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

 $\mbox{As of October 4^{th} , 2021} \label{eq:association} \mbox{()=Increase from September 27^{th} , 2021}$

Lineage	Domestic cases	Border-screening cases
B.1.1.7	48,151	339
(Alpha)	(+295)	(+0)
B.1.351	24	92
(Beta)	(+0)	(+0)
P.1	105	29
(Gamma)	(+1)	(+0)
B.1.617.2	50,426	1,002
(Delta*)	(+7,705)	(+62)
B.1.617.1	8	19
(Kappa)	(+0)	(+0)

(Provisional figures)

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