



# COVID-19



WEAR A MASK



STAY 6 FEET APART



AVOID CROWDS



GET A VACCINE

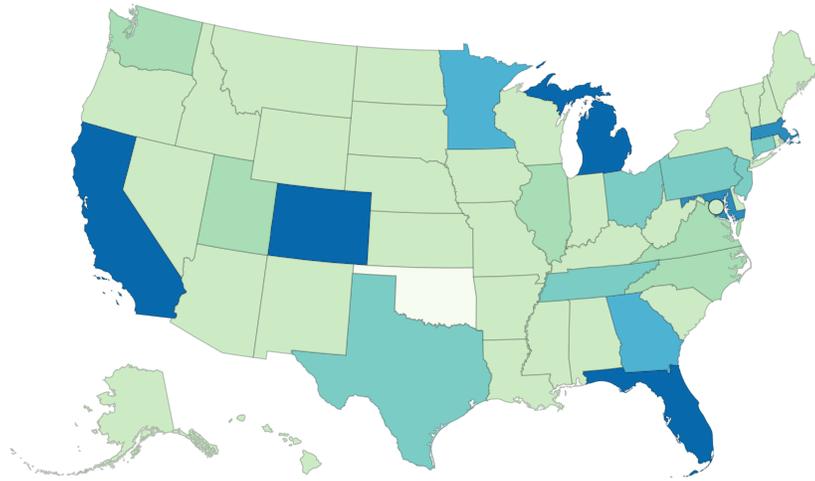
## US COVID-19 Cases Caused by Variants

Updated Apr. 1, 2021

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Variant	Reported Cases in US	Number of Jurisdictions Reporting
B.1.1.7	12,505	51
B.1.351	323	31
P.1	224	22

### Cases of Variants of Concern in the United States\*†



**Number of Cases**

- 0 to 0
- 1 to 150
- 151 to 300
- 301 to 450
- 451 to 600
- 601 to 750
- 751+

**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
Alabama	105	0	1
Alaska	3	5	0
American Samoa	0	0	0
Arizona	95	5	0
Arkansas	7	0	0
California	822	33	10
Colorado	894	0	23
Connecticut	379	2	7
Delaware	17	0	1
Florida	2351	55	16
Georgia	592	1	22
Guam	0	0	0
Hawaii	13	0	5
Idaho	30	0	2
Illinois	218	19	3
Indiana	137	1	2
Iowa	89	0	0
Kansas	27	0	0
Kentucky	64	0	0

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● Kentucky			
● Louisiana	51	0	0
● Maine	15	1	4
○ Marshall Islands	0	0	0
● Maryland	611	1	39
● Massachusetts	712	58	12
● Michigan	1237	0	5
○ Micronesia	0	0	0
● Minnesota	526	2	8
● Mississippi	42	0	1
● Missouri	16	0	0
● Montana	8	0	0
● Nebraska	51	2	0
● Nevada	62	0	1
● New Hampshire	101	0	0
● New Jersey	389	3	1
● New Mexico	21	0	0
● New York	136	0	1
● North Carolina	178	0	29
● North Dakota	7	0	0
○ Northern Marianas	0	0	0
● Ohio	398	10	3
○ Oklahoma	0	1	0
● Oregon	18	1	0
○ Palau	0	0	0
● Pennsylvania	431	0	3
● Puerto Rico	1	0	0
● Rhode Island	7	0	0
● South Carolina	86	0	68
● South Dakota	14	0	1
● Tennessee	318	4	1
● Texas	414	3	3
● Utah	207	3	0
● Vermont	12	0	0

Virgin Islands	0	0	0
Virginia	190	0	30
Washington	223	13	17
Washington D.C.	40	0	2
West Virginia	53	0	0
Wisconsin	78	1	2
Wyoming	9	0	0

[Download Data \(CSV\)](#)
[Download Accessible Data](#)  [CSV – 729 B]

CDC is closely monitoring these [variants of concern \(VOC\)](#). These variants 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).

The map above displays reported data on three VOCs, B.1.1.7, B.1.351, and P.1. Variants B.1.427 and B.1.429 have recently been classified as VOCs but are currently not featured on this map. Additional data is available on [Variant Proportions in the US](#).

03/14/21 – To maintain a limited number of colors and allow for an easier comparison between states, the data ranges have changed from 1-100 to 1-150.

\*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 Apr. 1, 2021

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