

Virus	Clade	Tier	IC ₅₀ (nM)							
			dsDNA	PG9-60bp-3BNC60	PG9-3BNC60 heterodimer	Mix of PG9, 3BNC60, dsDNA	PG9 Fab	PG9 IgG	3BNC60 Fab	3BNC60 IgG
6535.3	B	1B	>1000	<0.03 (>767x)		23	18	4.0	27	4.0
SC422661.8	B	2	>1000	<0.03 (>47x)		1.4	14	6.9	1.0	0.3
PVO.4	B	3	>1000	<0.03 (>40x)	40	1.2	59	57	1.3	0.1
TRO.11	B	2	>1000	<0.03 (>40x)	2.6	1.2	54	71	1.3	0.2
TRJO4551.58	B	3	>1000	<0.03 (>187x)		5.6	32	7.6	4.8	<0.1
CAAN5342.A2	B	2	>1000	0.07 (229x)	>10	16	25	26	6.2	3.6
THRO4156.18	B	2	>1000	0.21 (152x)	>10	32	<0.1	110	23	3.5
RHPA4259.7	B	2	>1000	<0.03 (>20x)		0.6	0.2	<0.1	1.5	0.3
Du156.12	C	2	>1000	<0.03 (>53x)		1.6	0.6	0.3	9.4	0.3
Du172.17	C	2	>1000	<0.03 (>273x)		8.2	2.2	3.1	210	16
Du422.1	C	2	>1000	<0.03 (>160x)	9.3	4.8	0.6	1.8	>200	>330
ZM197M.PB7	C	1B	>1000	0.07 (143x)		10	5.4	5.1	15	3.8
ZM214M.PL15	C	1B	>1000	0.21 (39x)	16	8.2	>200	>330	6.9	1.7
ZM233M.PB6	C	2	>1000	<0.03 (>3x)		<0.1	<0.1	<0.1	14	1.4
ZM249M.PL1	C	2	>1000	<0.03 (>27x)		0.8	0.2	0.3	2.7	0.3
ZM53M.PB12	C	2	>1000	<0.03 (>33x)	0.63	1.0	0.4	0.5	5.2	1.3
ZM109F.PB4	C	1B	>1000	<0.03 (>280x)		8.4	6.8	2.1	39	0.3
ZM135M.PL10a	C	2	>1000	<0.03 (>140x)		4.2	160	>330	3.7	0.4
CAP45.2.00.G3	C	2	>1000	<0.03 (>3x)		<0.1	<0.1	<0.1	>200	50
CAP210.2.00.E8	C	2	>1000	<0.03 (>140x)		4.2	1.6	2.0	>200	59
Q842.d12	A	2	>1000	<0.03 (>7x)		0.2	<0.1	<0.1	0.21	0.09
Q259.d2.17	A	2	>1000	<0.03 (>33x)		1.0	0.4	0.2	3.7	0.03
3718.v3.c11	A	2	>1000	<0.03 (>27x)		0.8	<0.1	<0.1	>200	>330
0330.v4.c3	A	2	>1000	<0.03 (>7x)		0.2	<0.1	<0.1	0.62	0.2
3415.v1.c1	A	2	>1000	<0.03 (>67x)		2.0	1.0	1.2	4.6	0.9
Geometric Mean IC₅₀ (nM)				0.038						
Arithmetic Mean IC₅₀ (nM)				0.049						
Mean Fold Improvement Over Mixture (Geometric)				68						
Mean Fold Improvement Over Mixture (Arithmetic)				121						