



Flash survey on SARS-CoV-2 variants in urban wastewater in Italy
20th Report
(Study period: April 3rd to April 7th, 2023)

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Main findings:

- During the week of April 3rd to April 7th, 2023, a total of 133 wastewater samples were collected from 16 Regions and 2 Autonomous Provinces (AP).
- Mutations characteristic of the Omicron variant were identified in 11 of these Regions/AP, while no sequencing data was obtained from the remaining areas.
- Analysis of the sequences obtained by Sanger sequencing showed that 78.9% of the positive samples had amino acid substitutions belonging to the Omicron XBB.1.5*/XBB.1.9* lineages while 18.4% of the positive samples had mutations representative of Omicron CH.1.1* lineage.

Introduction

On March 17th, 2021, the European Union Commission issued Recommendation 2021/472, encouraging Member States to establish a systematic surveillance of SARS-CoV-2 and its variants in wastewater by October 1st, 2021. Responding to this recommendation, the Istituto Superiore di Sanità (ISS) initiated a series of "flash surveys". These surveys involve monthly sampling campaigns carried out over a short period of time at different locations throughout Italy. The primary objective of these flash surveys is to gather supplementary information on SARS-CoV-2 variants in the population to complement the data obtained through clinical surveillance. The aim of this report is to summarize the results from the 20th national flash survey on SARS-CoV-2 variants in wastewater samples in Italy, which was conducted from April 3rd to April 7th, 2023.

Methodology

During the period from April 3rd to April 7th, 2023, the 20th national flash survey on SARS-CoV-2 variants in wastewater samples was conducted in Italy. The survey involved the collection of 133 wastewater samples from 129 wastewater treatment plants (WTPs) located in 16 Regions and 2 Autonomous Provinces. Information on the WTPs participating in the SARS-CoV-2 surveillance in urban wastewater in Italy can be found on the ISS website¹. The samples collected during the survey were processed and the virus concentration was determined by laboratories within the SARI network using the protocol "Sorveglianza di SARS-CoV-2 in reflui urbani - Protocollo progetto SARI - rev.3"². The purified RNAs extracted from the samples were then sent to ISS for variant detection.

A real-time RT-PCR assay was used to screen for the presence of the Omicron variant³. In addition, a long-nested PCR assay of approximately 1600 base pairs was used, covering amino acid residues 58 to 573 of the spike protein. The assay was designed to detect multiple nucleotide changes characteristic of Variants of Concern (VoCs) and Variants of Interest (Vols) in the spike protein⁴. Following amplification of the target DNA sequences, Sanger sequencing was performed on individual samples. For variant classification we adopted a lineage classification

¹ Surveillance of SARS-CoV-2 in urban wastewater in Italy 1° Report (Study period: 01 October 2021 - 31 March 2022) [8e5e2edb-bae0-f1b0-ee6e-08255c76484f \(iss.it\)](https://doi.org/10.5281/zenodo.5758724)

² DOI [10.5281/zenodo.5758724](https://doi.org/10.5281/zenodo.5758724).

³ La Rosa G, Iaconelli M, Veneri C, Mancini P, Bonanno Ferraro G, Brandtner D, Lucentini L, Bonadonna L, Rossi M, Grigioni M; SARI network; Suffredini E. The rapid spread of SARS-CoV-2 Omicron variant in Italy reflected early through wastewater surveillance. Sci Total Environ. 2022 Sep 1;837:155767. doi: 10.1016/j.scitotenv.2022.155767. Epub 2022 May 6. PMID: 35533857; PMCID: PMC9074219.

⁴ G La Rosa, P. Mancini, G. Bonanno Ferraro, C. Veneri, M. Iaconelli, L. Lucentini, L. Bonadonna, S. Brusaferro, D. Brandtner, A. Fasanella, L. Pace, A. Parisi, D. Galante, E. Suffredini. Rapid screening for SARS-CoV-2 variants of concern in clinical and environmental samples using nested RT-PCR assays targeting key mutations of the spike protein, Water Research, 2021, Volume 197, 1 June 2021, 117104. <https://doi.org/10.1016/j.watres.2021.117104>.

based on 'outbreak.info'⁵ rather than specifying sublineages. This choice was made because there are many sublineages that evolve rapidly, often converging on specific amino acid substitutions. In some cases, the differences between sublineages can be as small as a single nucleotide mutation in our target region, making a reliable assignation to sublineages, based solely on the mutations observed in the spike region, not feasible.

Results

Real Time qPCR

Out of the 133 samples collected, a total of 127 (95.5%) tested positive for SARS-CoV-2 using the real-time RT-qPCR method employed for environmental surveillance (Table 1). The viral concentrations detected in these samples varied, ranging from 1.84E+02 to 2.24E+05 genome copies (g.c.) per liter of wastewater. Additionally, 125 out of the 133 samples (94.0%) were found to be positive for the omicron variant using the RT-qPCR assay, with cycle threshold (CT) values ranging from 32.0 to 39.5.

Sanger Sequencing

Table 1 provides a summary of the results obtained from the real-time PCR assays, long nested PCR, and sequencing methods. A total of 38 samples (28.6%) from 11 Regions/AP were successfully amplified using the long-nested PCR assay. Sanger sequencing confirmed that all sequences corresponded to the omicron variant.

Analysis of the wastewater samples revealed the presence of two SARS-CoV-2 lineages, as shown in Tables 1 and 2. Among these lineages, the XBB.1.5*/XBB.1.9* lineages (which were not distinguishable in the sequenced region), were dominant, detected in 78.9% of the positive samples (30 samples). In addition, the CH.1.1* lineage of the omicron variant was identified in 18.4% of the samples (7 samples). One sample could not be assigned due to partial sequence. For ease of understanding, the mutations have been grouped into two panels or 'mutation packages' as follows:

- **Package A (assigned to the lineages Omicron XBB.1.5*/XBB.1.9*)** = V83A, G142D, DEL144, H146Q/H146K, Q183E, V213E, G252V, G339H, R346T, L368I, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, V445P, G446S, N460K, S477N, T478K, E484A, F486P, F490S, Q498R, N501Y, Y505H
- **Package B (assigned to the lineage Omicron CH.1.1*)** = G142D, K147E, W152R, F157L, I210V, V213G, G257S, G339H, R346T, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, K444T, G446S, L452R, N460K, S477N, T478K, E484A, F486S, Q498R, N501Y, Y505H

'Package A' combined with the 'F456L' mutation, suggests the presence of the Omicron EG.5 sublineage. This sublineage is characterized by having the 'F456L' mutation in approximately 80.5% of its sequences (<https://outbreak.info/situation-reports>, date: 05/10/2023).

Package A was detected in nine Regions (Abruzzo, Emilia-Romagna, Friuli-Venezia Giulia, Liguria, Lombardia, Molise, Puglia, Sicilia, Veneto) and two A.P. (Bolzano and Trento). Characteristic amino acid substitutions of package B (lineage CH.1.1*) were observed in Puglia and Sicilia.

⁵ <https://outbreak.info/situation-reports>, date: 05/10/2023

Table 1. PCR and sequencing results

Sample ID	Region/A.P.	City	WTP	RT-qPCR (c.g./L)	RT-PCR Omicron-ID 999 (CT value)	mutations found by Sanger sequencing (long PCR ID_980)	SARS-CoV-2 lineages
108	Abruzzo	Pescara	Villa Carmine	1,95E+02	36.30		
109		Pescara	Via Raiale	1,84E+02	38.37		
110		Chieti	S. Martino	<LOD	38.06		
111		L'Aquila	Pile	6,34E+02	34.47	Package A	Omicron XBB.1.5*/XBB.1.9*
112		Teramo	Villa Pavone	5,68E+02	36.13		
1	Basilicata	Potenza	Tiera di Vaglio	1,16E+04	38.90		
2		Matera	Pantano	7,39E+03	38.10		
3	Campania	Napoli	Napoli EST	3,09E+03	36.05		
4		Napoli	Napoli OVEST - Ingresso Principale	5,08E+04	37.19		
5		Napoli	Napoli OVEST - ex ingresso Camaldoli	1,72E+04	39.14		
6		Ferrara	Ferrara - Linea 1	2,93E+04	ND		
7		Ferrara	Ferrara - Linea 2	9,78E+03	ND		
8	Emilia- Romagna	Modena	Carpi	2,11E+04	37.11		
9		Piacenza	Borgoforte	3,73E+04	ND		
10		Parma	Parma ovest	8,13E+03	37.02		
11		Reggio Emilia	Mancasale	3,85E+04	33.46	Package A + F456L ^a	Omicron XBB.1.5*/XBB.1.9*
117		Bologna	IDAR	1,21E+05	35.06	Package A	Omicron XBB.1.5*/XBB.1.9*
118		Ravenna - Forlì-Cesena	Ravenna	1,10E+05	34.36		
119		Modena	Naviglio	2,93E+04	36.43		
120		Ravenna	Faenza	3,06E+04	36.09		
121		Bologna	Imola	3,76E+04	35.37	Package A	Omicron XBB.1.5*/XBB.1.9*
124		Forlì-Cesena	Forlì	5,44E+04	34.18	Package A	Omicron XBB.1.5*/XBB.1.9*

125	19952		Forlì-Cesena	Cesena	2,24E+05	37.01	Package A	Omicron XBB.1.5*/XBB.1.9*
127	20039		Rimini - Forlì-Cesena	S. Giustina	2,81E+04	35.54	Package A	Omicron XBB.1.5*/XBB.1.9*
122	19909	Friuli-Venezia Giulia	Pordenone	Cordenons	1,74E+05	35.89	Package A	Omicron XBB.1.5*/XBB.1.9*
123	19910		Udine	Udine	1,26E+05	33.02		
126	20046		Trieste	Servola	3,60E+04	33.67	Package A	Omicron XBB.1.5*/XBB.1.9*
12	19620		Viterbo	Viterbo - Strada Bagni	9,40E+03	35.72		
13	19621		Roma	Anzio - Colle Cocco	1,66E+03	33.79		
14	19622		Latina	Aprilia (Via del Campo)	7,38E+03	37.26		
15	19623		Latina	Latina Loc Latina Est	2,14E+03	36.72		
16	19625		Roma	Pomezia - Via Cincinnato	<LOD	35.90		
17	19626	Lazio	Roma	Velletri (LA CHIUSA-SORBO)	2,31E+03	38.01		
18	19627		Roma	Guidonia - Ponte Lucano	1,10E+04	33.92		
19	19689		Roma	Civitavecchia Fiumaretta	<LOD	36.61		
103	19882		Roma	Roma Est (linea 1 + linea 2)	2,10E+04	36.96		
104	19883		Roma	Roma Nord	1,61E+04	35.58		
105	19884		Roma	Roma Sud	1,05E+04	36.71		
106	19885		Roma	Ostia	2,47E+04	37.16		
107	19886		Roma	Fregene	4,42E+04	38.76		
20	19766		La Spezia	Camisano	2,21E+04	ND		
21	19767		La Spezia	Silea	2,69E+04	34.29		
22	19768		La Spezia	La Spezia	5,68E+04	33.08		
23	19770	Liguria	Genova	Pegli	3,81E+04	34.02		
24	19771		Genova	Voltri	2,10E+04	33.06		
25	19772		Genova	Quinto	2,19E+04	35.01		
26	19773		Genova	Rapallo	3,94E+04	34.78		
27	19774		Genova	Sestri P	1,24E+04	32.85		
28	19775		Genova	Sturla	1,71E+04	35.16		

29	19776		Imperia	Imperia	1,54E+04	34.67		
30	19777		Imperia	Sanremo - località Capo Verde	1,57E+04	34.34		
31	19778		Genova	Darsena	2,16E+04	34.53	Package A	Omicron XBB.1.5*/XBB.1.9*
32	19779		Genova	Punta Vagno Genova	8,37E+03	33.50		
33	19780		Genova	Valpolcevera	1,96E+04	34.68		
34	19781		Savona	Borghetto Santo Spirito	7,74E+03	36.99		
35	19851		Genova	Punta Vagno Genova	3,70E+04	37.30	Package A	Omicron XBB.1.5*/XBB.1.9*
116	19769		Savona	Savona	2,34E+04	36.18	Package A	Omicron XBB.1.5*/XBB.1.9*
36	19745		Sondrio	Sondrio	1,35E+04	38.32		
37	19749		Milano	Milano Nosedo	1,12E+05	36.15		
38	19750		Milano	Milano San Rocco	1,20E+05	35.53		
39	19751		Como	Como	8,99E+04	ND	Package A	Omicron XBB.1.5*/XBB.1.9*
40	19752		Pavia	Pavia	7,75E+04	37.40		
41	19753	Lombardia	Como - Lecco - Milano - Monza e della Brianza	Monza	5,90E+04	38.34		
42	19754		Pavia	Vigevano	5,26E+04	36.20		
43	19783		Bergamo	Bergamo	9,16E+03	35.44		
44	19785		Cremona	Citta di Cremona	1,88E+04	36.12		
45	19786		Brescia	Verziano	8,40E+03	36.93		
46	19823		Pesaro- Urbino	Borgheria	3,23E+03	37.40		
47	19824		Pesaro- Urbino	Ponte Metauro	1,45E+03	37.51		
48	19825	Marche	Pesaro- Urbino	Ponte Sasso	7,02E+03	33.48		
49	19826		Ancona	Zipa	3,00E+04	35.02		
50	19827		Ancona	Falconara	1,11E+04	37.21		
51	19828		Ancona	Camerano	5,51E+03	33.69		
113	19974	Molise	Campobasso	Campobasso - San Pietro	4,47E+02	37.11		

114	19975	Campobasso	Termoli - località Porto	<LOD	36.62		
115	19976	Campobasso	Termoli - località Pantano Basso	1,97E+02	35.57	Package A	Omicron XBB.1.5*/XBB.1.9*
131	19876	Bolzano	IDA Bolzano	3,66E+04	35.50	Partial sequence, from G339H to Y505H ^b	Not assigned ^b
132	19877	P.A. Bolzano	Bolzano	IDA Merano	6,71E+04	34.01	
133	19878		Bolzano	IDA Termeno	9,20E+04	33.15	Package A
52	19690		Trento	Trento nord	1,05E+05	35.42	Package A
53	19691	P.A. Trento	Trento	Trento sud	6,76E+04	33.65	
54	19692		Trento	Rovereto	1,26E+05	38.14	Package A
55	19650		Torino	Castiglione Torinese	9,15E+02	ND	
56	19651	Piemonte	Biella	Biella Nord	4,00E+02	38.75	
57	19652		Biella	Biella Sud	2,23E+03	37.68	
58	19653		Novara	Novara	<LOD	36.24	
59	19733		Alessandria	Alessandria	1,11E+04	34.70	
60	19734		Asti	Asti	1,23E+03	38.14	
61	19735		Cuneo	Cuneo	1,00E+03	36.80	
62	19618		Bari	Bari Ovest	1,17E+03	35.13	Package B
63	19619		Bari	Bari Est	7,80E+02	34.14	Package B
64	19624		Bari	Molfetta	6,06E+02	36.14	
65	19632	Puglia	Bari	Altamura	7,29E+02	36.47	Package A
66	19634		Brindisi	Brindisi Fiume Grande	3,27E+03	35.36	Package B
67	19635		Lecce	Lecce	1,81E+04	33.00	Package A
68	19637		Taranto	Taranto Bellavista	2,76E+03	35.72	Package B
69	19638		Taranto	Taranto Gennarini	9,51E+02	37.55	
70	19703		Barletta-Andria-Trani	Andria	7,92E+03	35.63	Package B
71	19704		Barletta-Andria-Trani	Barletta	1,30E+04	35.32	
72	19705		Bari	Bitonto	1,55E+04	34.82	Package A
							Omicron XBB.1.5*/XBB.1.9*

73	19706		Foggia	Cerignola	1,44E+04	35.25	Package A	Omicron XBB.1.5*/XBB.1.9*
74	19707		Foggia	Foggia	1,50E+04	35.02	Package A	Omicron XBB.1.5*/XBB.1.9*
75	19708		Foggia	Manfredonia	6,17E+03	ND		
98	19728		Barletta-Andria-Trani	Bisceglie	3,63E+03	37.14	Package A	Omicron XBB.1.5*/XBB.1.9*
99	19729		Barletta-Andria-Trani	Trani	1,58E+03	35.10		
76	19788		Ragusa	Modica	5,03E+02	37.54		
79	19791		Caltanissetta	Gela Macchitella	1,53E+03	ND		
80	19792		Messina	Mili Marina	2,25E+02	38.21		
81	19793		Messina	Mili Marina	1,06E+03	34.88		
82	19839		Agrigento	Agrigento	2,41E+04	37.43	Package B	Omicron CH.1.1*
83	19841		Enna	Enna	4,88E+03	36.36		
84	19842		Palermo	Bagheria	1,17E+04	37.94	Package A	Omicron XBB.1.5*/XBB.1.9*
85	19843	Sicilia	Palermo	Acqua dei Corsari	7,20E+03	37.08	Package B	Omicron CH.1.1*
86	19844		Palermo	Fondo Verde	3,20E+03	35.09		
87	19846		Caltanissetta	Caltanissetta e San Cataldo	1,35E+04	39.53	Package A	Omicron XBB.1.5*/XBB.1.9*
100	19700		Trapani	Mazara del Vallo	6,34E+03	37.11		
101	19701		Trapani	Marsala	3,89E+03	39.30		
102	19702		Trapani	Trapani	1,05E+04	39.02		
128	19763		Catania	Pantano d'Arci	2,10E+04	36.77		
129	19764		Catania	Giarre	1,80E+04	37.43		
130	19765		Siracusa	Siracusa	2,54E+04	39.03		
135	19697	Umbria	Perugia	Perugia - Pian della Genna	6,95E+04	34.49		
134	20007	Val d'Aosta	Aosta	Brissogne	4,25E+02	37.02		
88	19673		Padova	Padova Ca' Nordio - centro storico	8,33E+04	35.39	Package A	Omicron XBB.1.5*/XBB.1.9*
89	19674	Veneto	Padova	Padova Ca' Nordio - zip	8,95E+02	37.48		
90	19675		Padova	Padova Guizza	1,44E+05	33.58	Package A	Omicron XBB.1.5*/XBB.1.9*
91	19676		Padova	Abano Terme	1,01E+05	37.22		

92	19718	Treviso	Treviso	1,22E+04	33.74		
94	19720	Venezia	Venezia Fusina	7,53E+03	33.18	Package A	Omicron XBB.1.5*/XBB.1.9*
93	19719	Vicenza	Vicenza Casale	<LOD	34.89		
95	19756	Verona	Verona_collettore 1M	1,40E+04	32.36		
96	19757	Verona	Verona_collettore 3M	4,45E+04	32.01	Package A	Omicron XBB.1.5*/XBB.1.9*
97	19758	Verona	Verona_collettore 8M	2,68E+04	35.56	Package A	Omicron XBB.1.5*/XBB.1.9*

^a 'Package A' combined with the '**F456L**' mutation, suggests the presence of the Omicron EG.5 sublineage. This sublineage is characterized by having the 'F456L' mutation in approximately 80.5% of its sequences. The Omicron XBB.1.5* and Omicron XBB.1.9* lineages may also carry the F456L mutation, but at lower rates of 3.0% and 26.6%, respectively (<https://outbreak.info/situation-reports>, date: 05/10/2023);

^b Partial sequence due to mixed electropherograms and/or high signal noise; within brackets the region for which a sequence was provided;

ND: Not Detected

Table 2. Sanger sequencing results

Limitations of the study

The geographic and population coverage of this flash survey was not comprehensive, as it included 18 out of 21 of the Italian Regions/Autonomous Provinces.

It is important to note that the molecular analytical methods used for complex environmental matrices, such as wastewater, can be challenged by factors such as low viral concentrations, insufficient analyte recovery, and/or inhibition of PCR amplification. As a result, both the detection/quantification and PCR amplification for sequencing may yield false negatives, making it difficult to achieve molecular characterization and variant detection for all samples.

Partial sequencing of the Spike region does not provide conclusive results for sublineage assignment. Our decision to adopt a broader lineage classification from 'outbreak.info' for variant classification, rather than specifying sublineage assignments, was influenced by the rapid evolution of numerous sublineages, often with minor differences, that hampered the reliable assignation to sublineages based solely on mutations observed in the spike region.

Conclusions and final considerations

This report is part of a monthly series focusing on SARS-CoV-2 and its variants in wastewater samples in Italy, in accordance with the EU Commission Recommendation 2021/472. The primary objective is to provide supplementary information on SARS-CoV-2 variants in the population, complementing data obtained through clinical surveillance. The results of this survey indicate that the Omicron variant is the only SARS-CoV-2 variant in Italy, with the XBB.1.5*/XBB.1.9* lineages being the most common. However, mutations characteristic of the Omicron CH.1.1* lineage have also been detected.

The sequencing SARS-CoV-2 in wastewater samples provides valuable additional information alongside the sequencing of clinical cases. This approach provides a more comprehensive and complete understanding of the circulating variants in the country, contributing to a better characterization of the spread and evolution of the virus.

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