

Surveillance Data **Synopsis** 

• RSV activity has

consin.

consin.

The Delta

increased in Wis-

• SARS-CoV-2 activ-

ity continues to

increase in Wis-

[B.I.617.2] SARS-

**CoV-2** variant of

concern is by far

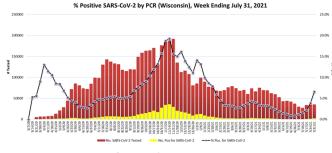
the most common lineage detected.

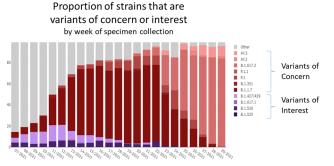
# Laboratory Surveillance Report

## SARS-CoV-2 Surveillance Updates

#### SARS-CoV-2 Update

- In Wisconsin, positivity was 6.6% of the 35,538 specimens tested by PCR and reported to WSLH.
- In the US, 7 day average percent positivity increased to 9.5%.
- Wisconsin genomic sequencing data showed the Delta [B.I.617.2] variant of concern was by far the predominant lineage detected.
- The WSLH sequencing dashboard is available at https:// dataportal.slh.wisc.edu/sc2dashboard





To enhance surveillance activities, the WSLH asks labs to <u>please send</u>:

- Ι. Please send all positive influenza specimens for further characterization.
- 2. Please send a maximum of 10 SARS-CoV-2 specimens per week.

<b>Other Surveil</b>	lance	Data-	<b>Wisconsin</b>

Week Ending July 31, 2021*						
		0/ <b>D</b>				

Resp. Pathogen PCR	# Tested	% Positive
SARS-CoV-2	35538	6.6个
Parainfluenza	823	11.1↓
Rhinovirus/ Enterovirus	776	13.5
RSV	1602	8.8个
Seasonal coronaviruses	72	2.8
Adenovirus	72	1.4
Influenza	2078	<1
Human metapneumovirus	825	0
B. pertussis	272	0

### **Respiratory**

There continues to be an increase in SARS-CoV-2 detections reported in Wisconsin.

#### Gastropathogens

- Campylobacter and Cylcospora were the predominant gastropathogens reported.
- Others detected included: EPEC (14.9%), EAEC (3.1%), ETEC (1.3%), Aeromonas (1.2%) and P. shigelloides (<1%).

Week Ending July 31, 2021*				
GI Pathogen PCR	# Tested	% Positive		
Campylobacter	717	3.3		
Cyclospora	161	3.1		
Norovirus	495	3		
Salmonella	717	2		
STEC	639	2		
Rotavirus	487	1.2		
Cryptosporidium	468	1.1		
Giardia	468	<1		
Shigella	626	<1		
Sapovirus	173	0		
E. coli 0157	321	0		

\* On a weekly basis, participating Wisconsin clinical laboratories voluntarily report to WSLH the total number of tests performed, the method used for detection, and the number of those tests with positive results.