

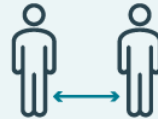


COVID-19

ACT NOW!



WEAR A MASK



STAY 6 FEET APART



AVOID CROWDS

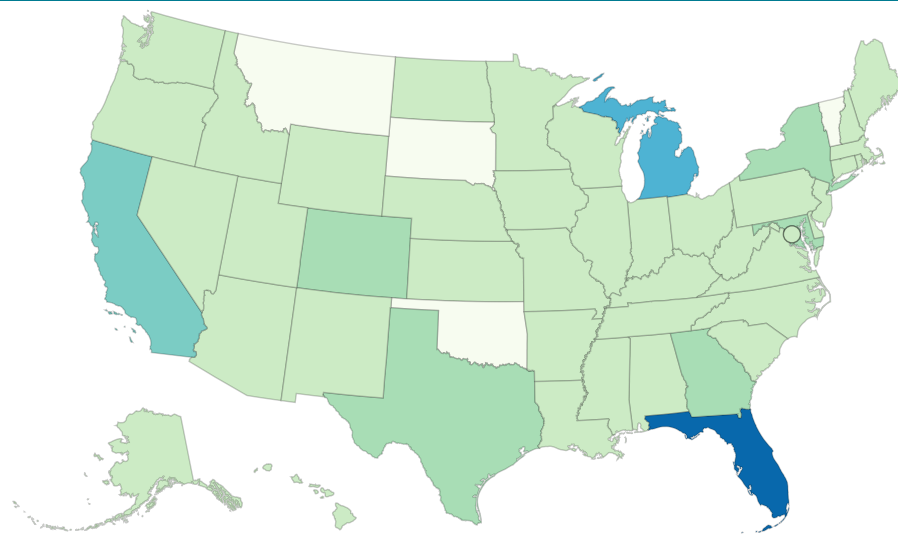
US COVID-19 Cases Caused by Variants

Updated Mar. 4, 2021

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Variant	Reported Cases in US	Number of Jurisdictions Reporting
B.1.1.7	2672	48
B.1.351	68	17
P.1	13	7

Emerging Variant Cases in the United States*†



Number of Cases

- 0 to 0
- 1 to 100
- 101 to 200
- 201 to 300
- 401 to 500
- 601 to 700

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	37	0	0
<input checked="" type="radio"/> Alaska	2	1	0
<input type="radio"/> American Samoa	0	0	0
<input checked="" type="radio"/> Arizona	19	0	0
<input checked="" type="radio"/> Arkansas	1	0	0
<input checked="" type="radio"/> California	250	0	3
<input checked="" type="radio"/> Colorado	114	0	0
<input checked="" type="radio"/> Connecticut	42	0	1
<input checked="" type="radio"/> Delaware	12	0	0
<input checked="" type="radio"/> Florida	642	5	1
<input checked="" type="radio"/> Georgia	155	0	1
<input type="radio"/> Guam	0	0	0
<input checked="" type="radio"/> Hawaii	6	0	0
<input checked="" type="radio"/> Idaho	2	0	2
<input checked="" type="radio"/> Illinois	69	0	1
<input checked="" type="radio"/> Indiana	19	0	0
<input checked="" type="radio"/> Iowa	14	0	0
<input checked="" type="radio"/> Kansas	14	0	0
<input checked="" type="radio"/> Kentucky	5	0	0
<input checked="" type="radio"/> Louisiana	18	0	0
<input checked="" type="radio"/> Maine	2	0	0
<input type="radio"/> Marshall Islands	0	0	0
<input checked="" type="radio"/> Maryland	109	1	12
<input checked="" type="radio"/> Massachusetts	57	0	4
<input checked="" type="radio"/> Michigan	421	0	0
<input type="radio"/> Micronesia	0	0	0
<input checked="" type="radio"/> Minnesota	78	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 checked="" type="radio"/> Nebraska	1	0	0
<input checked="" type="radio"/> Nevada	10	0	1
<input checked="" type="radio"/> New Hampshire	1	0	0
<input checked="" type="radio"/> New Jersey	72	2	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	22	0	0
<input type="radio"/> Oklahoma	0	1	0

<input checked="" type="radio"/> Oregon	11	1	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	30
<input type="radio"/> South Dakota	0	0	0
<input checked="" type="radio"/> Tennessee	18	0	1
<input checked="" type="radio"/> Texas	101	0	1
<input checked="" type="radio"/> Utah	23	0	0
<input type="radio"/> Vermont	0	0	0
<input type="radio"/> Virgin Islands	0	0	0
<input checked="" type="radio"/> Virginia	20	0	4
<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	14	0	0
<input checked="" type="radio"/> Wyoming	9	0	0

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

[Download Accessible Data](#)  [CSV – 674 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 Mar. 4, 2021

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