

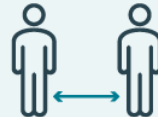


COVID-19

ACT NOW!



WEAR A MASK



STAY 6 FEET APART



AVOID CROWDS

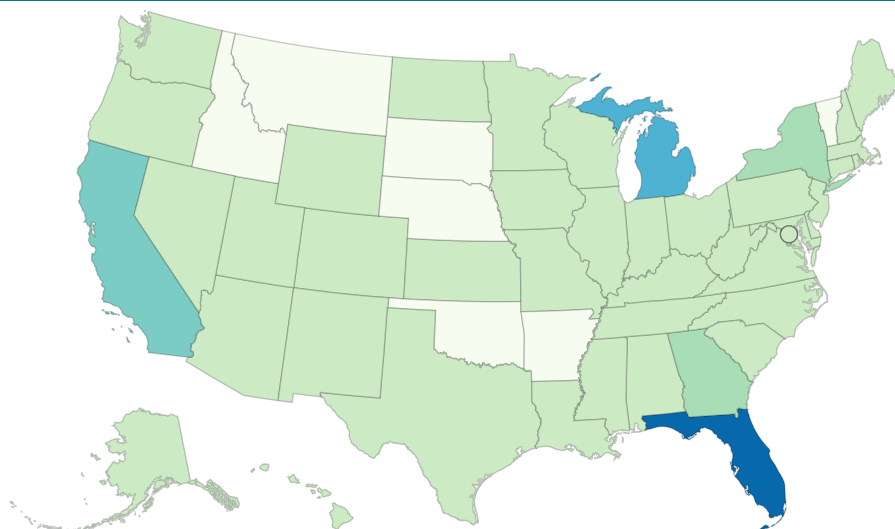
US COVID-19 Cases Caused by Variants

Updated Feb. 25, 2021

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Variant	Reported Cases in US	Number of Jurisdictions Reporting
B.1.1.7	2102	45
B.1.351	49	15
P.1	6	5

Emerging Variant Cases in the United States*†



Number of Cases

- 0 to 0
- 1 to 100
- 101 to 200
- 201 to 300
- 301 to 400
- 401 to 500

Filters

Variant B.1.1.7

Territories AS GU MH FM MP PW PR VI



Data Table

Location	B.1.1.7 Variant	P.1 Variant	B.1.351 Variant
<input checked="" type="radio"/> Alabama	35	0	0
<input checked="" type="radio"/> Alaska	2	1	0
<input type="radio"/> American Samoa	0	0	0
<input checked="" type="radio"/> Arizona	14	0	0
<input type="radio"/> Arkansas	0	0	0
<input checked="" type="radio"/> California	204	0	2
<input checked="" type="radio"/> Colorado	86	0	0
<input checked="" type="radio"/> Connecticut	42	0	1
<input checked="" type="radio"/> Delaware	7	0	0
<input checked="" type="radio"/> Florida	500	1	0
<input checked="" type="radio"/> Georgia	119	0	0
<input type="radio"/> Guam	0	0	0
<input checked="" type="radio"/> Hawaii	2	0	0
<input type="radio"/> Idaho	0	0	1
<input checked="" type="radio"/> Illinois	42	0	1
<input checked="" type="radio"/> Indiana	16	0	0
<input checked="" type="radio"/> Iowa	10	0	0
<input checked="" type="radio"/> Kansas	2	0	0
<input checked="" type="radio"/> Kentucky	5	0	0
<input checked="" type="radio"/> Louisiana	17	0	0
<input checked="" type="radio"/> Maine	2	0	0
<input type="radio"/> Marshall Islands	0	0	0
<input checked="" type="radio"/> Maryland	68	1	9
<input checked="" type="radio"/> Massachusetts	51	0	2
<input checked="" type="radio"/> Michigan	336	0	0
<input type="radio"/> Micronesia	0	0	0
<input checked="" type="radio"/> Minnesota	57	2	0
<input checked="" type="radio"/> Mississippi	1	0	0
<input checked="" type="radio"/> Missouri	1	0	0
<input type="radio"/> Montana	0	0	0
<input type="radio"/> Nebraska	0	0	0
<input checked="" type="radio"/> Nevada	7	0	1
<input checked="" type="radio"/> New Hampshire	1	0	0
<input checked="" type="radio"/> New Jersey	55	0	0
<input checked="" type="radio"/> New Mexico	4	0	0
<input checked="" type="radio"/> New York	136	0	1
<input checked="" type="radio"/> North Carolina	40	0	3
<input checked="" type="radio"/> North Dakota	4	0	0
<input type="radio"/> Northern Marianas	0	0	0
<input checked="" type="radio"/> Ohio	19	0	0
<input type="radio"/> Oklahoma	0	1	0

<input checked="" type="radio"/> Oregon	9	0	0
<input type="radio"/> Palau	0	0	0
<input checked="" type="radio"/> Pennsylvania	39	0	0
<input checked="" type="radio"/> Puerto Rico	1	0	0
<input checked="" type="radio"/> Rhode Island	3	0	0
<input checked="" type="radio"/> South Carolina	7	0	21
<input type="radio"/> South Dakota	0	0	0
<input checked="" type="radio"/> Tennessee	13	0	1
<input checked="" type="radio"/> Texas	60	0	1
<input checked="" type="radio"/> Utah	12	0	0
<input type="radio"/> Vermont	0	0	0
<input type="radio"/> Virgin Islands	0	0	0
<input checked="" type="radio"/> Virginia	12	0	3
<input checked="" type="radio"/> Washington	39	0	1
<input checked="" type="radio"/> Washington D.C.	4	0	1
<input checked="" type="radio"/> West Virginia	3	0	0
<input checked="" type="radio"/> Wisconsin	6	0	0
<input checked="" type="radio"/> Wyoming	9	0	0

[Download Data \(CSV\)](#)

[Download Accessible Data](#)  [CSV – 665 KB]

The emerging variants CDC is closely monitoring have mutations in the virus genome that alter the characteristics and cause the virus to act differently in ways that are significant to public health (e.g., causes more severe disease, spreads more easily between humans, requires different treatments, changes the effectiveness of current vaccines). It's important to understand that genetic mutations are expected, and some variants can spread and become predominant while others subside.

02/11/21 – To maintain a limited number of colors and allow for an easier comparison between states, the data ranges have changed from 1-50 to 1-100.

*The cases identified above are based on a sampling of SARS-CoV-2-positive specimens and do not represent the total number of B.1.1.7, B.1.351, and P.1 lineage cases that may be circulating in the United States and may not match numbers reported by states, territories, tribes, and local officials.

†Numbers will be updated on Sunday, Tuesday, and Thursday by 7:00 pm.

Learn more about [Genomic Surveillance for SARS-CoV-2 Variants](#).

Last Updated Feb. 25, 2021

Content source: [National Center for Immunization and Respiratory Diseases \(NCIRD\), Division of Viral Diseases](#)