National Communicable Diseases Surveillance Report

Fortnight 23, 2021 Summary Notes for Selected Diseases

08 November to 21 November 2021

<u>Infectious and congenital syphilis</u>

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) predominately 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 largely 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.

For further information on national activities related to 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 (24/08/2021 to 21/11/2021).

²The quarterly (90 day) five year rolling mean is the average of 5 intervals of 90 days up to 21/11/2021. 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 (22/11/2020 to 21/11/2021).

⁴The yearly (365 day) five year rolling mean is the average of 5 intervals of 365 days up to 21/11/2021. 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.

	T FN22/2024										Notification received date												
ADT FN23/2021				State or Territory								Totals for Australia				Historical 90 Day Period				Historical Yearly Period			
Disease group	Disease name	Disease code	АСТ	NSW	M	old	SA	Tas	Vic	WA	This reporting period	Previous reporting Period	Same reporting period last year	Current year YTD	Past Quarter	Quarterly rolling 5 year	Ratio past quarter/5	Exceeds quarterly rolling	Past Year	Yearly rolling 5 year mean	Ratio past year/5 year	Exceeds yearly rolling	
											08/11/2021	25/10/2021	08/11/2020	01/01/2021	24/08/2021	mean	year mean*	mean +2 SD by	22/11/2020	22/11/2015	mean*	mean +2 SD by	
Bloodborne diseases	Hepatitis B (newly acquired)	039	-		-	-	1	-	-	-	21/11/2021	07/11/2021	21/11/2020	21/11/2021 75	21/11/2021	33.4	0.3	-	21/11/2021 87	21/11/2020 149.2	0.6	-	
	Hepatitis B (unspecified)	052	2	96	-	48	-	3	60	15	224	227	205	4,338	1,268	1,386.6	0.9	-	4,808	5,739.2	0.8	-	
	Hepatitis C (newly acquired) Hepatitis C (unspecified)	040 053	- 5	- 98	- 1	10 68	-	- 4	- 52	- 38	10 266	21 297	19 272	648 6,230	157 1,626	179.6 2,215.6	0.9	-	711 7,029	701.2 9,452.2	1.0 0.7	-	
	Hepatitis D	050	-	1	-	-	1	-	2	1	5	3	-	80	21	18.8	1.1	-	88	70.4	1.3		
Gastrointestinal diseases	Botulism Campylobacteriosis	045 005	- 38	482	7	418	166	- 46	508	150	1,815	1,647	1,645	32,550	9,003	8,063.4	1.1	-	3 36,731	1.0 30,654.0	3.0 1.2	-	
	Cryptosporidiosis	061	-	13	-	27	1	2	23	4	70	57	50	1,625	329	473.0	0.7	-	1,761	3,772.0	0.5	-	
	Haemolytic uraemic syndrome (HUS) Hepatitis A	055 038	-	-	-	-	-	-	-	- 1	- 1	1	-	19	10	2.8 42.6	1.4 0.2	0.3	8 19	15.4 231.2	0.5 0.1	-	
	Hepatitis E	051	-	-	-	-	-	-	-	-	-	-	-	10	1	8.0	0.1	-	10	46.0	0.2	-	
	Listeriosis Paratyphoid	018 080	-	- 2	-	- 1	-	-	-	- 1	- 4	1	- 4	43	14	15.4 12.4	0.9	-	50	65.6 79.8	0.8	-	
	Salmonellosis	030	2	148	25	192	40	6	98	27	538	396	309	9,639	2,144	2,689.4	0.8	-	10,803	15,228.8	0.7	-	
	Shigellosis STEC	031 054	-	3 8	9	1	3 11	- 1	3	- 3	19 29	13 34	40 21	410 527	105 155	476.8 128.8	0.2 1.2	-	476 592	2,103.2 515.0	0.2 1.1	-	
	Typhoid Fever	035	-	-	-	-	-	·	-	-	-	-	1	12	4	25.2	0.2	-	17	144.6	0.1	-	
Quarantinable diseases	Avian influenza in humans (AIH) Cholera	076 008	-	-	-	-	-	-	-	-	-	-	-	- 1	- 1	0.4	2.5	-	- 1	1.2	0.8	-	
	COVID-19	081	147	2,996	36	14	4	-	14,856	7	18,060	21,417	160	171,386	153,695	599.8	256.2	150,412.8	172,017	5,588.2	30.8	141,437.6	
	Middle East respiratory syndrome coronavirus (I Plague	079 025	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Rabies	028	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Severe acute respiratory syndrome (SARS) Smallpox	071 069	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Viral haemorrhagic fever (NEC)	036	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
Sexually transmissible infections	Yellow fever Chlamydial infection	041	- 57	- 774	- 52	- 920	- 252	- 58	- 452	- 458	3,022	3,073	3,356	77,318	19,021	24,056.6	0.8	-	- 86,322	100,474.0	0.9	-	
	Donovanosis	010	-	-	-	-	-	-	-	-		-	-	-	- 15,021	-	0.8	-		-	0.9	-	
	Gonococcal infection Syphilis < 2 years	011	17	220 28	22	264 14	46 8	4	76 49	131 27	780 136	935 186	1,015 195	23,543 4,939	5,554 1,217	7,085.8 1,244.8	0.8 1.0	-	26,241 5,458	29,554.2 4,824.4	0.9	-	
	Syphilis > 2 years or unspecified duration	067	-	28	3	3	-	1	55	4	68	48	75	1,595	353	527.4	0.7	-	1,812	2,147.8	0.8	-	
Vaccine preventable diseases	Syphilis congenital Diphtheria	047	-	-	-	-	-	-	-		-	-	1	14	2	2.2 1.6	0.9	-	17 8	7.6 8.4	2.2 1.0	-	
	Haemophilus influenzae type b	012	-	-	-	1	-	-	-	-	1	-	-	17	4	4.4	0.9	-	19	19.2	1.0	-	
	Influenza (laboratory confirmed) Measles	062 021	-	9	1	14	6	2	2	3	37	17	29	655	163	54,863.0 31.4	0.0	-	746	147,544.6 120.2	0.0	-	
	Mumps	043	-	-	-	-	-	-	-	-	-	-	1	20	7	100.0	0.1	-	23	543.0	0.0	-	
	Pertussis	024 065	- 2	- 4	-	8 10	- 6	-	14 5	2	24	23 41	24 48	523	134 300	3,206.2	0.0	-	584	13,026.4	0.0	-	
	Pneumococcal disease (invasive) Poliovirus infection	026	-	- 4	- 4	-	-	- 1	-	-	41 -	- 41	- 48	1,252	-	548.2	0.5	-	1,386	1,802.2	0.8	-	
	Rotavirus	077 029	-	13	46	39	49	1	5	75	228	205	34	1,825	985	1,641.8	0.6	-	1,949	4,304.6	0.5	-	
	Rubella Rubella congenital	046	-	-	-	-	-	-	-	-	-	-	-	-	-	1.2	0.8	-	-	11.8	0.3	-	
	Tetanus	033 073	- 4	- NN	- 2	-	- 17	- 2	- 15	- 23	- 63	-	- 141	1 761	2	0.8		-	7 2 100	4.0 3,632.2	1.8	0.6	
	Varicella zoster (chickenpox) Varicella zoster (shingles)	074	23	NN	13	7	98	_	15 63	80	63 300	65 398	141 604	1,761 9,526	411 2,321	1,090.6 3,189.0	0.4	-	2,108 11,094	12,287.0	0.6 0.9	-	
	Varicella zoster (unspecified) Barmah Forest virus infection	075 048	-	NN 6	- 6	396 8	43	22	298	140	903 16	927 9	599 26	17,815 338	5,567 69	3,467.0 75.4	1.6 0.9	1,180.4	19,521 389	14,079.0 412.2	1.4 0.9		
Vectorborne diseases	Chikungunya virus infection	078	-	-	-	- 8	-	-	-	-	-	-	-	2	-	23.2	-	-	2	76.4	0.9	-	
	Dengue virus infection Flavivirus infection (unspecified)	003 001	-	-	-	-	-	-	-	-	-	1	- 1	5	2	208.0 7.6		-	5 5	1,237.0 33.4	0.0 0.1	-	
	Japanese encephalitis virus infection	059	-	-	-	-	-	-	-	-	-	-	-	1	-	0.2		-	1	1.2		-	
	Malaria Murray Valley encephalitis virus infection	020 049	-	-	-	2	-	-	-	-	2	-	-	41 1	9	75.6	0.1	-	47 1	329.6 0.2	0.1 5.0	-	
	Ross River virus infection	002	-	1	1	11	- 6	-	1	12	32	45		3,041	242	479.6	0.5	-	3,403	4,609.0	0.7	-	
Zoonoses	West Nile/Kunjin virus infection Anthrax	060 058	-	-	-	-	-	-	-	-	-	-	-	-	-	0.2	-	-	-	1.2	-	-	
	Anthrax Australian bat lyssavirus infection	063	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Brucellosis Lentocnirosis	004	-	-	-	- 1	-	-	-	-	- 1	- 1	1		3	4.6		-	18	18.8	1.0	-	
	Leptospirosis Lyssavirus infection (NEC)	017 064	-	-	-	- 1	-	-	-	-	- 1	1	-	244	- 23	16.8	1.4	-	253 -	118.8	2.1	-	
	Ornithosis	023	-	1	-	-	-	-	-	-	1	- 14	2		4	9.6		-	36	25.8	1.4	-	
	Q fever Tularaemia	027 070	-	- 4	-	- 11	-	-	-	-	15 -	- 14	16 -	439	- 88	121.4	0.7	-	472	526.8 0.4	0.9	-	
Other notifiable diseases	iGAS^	082	NN	NN	3	10		NN	NN	1	16	13		168	92	- 100.6	-	-	168	-	- 13		
	Legionellosis Leprosy	015 016	- 1	1	-	- 1	-	-	- 5	- 3	14 1	22 1		448 13	96 6	108.6 3.4	0.9 1.8	-	534 14	423.4 10.6	1.3 1.3	-	
	Meningococcal disease (invasive)	022	-	1	-	-	2	_	-	-	3	1	1	64	16	81.4	0.2	-	72	244.0	0.3	-	
	RSV^ Tuberculosis	083 034	NN -	NN 22	36 1	24 9	- 8	NN 3	NN 19	5	69 59	79 38	- 67	743 1,282	482 317	393.0	- 0.8	-	743 1,486	1,465.2	1.0		
Footnotes:			300	4,914	241	2,502	763	169	16,643	1,217	26,746	30,142	9,051	373,269	205,240				397,932		. '		

^{**} Ratio 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 period 5 year rolling mean.

NN = Not Notifiable, NEC = Not Elsewhere Classified

NN = NOT NOTIFICATION, NEC. = NOT ESSEWNERE Classified

A 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.

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 (22/11/2021). 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.