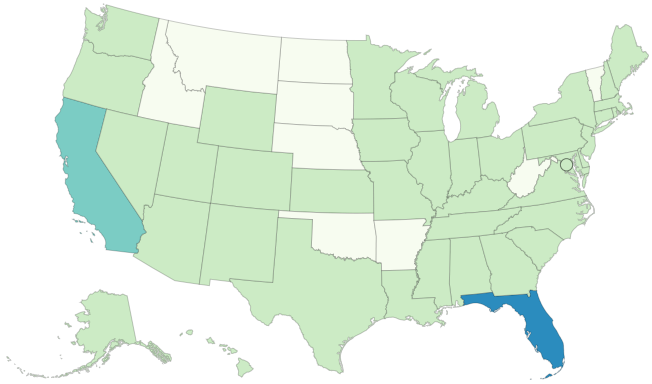
COVID-19 ACT NOW! WEAR A MASK STAY 6 FEET APART AVOID CROWDS

US COVID-19 Cases Caused by Variants

Updated Feb. 16, 2021 Print

Variant	Reported Cases in US	Number of States Reporting
B.1.1.7	1277	42
B.1.351	19	10
P.1	3	2

Emerging Variant Cases in the United States*†



Number of Cases

0 to 0
1 to 100
101 to 200
401 to 500

Filters

Variant B.1.1.7

Territories AS GU MH FM MP PW PR VI

Data Table

Locati...	B.1.1.7 Variant	P.1. Variant	B.1.351 Variant
Alaba...	7		
Alas...	1		

American Sa...			
Arizo...	5		
Arkan...			
Califor...	186		2
Colora...	67		
Connecti...	42		1
Delaw...	3		
Flori...	416		
Geor...	45		
Gua...			
Haw...	2		
Ida...			
Illin...	25		1
India...	13		
Iowa	8		
Kans...	1		
Kentu...	5		
Louisia...	15		
Mai...	2		
Marshall Isl...			
Maryla...	22		7
Massachu...	34		1
Michig...	67		
Microne...			
Minnes...	18	2	
Mississi...	1		
Misso...	1		
Monta...			
Nebra...			
Neva...	6		
New Hamp...	1		
New Jer...	38		
New Me...	2		
New Y...	70		
North Car...	23		1
North Da...			
Northern Mar			

Oh...	1			
Oklaho...		1		
Oreg...	4			
Pal...				
Pennsylv...	26			
Puerto ...				
Rhode Isl...	3			
South Car...	1			2
South Da...				
Tennes...	5			
Tex...	60		1	
Ut...	12			
Verm...				
Virgin Isla...				
Virgi...	11		2	
Washing...	15			
Washington...	3		1	
West Virg...				
Wiscon...	5			
Wyomi...	5			

Download Data (CSV)

Download Accessible Data [XLS – 1 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. 16, 2021
Content source: National Center for Immunization and Respiratory Diseases (NCIRD), Division of Viral Diseases