



COVID-19

ACT NOW!



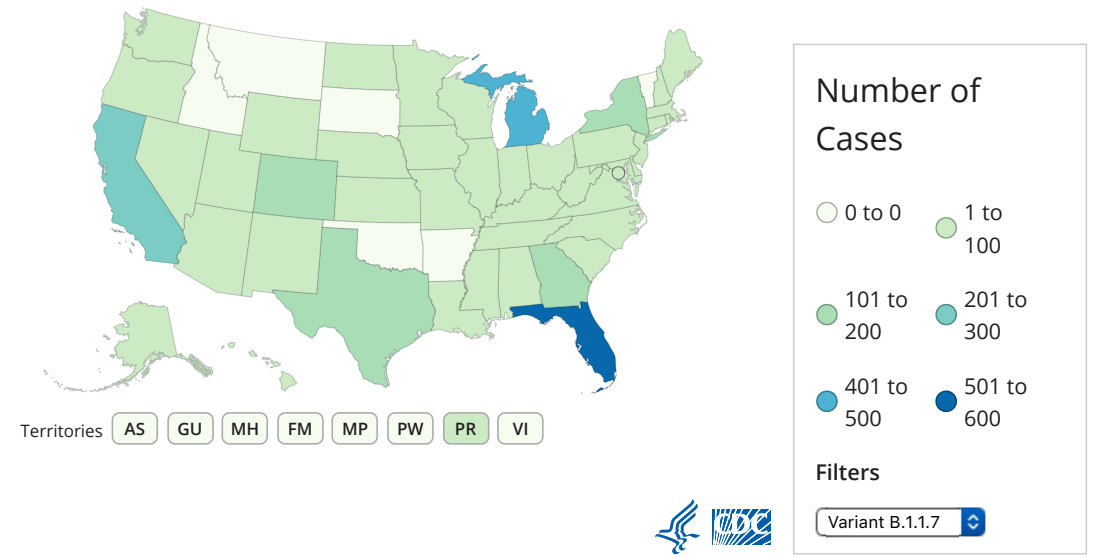
US COVID-19 Cases Caused by Variants

Updated Mar. 2, 2021

[Print](#)

Variant	Reported Cases in US	Number of Jurisdictions Reporting
B.1.1.7	2506	46
B.1.351	65	17
P.1	10	5

Emerging Variant Cases in the United States*†



Data Table			
Location	B.1.1.7 Variant	P.1 Variant	B.1.351 Variant
Alaba...	37	0	0
Alaska	2	1	0
American Sa...	0	0	0

Arizo...	14	0	0
Arkans...	0	0	0
Califor...	212	0	3
Colora...	103	0	0
Connecti...	42	0	1
Delawa...	12	0	0
Flori...	599	5	1
Geor...	137	0	1
Guam	0	0	0
Hawaii	6	0	0
Idaho	0	0	1
Illinois	69	0	1
India...	19	0	0
Iowa	14	0	0
Kans...	10	0	0
Kentuc...	5	0	0
Louisia...	18	0	0
Maine	2	0	0
Marshall Isla...	0	0	0
Maryla...	89	1	10
Massachus...	57	0	4
Michig...	421	0	0
Microne...	0	0	0
Minnes...	78	2	0
Mississi...	1	0	0
Misso...	1	0	0
Monta...	0	0	0
Nebras...	1	0	0
Neva...	10	0	1
New Hamps...	1	0	0
New Jer...	55	0	0
New Mex...	4	0	0
New Y...	136	0	1
North Caro...	40	0	3
North Dak...	4	0	0
Northern Mari...	0	0	0
Ohio	19	0	0
Oklaho...	0	1	0
Oreg...	11	0	0
Palau	0	0	0
Pennsylva...	39	0	0
Rhode I...	1	0	0

	U	U	U
<input checked="" type="radio"/> Puerto R...	1	0	0
<input checked="" type="radio"/> Rhode Isl...	3	0	0
<input checked="" type="radio"/> South Caro...	7	0	30
<input type="radio"/> South Dak...	0	0	0
<input checked="" type="radio"/> Tennes...	18	0	1
<input checked="" type="radio"/> Texas	101	0	1
<input checked="" type="radio"/> Utah	23	0	0
<input type="radio"/> Vermo...	0	0	0
<input type="radio"/> Virgin Isla...	0	0	0
<input checked="" type="radio"/> Virgi...	20	0	4
<input checked="" type="radio"/> Washing...	39	0	1
<input checked="" type="radio"/> Washington ...	4	0	1
<input checked="" type="radio"/> West Virgi...	3	0	0
<input checked="" type="radio"/> Wiscon...	10	0	0
<input checked="" type="radio"/> Wyomi...	9	0	0

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

[Download Accessible Data](#)  [CSV - 673 B]

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. 2, 2021

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