Variant of Concern (VOC) and Variant of Interest (VOI) cases confirmed by the whole genome sequencing at the NIID* and other institutions

*The National Institute of Infectious Diseases

As of September 13^{th} , 2021 ()=Increase from September 6^{th} , 2021

Lineage	Domestic cases	Border-screening cases
B.1.1.7	46,734	337
(Alpha)	(+775)	(+0)
B.1.351	24	92
(Beta)	(+0)	(+0)
P.1	95	29
(Gamma)	(+1)	(+0)
B.1.617.2	27,820	834
(Delta*)	(+6,919)	(+65)
B.1.617.1	8	19
(Kappa)	(+0)	(+0)

(Provisional figures)

^{*}Includes strains with equivalent mutations (e.g.AY.1).