemiological link between any of the cases reported where genotyping was available.

2014 AMSP		Num	oer per sero	group	
Genotype PorA	Total	В	С	W135	Y
P1.5,2	9	0	0	9	0
P1.5-1,10-1	8	1	0	1	6
P1.5-1,10-4	5	1	0	1	3
P1.5-1,10-46	2	1	0	0	1
P1.5-1,10-8	2	0	2	0	0
P1.5-1,2-2	1	0	0	0	1
P1.5-1,2-52	1	1	0	0	0
P1.5-2,10-1	1	0	0	0	1
P1.5-2,2	1	0	1	0	0
P1.7,16-26	7	7	0	0	0
P1.7,16-44	2	2	0	0	0
P1.7,30	1	1	0	0	0
P1.7,30-3	1	1	0	0	0
P1.7-1,10-27	1	1	0	0	0
P1.7-2,4	14	14	0	0	0
P1.7-2,16-26	2	2	0	0	0
P1.12-6,13-13	1	1	0	0	0
P1.17,16-3	1	1	0	0	0
P1.18-1,3	6	3	0	3	0
P1.18-1,3-8	3	3	0	0	0
P1.18-1,30	1	1	0	0	0
P1.18-1,34	6	6	0	0	0
P1.19,15	1	1	0	0	0
P1.19-1,15-11	1	1	0	0	0
P1.19-1,26	1	1	0	0	0
P1.19-2,13-1	1	1	0	0	0

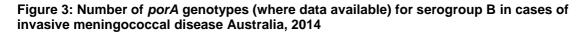
Table 5: Laboratory confirmed cases of invasive meningococcal disease, Australia, 2014, by *porA* gentoype

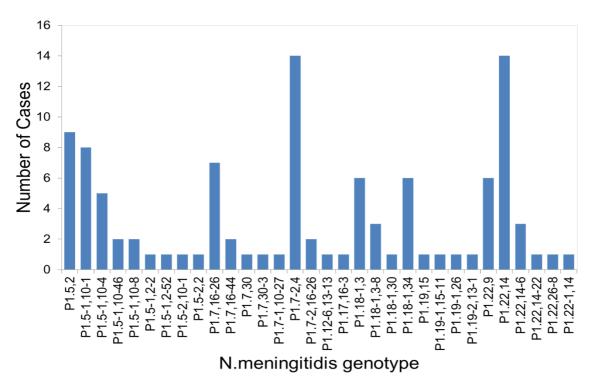
2014 AMSP		Numb	er per sero	group	
Genotype PorA	Total	В	С	W135	Y
P1.22,9	6	6	0	0	0
P1.22,14	14	14	0	0	0
P1.22,14-6	3	3	0	0	0
P1.22,14-22	1	1	0	0	0
P1.22,26-8	1	1	0	0	0
P1.22-1,14	1	1	0	0	0
Total	106	77	3	14	12

 Table 6: Distribution of porA genotype laboratory confirmed cases of invasive meningococcal disease, Australia, 2014, by state or territory

2014 AMSP	Number per serogroup per state									
Genotype PorA	NSW	Qld	Vic.	SA	WA	ACT	Tas.	NT		
P1.5,2	2 W135	2 W135	2 W135		2 W135		2 W135			
P1.5-1,10-1	6 Y,1 W135		1 B							
P1.5-1,10-4	1 W135	1 Y	1 Y		1 Y			1 B		
P1.5-1,10-46			1 B			1Y				
P1.5-1,10-8					2 C					
P1.5-1,2-2		1 Y								
P1.5-1,2-52	1 B									
P1.5-2,10-1	1 Y									
P1.5-2,2		1 C								
P1.7,16-26	3 B	3 B	1 B							
P1.7,16-44	1 B	1 B								
P1.7,30					1 B					
P1.7,30-3				1 B						
P1.7-1,10-27		1 B								
P1.7-2,4	1 B	7 B	4 B		1 B			1 B		
P1.7-2,16-26		1 B	1 B							
P1.12-6,13-13					1 B					
P1.17,16-3	1 B									
P1.18-1,3	1 W135		3 B, 2 W135							
P1.18-1,3-8		3 B								
P1.18-1,30		1 B								
P1.18-1,34		4 B	1 B		1 B					
P1.19,15			1 B							
P1.19-1,15-11	1 B									

2014 AMSP		Number per serogroup per state						
Genotype PorA	NSW	Qld	Vic.	SA	WA	ACT	Tas.	NT
P1.19-1,26		1 B						
P1.19-2,13-1					1 B			
P1.22,9	2 B		4 B					
P1.22,14	2 B	5 B	3 B		4 B			
P1.22,14-6	2 B	1 B						
P1.22,14-22	1 B							
P1.22,26-8				1 B				
P1.22-1,14		1 B						





Antibiotic susceptibility testing

Testing for antimicrobial susceptibility was performed for 95/165 (58%) of the IMD cases in 2014. All isolates tested were susceptible to ceftriaxone and ciprofloxacin. There were two isolates that were resistant to rifampicin. Using the defined criteria, 11/95 (11.6%) isolates were fully sensitive to penicillin (MIC 0.03 mg/L or less), and 84 (88%) isolates were less sensitive to penicillin (MIC=0.06–0.5 mg/L). No isolates were resistant to penicillin. The proportion of strains less sensitive to penicillin was the highest recorded by the AMSP.

Discussion

In 2014, there were 165 cases of laboratory confirmed IMD, representing 97% of the number of notifications to the NNDSS.(2) This is both the second lowest number of cases reported since laboratory based surveillance for confirmed IMD cases (AMSP) began in 1994, and since notification data collection commenced in 1991. This represents less than one-third of the number reported in Australia in 2002 (n=580), when IMD rates peaked in Australia. The introduction of the serogroup C vaccine to the national immunisation schedule in 2003 has led to a steady decline in the total number of both serogroup C, and the overall number of cases of IMD. The primary peak in IMD infection continues to be in children less than 5 years, as reported in previous years, with a secondary peak in adolescents.

The majority of IMD cases in Australia are caused by serogroup B. The proportion and number of IMD cases caused by serogroup C was lowest reported by the AMSP since the beginning of the program. The number of IMD cases caused by serogroup Y was similar to previous years. The number and proportion of cases caused by serogroup W135 was the highest reported by the AMSP. The proportion of IMD cases caused by serogroups Y and W135 has increased in recent years, coincident with the overall reduction in numbers of IMD cases, and are the predominant serogroups causing IMD in those aged 65 years or older.

As in previous years, genotypic data found no evidence of a substantial number of cases of IMD caused by *N. meningitidis* that have undergone genetic recombination. There have been concerns that the emergence of new and invasive subtypes following extensive vaccine use would occur given the capacity for genetic recombination within meningococci.(5) Therefore the monitoring of meningococcal genotypes is an important part of the NNN program.

All isolates were susceptible to ceftriaxone and ciprofloxacin; whilst there were two IMD isolates that were resistant to rifampicin. The proportion of IMD isolates with penicillin MIC values in the less sensitive category in 2014 was 88%, and was the highest proportion recorded by the AMSP. In previous years the range was 62% to 75% in 1996 to 2006; 67% to 79% in 2007 to 2009; and 78% to 85% in 2010 to 2013. Thus indicating a right shift in penicillin MIC values of IMD isolates, however, in Australia, the incidence of penicillin resistance in *N. meningitidis* is very low.

In early 2014, a recombinant multi-component meningococcal B vaccine became available in Australia.(6) This vaccine is not on the immunisation register but is available for purchase privately. Therefore uptake is elective and the impact of its introduction is yet to be determined in this country. The AMSP continues to monitor the phenotypic and genotypic features of N. meningitidis causing IMD to inform treatment protocols and monitor prevention strategies.

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