National Communicable Diseases Surveillance Report

Fortnight 14, 2021 Summary Notes for Selected Diseases

05 July to 18 July 2021

Infectious and congenital syphilis

Increases in infectious syphilis notifications are attributed to an on-going outbreak occurring in Aboriginal and Torres Strait Islander people residing in northern and central Australia, continued increases among men who have sex with men (MSM) in urban areas, and increases in women (Aboriginal and Torres Strait Islander and non-Indigenous) residing in urban areas of Australia.

Outbreak in northern and central Australia

In January 2011, an increase of infectious syphilis notifications among Aboriginal and Torres Strait Islander people was identified in the North West region of Queensland, following a steady decline at a national level in remote communities. Subsequent increases in infectious syphilis notifications were reported in the Northern Territory in 2013, Western Australia in 2014 and South Australia in 2016, following sustained periods of low notification rates. The outbreak is of significant public health concern given the: elevated rates of infectious syphilis among women of child-bearing age, increasing the risk of congenital syphilis; and the concomitant risk of HIV transmission. For the latest information on the infectious syphilis outbreak and related national activities, refer to the Department's website.

Increases among MSM

Since 2010 increases in notifications of infectious syphilis have been reported in MSM, predominately 20-39 years of age, residing in urban areas of Australia.

Increases among women (Aboriginal and Torres Strait Islander and non-Indigenous) Since 2016, increases in notifications of infectious syphilis have been reported in women (Aboriginal and Torres Strait Islander and non-Indigenous) aged predominately 20-39 years of age residing in urban areas in Australia. As noted in the outbreak in northern and central Australia, increases in women of childbearing age is of significant public health concern given the increased risk of congenital syphilis.

Syphilis response

On 23 March 2021, the Australian Health Protection Principal Committee (AHPPC) endorsed the *National strategic approach for responding to rising rates of syphilis in Australia 2021* (Strategic Approach) prepared through the Communicable Diseases Network Australia (CDNA) and BBV STI Standing Committee (BBVSS). The Strategic Approach builds on and intersects with existing national activities related to syphilis and provides specific focus for efforts towards rising rates of syphilis and adverse outcomes in Australia.

The CDNA and BBVSS are, in collaboration, developing priority public health actions, including those related to workforce and community engagement, to ensure progress is made towards reducing the incidence of syphilis and elimination of congenital syphilis in Australia. These actions will be provided to AHPPC for endorsement in the coming months. For further information on national activities related to STIs, including syphilis, refer to the Department's website.

Interpretative Notes

Selected diseases are chosen each fortnight based on either exceeding two standard deviations from the 90 day and/or 365 day five year rolling mean or other disease issues of significance identified during the reporting period. All diseases reported are analysed by notification receive date. Data are extracted each Monday of a CDNA week.

Totals comprise data from all States and Territories. Cumulative figures are subject to retrospective revision so there may be discrepancies between the number of new notifications and the increment in the cumulative figure from the previous period.

¹The past quarter (90 day) surveillance period includes the date range (20/04/2021 to 18/07/2021).

²The quarterly (90 day) five year rolling mean is the average of 5 intervals of 90 days up to 18/07/2021. The ratio is the notification activity in the past quarter (90 days) compared with the five year rolling mean for the same period.

³The past year (365 day) surveillance period includes the date range (19/07/2020 to 18/07/2021).

⁴The yearly (365 day) five year rolling mean is the average of 5 intervals of 365 days up to 18/07/2021. The ratio is the notification activity in the past year (365 days) compared with the five year rolling mean for the same period.

The five year rolling mean and the ratio of notifications compared with the five year rolling mean should be interpreted with caution. Changes in surveillance practice, diagnostic techniques and reporting may contribute to increases or decreases in the total notifications received over a five year period. Ratios are to be taken as a crude measure of current disease activity and may reflect changes in reporting rather than changes in disease activity.

ADT EN14/2021											Notification received date												
ADT FN14/2021				State or Territory								Totals for Australia				Historical 90 Day Period				Historical Yearly Period			
Disease group	Disease name	sease code	ACT				SA				This reporting	Previous	Same	Current		Ougstosly		Exceeds		Yearly		Exceeds	
				NSW	۲	Old		Tas	,u		This reporting period	reporting	reporting period last	Current year YTD	Past Quarter	Quarterly rolling	Ratio past	quarterly	Past Year	rolling 5 year	Ratio past	yearly	
									>			Period	year			5 year	quarter/5 year mean*	rolling mean +2 SD		mean	year/5 year mean*	rolling mean +2 SD	
		ق									05/07/2021 18/07/2021	21/06/2021 04/07/2021	05/07/2020 18/07/2020	01/01/2021 18/07/2021	20/04/2021 18/07/2021	mean		by	19/07/2020	19/07/2015		by	
Bloodborne diseases	Hepatitis B (newly acquired)	039	-	-	-	2	-	-	3	-	5	4	7	55	30	38.6	0.8	-	18/07/2021 100	18/07/2020 150.8	0.7	-	
	Hepatitis B (unspecified)	052	3		-	27	1	2	46	15	132	147	194	2,560	1,137	1,453.8	0.8	-	4,768	5,866.0	0.8	-	
	Hepatitis C (newly acquired) Hepatitis C (unspecified)	040 053	- 7	79	- 4	- 78	- 1	12	- 52	- 24	257	300		300 4,064	83 1,911	162.4 2,374.2	0.5	-	587 7,412	714.4 9,636.0	0.8	-	
	Hepatitis D	050	-	2	-	-	-	-	-	-	2	3		44	20	16.2	1.2	-	85	67.4	1.3	-	
Gastrointestinal diseases	Botulism	045	-	-	-	-	-	-	1	-	1	-	-	2	1	0.2	5.0	-	3	1.0	3.0	-	
	Campylobacteriosis Cryptosporidiosis	005 061	- 24	366 10	9	392 40	109	25 1	289 9	116 1	1,330 69	1,327 62	999 36	20,195 1,109	8,384 517	6,384.0 795.0	1.3 0.7	247.9	35,659 1,624	30,016.6 3,852.6	1.2 0.4	-	
	Haemolytic uraemic syndrome (HUS)	055	-	-	-	-	-	-	-	-	-	-	-	4	-	3.2	-	-	11	16.0	0.7	-	
	Hepatitis A Hepatitis E	038 051	-	1	-	1	-	-	-	-	1	1		11 8	7 8	43.8 9.2	0.2	-	18 11	237.2 48.2	0.1	-	
	Listeriosis	018	-	3	-	-	-	-	-	-	3	2		24	9	11.8	0.8	-	49	66.8	0.7	-	
	Paratyphoid	080	-	-	-	-	-	-	1	-	1		-	1	1	12.0	0.1	-	1	84.0	0.0	-	
	STEC Salmonellosis	054 030	- 4	83	18	- 78	9 18	- 4	31	19	21 255	7 298	10 252	332 6,960	123 2,169	101.4 3,225.2	1.2 0.7	-	555 10,467	492.4 15,571.0	1.1 0.7	-	
	Shigellosis	031	- 1	1	6	4	2	-	-	1	14	9	33	256	95	443.6	0.2	-	600	2,110.6	0.3	-	
	Typhoid Fever	035	-	-	-	-	-	-	-	-	-	-	3	6	1	23.0	0.0	-	15	149.8	0.1	-	
Quarantinable diseases	Avian influenza in humans (AIH) COVID-19	076 081	-	1,082	- 6	35	- 20	-	- 85	12	1,240	463	3,277	3,822	2,583	991.6	2.6	-	20,610	2,407.6	8.6	7,435.3	
	Cholera	008	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	1.2	-	-	
	MERS-CoV	079 025	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Plague Rabies	023	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Severe acute respiratory syndrome (SARS)	071	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Smallpox Viral haemorrhagic fever (NEC)	069 036	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Yellow fever	041	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
Sexually transmissible infections	Chlamydial infection	007	81	994	11	848	198	46	258	404	2,840	2,662	3,351	45,569	19,523	24,270.8	0.8	-	84,340	100,343.4	0.8	-	
	Donovanosis Gonococcal infection	010 011	19	262	- 10	210	- 48	12	223	117	901	1,130	1,095	15,438	6,866	7,095.4	1.0	-	27,396	28,605.2	1.0	-	
	Syphilis < 2 years	066	1	26	9	34	8	-	71	28	177	175	189	3,061	1,399	1,172.4	1.2	-	5,354	4,633.6	1.2	-	
	Syphilis > 2 years or unspecified duration Syphilis congenital	067 047	1	2	-	1	-	-	33	- 8	45 1	- 63	- 62	972 9	395 3	534.0 1.6	0.7 1.9	-	1,881 17	2,188.6 8.0	0.9 2.1	1.9	
Vaccine preventable diseases	Diphtheria	009	-	-	-	-	-	-	-	-	-	1		3	2	0.6	3.3	-	8	8.0	1.0	-	
	Haemophilus influenzae type b	012	-	-	-	-	-	-	-	-	-	2	3	12	5	5.4	0.9	-	21	19.4	1.1	-	
	Influenza (laboratory confirmed) Measles	062 021	- 4	- 5	-	13	-		5	2	31	28	95	436	212	34,561.8 13.4	0.0	-	903	164,345.4 123.8	0.0	-	
	Mumps	043	-	-	-	-	-	-	-	-	-	2			3	134.0	0.0	-	36	606.2	0.1	-	
	Pertussis	024	1	2	-	9	2	-	3	2	19	40		362	190	2,395.2	0.1	-	660	14,832.6	0.0	-	
	Pneumococcal disease (invasive) Poliovirus infection	065 026	- 2	28	- 4	- 15	- 6	-	10	15 -	- 81	100	- 60	801	496	483.2	1.0	-	1,341	1,863.0	0.7	-	
	Rotavirus	077	1	12	-	20	12	-	NN	12	62	53	57	692	355	782.8	0.5	-	1,174	4,646.4	0.3	-	
	Rubella Rubella congenital	029 046	-	-	-	-	-	-	-	-	-	-	-	1	-	3.4	-	-	2	13.2	0.2	-	
	Tetanus	033	-	-	-	-	-	-	-	-	-	-	1	3	-	1.0	-	-	- 6	4.0	1.5	-	
	Varicella zoster (chickenpox)	073	12		-	3	14	-	24	20	73	58			504	735.8	0.7	-	2,609	3,597.4	0.7	-	
	Varicella zoster (shingles) Varicella zoster (unspecified)	074 075	27 1	NN NN	17 8	15 357	85 45		61 153	65 90	282 666	279 558	633 491	5,869 9,970	2,051 4,758	2,927.6 3,370.0	0.7 1.4	443.2	12,928 16,347	11,594.0 14,108.2	1.1 1.2	-	
Vectorborne diseases	Barmah Forest virus infection	048	-	5	-	10	-	-	1	-	16	12			96	138.8	0.7	-	507	402.2	1.3	-	
	Chikungunya virus infection	078	-	-	-	-	-	-	-	-	-	-	- 2	2	- 1	9.8	-	-	4	80.4	0.0	-	
	Dengue virus infection Flavivirus infection (unspecified)	003 001	-	-	-	-	-	 -	-	-	-	-	- 2	3	1	307.2 6.8	0.0	-	3	1,310.2 32.6	0.0	-	
	Japanese encephalitis virus infection	059	-	-	-	-	-	-	-	-	-	-	-	1	1	0.2	5.0	-	1	1.2	0.8	-	
	Malaria Murray Valley encephalitis virus infection	020 049	-	-	-	-	-	-	-	- 2	2	2		22	11	66.2	0.2	1.0	45 1	342.8 0.2	0.1 5.0	-	
	Ross River virus infection	002		15	4	27	2	1	3	19	71	132		2,662	931	1,692.6	0.6	-	3,713	4,740.8	0.8		
	West Nile/Kunjin virus infection	060	-	-	-	-	-	-	-	-	-	-	-	-	-	1.0		-	-	1.6	-	-	
Zoonoses	Anthrax Australian bat lyssavirus infection	058 063	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Brucellosis	003	-	-	-	1	-	-	-	-	1	-	1		8	4.0	2.0	-	18	19.0	0.9	-	
	Leptospirosis	017	-	1	-	-	-	-	1	1	3	9	l		90	37.2	2.4	27.1	225	118.6	1.9	49.3	
	Lyssavirus infection (NEC) Ornithosis	064 023	-	- 2	-	-	-	-	-	-	- 2	-	- 2	- 15	- 8	6.6	1.2	-	- 53	22.4	2.4	9.5	
	Q fever	027	-	2	-	4	-	-	2	-	8	12	19	301	131	125.8	1.0	-	474	541.2	0.9	-	
	Tularaemia	070	-	- 5	-	- 2	-	-	- 1	-	-	- 15	-	- 207	- 112	0.4		-	- 525	0.4	- 1 2	-	
Other notifiable diseases	Legionellosis Leprosy	015 016	-	-	-	- 2	-	-	-	1	8	- 15	- 8	307 5	112	106.8 3.0	1.0 1.3	-	535 10	420.8 11.4	1.3 0.9	-	
	Meningococcal disease (invasive)	022	-	-	-	-	-	-	-	-	-	2	5	42	18	49.8	0.4	-	78	254.8	0.3	-	
	RSV^	083 034	1	- 24	-	82 6	-	- 1	- 15	- 14	83 60	9 57		92 787	92 360	339.4	1.4	-	92 1,591	1,436.2	1.1	-	
	Tuberculosis iGAS^	082	-	- 24	-	15	-	-	- 15	- 14	15	3		22	22	339.4	1.1	-	1,591	1,436.2	1.1	-	
Footnotes:			189	3,029	113	2,310	582	130	1,370	978	8,706	7,973	11,464	128,002	55,346				243,366				

The data in this report are reliant on the provision of data from states and territories to the Australian Government Department of Health. Backlogs in notifications at the state or territory level may contribute to delays in reporting to the NNDSS. Notifications for some high volume conditions are only uploaded quarterly by some jurisdictions, which can result in apparent large variability over time. The NNDSS is a dynamic dataset, with data in this report representing data available on (20/07/2021). Data in this report are subject to retrospective revision and may vary from data reported in published NNDSS reports and reports of notification data by states and territories.

Footnotes:

* Natio of the 90 day prior surveillance period to the past 90 day 5 year rolling mean, or ratio of the year period prior surveillance period to the year rolling mean.

NN = Not Notifiable, NEC = Not Elsewhere Classified

* RSV and iGAS were listed as nationally notifiable diseases as of 1 July 2021. However, notification numbers presented here do not represent a national picture, as these conditions are not yet notifiable in all states and territories.