

Virus detail	
Virus name:	hRSV/A/Spain/CN-HUNSC_ITER_150388052/2023
Accession ID:	EPI_ISL_18321533
Subtype:	A
AA Substitutions	F A103T, F T122A, G A57V, G G224E, G H90Y, G I134K, G I265L, G K209R, G K262E, G L101F, G L248I, G L310P, G N103S, G P71L, G P206Q, G S105P, G S243I, G T319I, G T320A, G V303A, G Y304H, L A2014del, L A2026del, L A2042I, L A2070L, L A2088I, L C1995G, L C2059D, L D1978A, L D1981L, L D1985N, L D2043Q, L D2048P, L D2081E, L E1725G, L E2045N, L E2092Q, L E2120del, L E2130L, L E2138M, L E2147L, L F1982L, L F2017del, L F2020del, L F2057W, L F2094K, L F2110L, L G1731D, L G2012del, L G2066S, L G2089F, L H2101N, L H2103Y, L H2125N, L H2135N, L I1980Y, L I1987L, L I1989L, L I2011del, L I2029del, L I2037del, L I2047V, L I2055L, L I2062V, L I2082S, L I2087L, L I2099C, L ins1990Y, L ins2047YY, L ins2055RVG, L ins2058NTILD, L ins2091VWLYNQIAL, L ins2102L, L K1983A, L K2027del, L K2034M, L K2041T, L K2044H, L K2064L, L K2065E, L K2073E, L K2075E, L K2097A, L L1438Q, L L1996I, L L2007del, L L2009del, L L2028del, L L2030I, L L2054V, L L2058I, L L2071T, L L2114K, L L2121F, L L2126I, L L2140F, L L2143G, L M2104L, L M2128T, L N1723G, L N1986S, L N2015del, L N2021del, L N2025del, L N2035Q, L N2050S, L N2068G, L N2096H, L N2105D, L N2111K, L N2115T, L N2124D, L N2141G, L P171L, L P2013del, L P2018del, L P2039S, L P2061K, L P2134M, L Q2024del, L R256K, L R2117del, L S1998H, L S2003T, L S2046G, L S2053K, L S2072Q, L S2076del, L S2079R, L S2084L, L S2095N, L S2118del, L S2131T, L S2137P, L S2142G, L T1988K, L T1992E, L T2010del, L T2033D, L T2063S, L T2069S, L T2119del, L T2132L, L T2144D, L T2145P, L V1994A, L V2005T, L V2008del, L V2016del, L V2019del, L V2022del, L V2023del, L V2077del, L V2078Y, L V2093L, L V2113L, L V2129A, L W2109V, L Y598H, L Y2060F, L Y2085C, L Y2123L, L Y2127D, M2-1 S176P, M2-2 S44N, M M73L, N V352A, NS2 D100G, P L55P, P T69I, SH F55L
Passage details/history:	Original
Sample information	
Collection date:	2023-02-14
Location:	Europe / Spain / Canary Islands
Host:	Human
Additional location information:	
Gender:	Male
Patient age:	75
Patient status:	Hospitalized
Specimen source:	Nasopharyngeal swab
Additional host information:	
Sampling strategy:	Non-sentinel-surveillance (hospital)
Outbreak:	
Last vaccinated:	
Treatment:	
Sequencing technology:	Illumina MiSeq
Assembly method:	Ampliseq-custom pipeline
Coverage:	960x
Comment:	Gap of 4 nucleotides when compared to the reference sequence. %UniqueMutations 3.18%.
Institute information	
Originating lab:	Servicio de Microbiología Complejo Hospitalario Universitario Nuestra Señora de Candelaria
Address:	Servicio de Microbiología Complejo Hospitalario Universitario Nuestra Señora de Candelaria Ctra. Gral. del Rosario 38010 Santa Cruz de Tenerife Spain
Sample ID given by the originating laboratory:	HUNSC_ITER_150388052
Submitting lab:	Institute of Technology and Renewable Energy (ITER) Genomics Division
Address:	Institute of Technology and Renewable Energy (ITER) Genomics Division Poligono Industrial de Granadilla, S/n, (parque Eolico) 38600 Granadilla de Abona Spain
Sample ID given by the submitting laboratory:	hRSV/A/CN-Spain/HUNSC_ITER_150388052/2023
Authors:	Julia,Alcoba-Florez;Rafaela,González-Montelongo;Diego,García-Martínez de Artola;Adrián,Muñoz-Barrera;Helena,Gil-Campesino;Oscar,Diez-Gil;Jose Miguel,Lorenzo-Salazar;Carlos,Flores
Submitter information	
Submitter:	Lorenzo Salazar, Jose Miguel
Address:	Poligono Industrial De Granadilla, S/n, (parque Eolico) 38600 Granadilla De Abona Spain
Submission Date:	2023-09-28

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