National Communicable Diseases Surveillance Report

Fortnight 22, 2021 Summary Notes for Selected Diseases

25 October to 07 November 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) 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 (10/08/2021 to 07/11/2021).

²The quarterly (90 day) five year rolling mean is the average of 5 intervals of 90 days up to 07/11/2021. 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 (08/11/2020 to 07/11/2021).

⁴The yearly (365 day) five year rolling mean is the average of 5 intervals of 365 days up to 07/11/2021. 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.

۸۵	T FN22/2021											Notification received date											
AL	ADT FN22/2021					ate or	Territo	ry			Totals for Australia				Historical 90 Day Period				Historical Yearly Period				
Disease group	Disease name	isease code	ACT	NSW	NT	Qid	8A	Tas	Vic	WA	This reporting period	Previous reporting Period	Same reporting period last year	Current year YTD	Past Quarter	Quarterly rolling 5 year mean	Ratio past quarter/5 year mean*	Exceeds quarterly rolling mean +2 SD	Past Year	Yearly rolling 5 year mean	Ratio past year/5 year mean*	Exceeds yearly rolling mean +2 SD	
		Q									25/10/2021 07/11/2021	11/10/2021 24/10/2021	25/10/2020 07/11/2020	01/01/2021 07/11/2021	10/08/2021 07/11/2021	mean		by	08/11/2020 07/11/2021	08/11/2015 07/11/2020		by	
Bloodborne diseases	Hepatitis B (newly acquired) Hepatitis B (unspecified)	039 052	- 2	- 77	-	1 43	-	- 1	- 58	- 22	203	- 187	- 188	72 4,084	11 1,183	33.2 1,409.6	0.3	-	95 4,760	148.2 5,742.0	0.6 0.8	-	
	Hepatitis C (newly acquired)	040	-	1	-	9	-	-	-	1	11	31	28	622	163	178.0	0.9	-	704	702.4	1.0	-	
	Hepatitis C (unspecified) Hepatitis D	053 050	1	116 2	3	78	- 1	- 6	45	30	280	250 4	279 3	5,944 73	1,602 17	2,235.4 21.2	0.7 0.8	-	7,015 81	9,468.2 70.6	0.7 1.1	-	
Gastrointestinal diseases	Botulism	045	-	-	-	-	-	-	-	-	-	-	-	2	-	-	0.0	-	3	1.0	3.0	-	
	Campylobacteriosis	005 061	46	487 9	11	355 13	135	42	294 18	154	1,524 51	1,452 40	1,498 48	30,560 1,548	8,220 314	7,720.8 443.4	1.1 0.7	-	36,373 1,734	30,527.0 3,793.8	1.2 0.5	-	
	Cryptosporidiosis Haemolytic uraemic syndrome (HUS)	055	-	-	-	-	1	-	-	-	1	- 40	1	7,548	314	3.2	0.7	-	7	15.6	0.5	-	
	Hepatitis A	038 051	-	1	-	1	-	-	-	-	2	2	-	20 10	12	43.2 7.4	0.3	-	20 10	230.8 46.6	0.1 0.2	-	
	Hepatitis E Listeriosis	051	-	1	-	-	-	-	-	-	1	2	1	39	11	14.8	0.1	-	50	46.6 65.8	0.2	-	
	Paratyphoid	080	-	-	-	-	-	-	-	-	-	-	-	3	- 4.700	10.0	-	-	3	80.2	0.0	-	
	Salmonellosis Shigellosis	030	7	119 1	21	134	24	1	36	24 5	367 13	381 20	330 26	9,068 391	1,766 109	2,551.4 468.4	0.7	-	10,541 497	15,283.0 2,100.8	0.7 0.2	-	
	STEC	054	-	7	-	2	7	-	6	7	29	16	27	493	131	119.8	1.1	-	579	512.6	1.1	-	
Quarantinable diseases	Typhoid Fever Avian influenza in humans (AIH)	035 076	-	-	-	-	-	-	-	-	-	-	- 1	- 12	- 4	24.0	0.2	-	18	144.4	0.1	-	
	Cholera	008	-	-	-	-	-	-	-	-	-	-	-	1	1	0.4	2.5	-	1	1.2	0.8	-	
	COVID-19 Middle East respiratory syndrome coronaviru	081 is (M 079	146 -	3, <u>222</u> -	- 4	11 -	-	-	17,960 -	- 1	21,344	30,479	124 -	152,685 -	144,177 -	1,283.8	112.3	137,151.9	153,413 -	5,451.6 -	28.1	123,581.1	
	Plague	025	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Rabies Severe acute respiratory syndrome (SARS)	028 071	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Smallpox	069	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
	Viral haemorrhagic fever (NEC) Yellow fever	036 041	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-		-	
Sexually transmissible infections	Chlamydial infection	007	45	806	66	937	230	55	344	411	2,894	2,951	3,358	74,120	19,099	24,070.4	0.8	-	86,480	100,506.4	0.9	-	
	Donovanosis Gonococcal infection	010 011	- 8	- 218	- 51	- 216	- 42	- 7	184	106	832	- 804	- 997	22,582	- 5,466	7,108.2	0.8	-	- 26,295	- 29,467.8	0.9	-	
	Syphilis < 2 years	066	1	28	3	39	2	-	45	28	146	181	193	4,749	1,223	1,239.4	1.0	-	5463	4808.8	1	-	
	Syphilis > 2 years or unspecified duration Syphilis congenital	067 047	-	2	-	3	2	-	33	3	43	43	67	1,509 13	341	525.2 2.4	0.6 1.3	-	1,802 17	2,145.0 7.8	0.8 2.2	-	
Vaccine preventable diseases	Diphtheria	009	-	-	-	-	-	-	-	-	-	-	-	6	2	1.6	1.3	-	8	8.2	1.0	-	
	Haemophilus influenzae type b Influenza (laboratory confirmed)	012 062	-	- 4	- 3	-	- 2	-	- 3	- 1	- 19	- 28	21	16 621	3 156	4.0 70,589.4	0.8	-	18 741	19.2 147,678.6	0.9	-	
	Measles	002	-	-	-	-	-	-	-	-	-	-	-	- 021	-	32.0	-	-	-	120.4	-	-	
	Mumps	043 024	-	- 4	- 1	- 4	-	-	- 8	-	- 22	2 19	2 13	20 497	7 132	103.0 3,059.8	0.1	-	24 582	553.0 13,321.0	0.0	-	
	Pertussis Pneumococcal disease (invasive)	065	-	8	2	10	9	1	7	2	39	44	39	1,210	322	576.4	0.6	-	1,392	1,804.0	0.8	-	
	Poliovirus infection Rotavirus	026 077	-	- 13	- 43	- 33	- 29	-	- 1	- 69	188	- 146	- 41	1,582	- 791	1,588.8	0.5	-	1,740	4.344.0	0.4	-	
	Rubella	029	-	-	-	-	-	-	-	-	-	-	-	2	1	1,566.6	0.5	-	3	12.0	0.4	-	
	Rubella congenital	046 033	-	-	-	-	-	-	-	-	-	-	-	- 5	- 2	-	2.5	-	- 7	- 4.2	1.7	-	
	Tetanus Varicella zoster (chickenpox)	033	- 8	NN	1	3	- 6	4	11	23	56	- 66		1,686	397	1,087.0	2.5 0.4	-	2,174	3,625.8	0.6	-	
	Varicella zoster (shingles)	074	33		7	8	88	9	77	99	321	351	661	9,103	2,295	3,161.4	0.7	-	11,275	12,212.0	0.9	-	
	Varicella zoster (unspecified) Barmah Forest virus infection	075 048	-	NN 2	- 9	436 4	72 -	- 29	219	135	904	871 10	548 21	16,888 320	5,466 60	3,467.2 76.6	1.6 0.8	-	19,193 397	14,062.2 409.2	1.4 1.0	-	
Vectorborne diseases	Chikungunya virus infection	078	-	-	-	-	- 1	-	-	-	- 1	- 1	-	2	-	21.8	-	-	2	77.4	0.0	-	
	Dengue virus infection Flavivirus infection (unspecified)	003	-	-	-	-	-	-	-	-	- 1	- 1	- 1	5 3	- 3	206.4	0.0	-	5 6	1,243.8 33.0	0.0	-	
	Japanese encephalitis virus infection	059	-	-	-	-	-	Ŀ	-	-	-	-	-	1	-	-	0.5	-	1	1.2	0.8	-	
	Malaria Murray Valley encephalitis virus infection	020 049	-	-	-	-	-	-	-	-	-	- 1	- 1	39 1	- 11	76.2	0.1	-	45 1	332.4 0.2	0.1 5.0	-	
	Ross River virus infection	002	-	8	3	11	1	-	1	16	40	39		3,002	251	467.2	0.5	-	3,487	4,619.4	0.8	-	
Zoonoses	West Nile/Kunjin virus infection Anthrax	060 058	-	-	-	-	-	-	-	-	-	-	-	-	-	0.2	-	-	-	1.4	-	-	
	Australian bat lyssavirus infection	063	-	-	-	-	-	·	-	-	-	-	-	-	-	-		-	-	-		-	
	Brucellosis Leptospirosis	004 017	-	-	-	-	-	-	-	- 1	- 1	- 8	- 2	17 243	24	5.0 18.6	0.8 1.3	-	19 255	19.0 119.0	1.0 2.1	74.4	
	Lyssavirus infection (NEC)	064	-	-	-	-	-		-	Ė	-	-	-	-	-	-		-	-	-		-	
	Ornithosis Q fever	023 027	-	- 4	-	- 8	-	-	-	-	- 12	- 12	4 19	21 420	4 88	9.6 123.2	0.4	-	37 469	25.6 527.4	1.4 0.9	-	
	Tularaemia	070	-	-	-	-	-		-	-	-	-	-	- 420	-	-	0.7	-	- 409	0.4	-	-	
Other notifiable diseases	iGAS	082	NN -	NN 8	- 2	8	1	_	NN 1	2 8	13 20	12		151 429	93	- 99.0	0.9		151 533	424.8	1 2		
	Legionellosis Leprosy	015 016	-	- 8	-	1	-	-	- 1	- 8	1	18			94	99.0 3.4	2.1	-	14	10.4	1.3	-	
	Meningococcal disease (invasive)	022	-	1	- 42	-	-	-	-	-	1	4		62	17	83.2	0.2	-	71	244.0	0.3	-	
	RSV Tuberculosis	083 034	NN -	NN 10	42	24 7	- 7	NN -	NN 10	9	78 37	90 41	70	673 1,222	454 299	392.0	0.8	-	673 1,493	1,466.2	1.0	-	
			301	5,149	235	2,376	660	160	19,352	1,156	29,389	38,477	8,814	344,971	194,087		•		378,611		ı		

Footnotes:

* 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

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 (08/11/2021). 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.