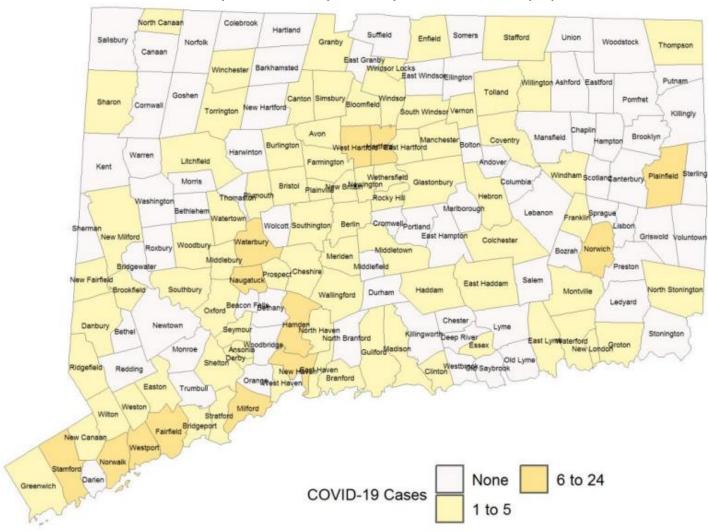


^{*}The significant increase in deaths reported 4/13/21 is due to delayed reporting, **not** a significant increase in death counts over that 7-day period.

Community Transmission of COVID-19

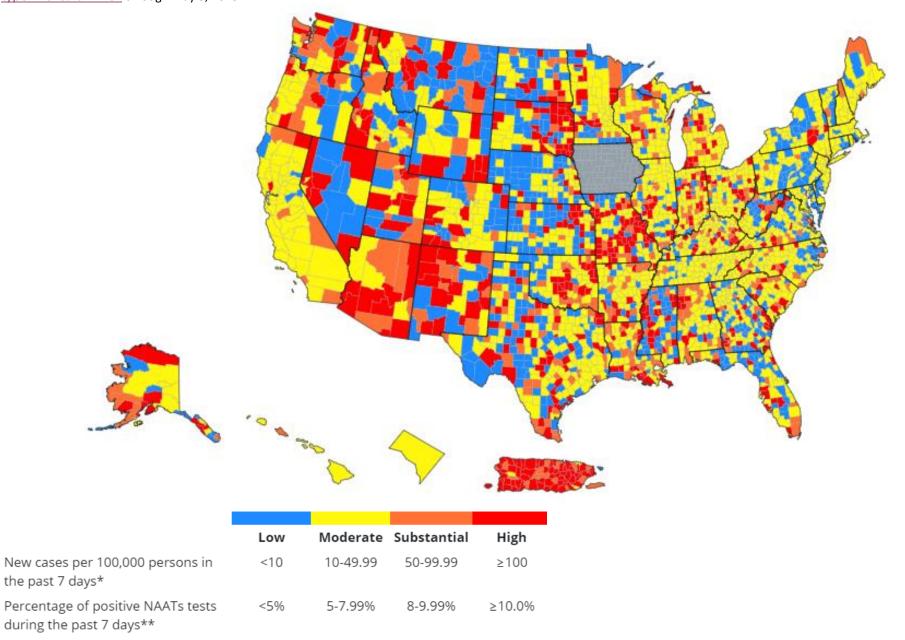
Statewide case rate for April 27 – May 3: 8.0 per 100,000 CT population



Map does not include 0 cases pending address validation

All data are preliminary and subject to change. Updated 5-10-2023.

Data obtained from: https://covid.cdc.gov/covid-data-tracker/#county-view?list_select_state=all_states&list_select_county=all_counties&data-type=Risk&null=Risk through May 3, 2023.



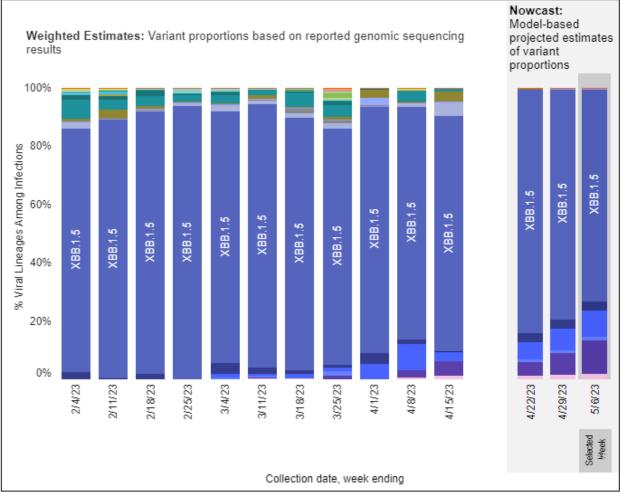
All data are preliminary and subject to change. Updated 5-10-2023.

Weighted and Nowcast Estimates in HHS Region 1 for Weeks of 1/29/2023 – 5/6/2023

Nowcast Estimates in HHS Region 1 for 4/30/2023 – 5/6/2023

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Hover over (or tap in mobile) any lineage of interest to see the amount of uncertainty in that lineage's estimate.



Region 1 - Connecticut, Maine, Massachusetts, New Hampshire, Rhode Island, and Vermont

WHO label	Lineage #	US C	lass %To	tal 95%PI
Omicron	XBB.1.5	VOC	72.9%	60.8-82.5%
	XBB.1.16	VOC	11.4%	4.2-25.4%
	XBB.1.9.1	VOC	9.0%	4.1-17.9%
	XBB.1.5.1	VOC	3.2%	2.3-4.4%
	XBB.2.3	VOC	1.9%	0.6-5.4%
	XBB.1.9.2	VOC	1.4%	0.4-3.9%
	FD.2	VOC	0.1%	0.0-0.4%
	CH.1.1	VOC	0.1%	0.0-0.1%
	BQ.1.1	VOC	0.0%	0.0-0.1%
	XBB	VOC	0.0%	0.0-0.1%
	BQ.1	VOC	0.0%	0.0-0.0%
	BA.2.12.1	VOC	0.0%	0.0-0.2%
	BA.1.1	VOC	0.0%	0.0-0.3%
	BA.2	VOC	0.0%	0.0-0.0%
	BA.2.75.2	VOC	0.0%	0.0-0.0%
	BA.5	VOC	0.0%	0.0-0.0%
	BN.1	VOC	0.0%	0.0-0.0%
	BA.2.75	VOC	0.0%	0.0-0.0%
	BF.7	VOC	0.0%	0.0-0.0%
	BA.5.2.6	VOC	0.0%	0.0-0.0%
	BF.11	VOC	0.0%	0.0-0.0%
Other	Other*		0.0%	0.0-0.0%

^{*} Enumerated lineages are US VOC and lineages circulating above 1% nationally in at least one week period. "Other" represents the aggregation of lineages which are circulating <1% nationally during all weeks displayed.

All data are preliminary and subject to change. Updated 5-10-2023.

[#] BA.1, BA.3 and their sublineages (except BA.1.1 and its sublineages) are aggregated with B.1.1.529. Except BA.2.12.1, BA.2.75, XBB and their sublineages, BA.2 sublineages are aggregated with BA.2. Except BA.2.75.2, CH.1.1 and BN.1, BA.2.75 sublineages are aggregated with BA.2.75. Except BA.4.6, sublineages of BA.4 are aggregated to BA.4. Except BF.7, BF.11, BA.5.2.6, BQ.1 and BQ.1.1, sublineages of BA.5 are aggregated to BA.5. Except the lineages shown and their sublineages, sublineages of XBB are aggregated to XBB. Except XBB.1.5.1 and FD.2, sublineages of XBB.1.5 are aggregated to XBB.1.5. For all the other lineages listed, their sublineages are aggregated to the listed parental lineages respectively. Previously, XBB.2.3 was aggregated to XBB. Lineages BA.2.75.2, XBB, XBB.1.5, XBB.1.5.1, FD.2, XBB.1.9.1, XBB.1.9.2, XBB.1.16, XBB.2.3, BN.1, BA.4.6, BF.7, BF.11, BA.5.2.6 and BQ.1.1 contain the spike substitution R346T.