

D_27.4	I	4632	0.12	2759	ND	0.13	ND	-M-----D-----Q-----R--T-----	MN--DNM--Q--K-----YT-----S-----
D_27.7	I	5419	0.56	2350	ND	0.17	ND	-M-----D-----Q-----R--T-----	MN--DNM--Q--K--SK---YT-----S-----
D_27.9	I	1570	0.18	466	ND	0.11	ND	-M-----DS-----Q-----R--T-----	MN--DNM--Q--K-----YT-----S-----
D_27.10	NI	ND	ND	ND	ND	ND	ND	-M-----DS-----Q-----R--T-----	MN--DNM--Q--K--K---YT-----S-----
D_59.1	NI	ND	ND	ND	ND	ND	ND	-M-----D-----Q-----R--T-----	MN--DNMA--Q--K--SK---YT--A-S-----
D_59.3	I	7750	0.38	2310	ND	0.25	ND	-M-----D-----Q-----R--T-----	MN--DNM--Q--K--SK---YT--A-S-----
D_59.4	I	2414	0.37	6572	ND	0.18	ND	-M-----D-----Q-----R--T-----	MN--DNM--Q--K--SK---YT--A-S-----
D_59.5	I	2566	0.44	4446	ND	0.28	ND	-M-----D-----Q-----R--T-----	MN--DNM--Q--K--SK---YT---SR-----
D_59.6	I	5682	0.47	2161	ND	0.03	ND	-M-----D-----Q-----R--T-----	MN--DNM--Q--E--SK---YT--A-S-----
D_94.3	I	7022	0.81	2587	ND	0.42	ND	-M-----D-----Q-----R--T-----	MN--DNM--Q--K--SK-S--YT--A-S-----
D_94.4	NI	ND	ND	ND	ND	ND	ND	-M-----D-----P-----Q-----R--T-----	MN--DNM--Q--K--SK---YT--A-S--D-----
D_94.5	NI	ND	ND	ND	ND	ND	ND	-M-----D-----Q-----R--T-----	MN--DNM--Q--K--SK---YT--A-S--D-----
D_94.6	NI	ND	ND	ND	ND	ND	ND	-M-----D-----Q-----R--T-----	MN--DNM--Q--K--SK-S--YT--A-S-----
D_94.7	NI	ND	ND	ND	ND	ND	ND	-M-----D-----Q-----R--T-----	MN--DNM--Q--K--SK-S--YT--A-S-----
E_-129.1	I	0.63	1.79	ND	8.52	ND	0.03	-----I-----V-----T-----	DE--KNM--Q-----DI-YN-----
E_-129.2	NI	ND	ND	ND	ND	ND	ND	-----I-----V-----T-----	DE--KNM--Q-----DI-YN-----
E_-129.3	NI	ND	ND	ND	ND	ND	ND	-----I-----V-----T-----	DE--KNM--Q--G-----DI-YN-----
E_-129.4	I	1.02	0.93	ND	5.09	ND	0.03	-----I-----V-----T-----	DE--KNM--Q-----DI-YN-----
E_-129.5	I	0.47	1.20	ND	3.97	ND	0.05	-----I-----V-----T-----	DE--KNM--Q-----DI-YN-----
E_42.1	I	683	1.66	ND	1338	ND	0.06	-----D-----I-----V-----T-----	QNE--NM--Q-----GI-YN--KA-----
E_42.2	NI	ND	ND	ND	ND	ND	ND	-----D-----I-----V-----T-----	QDE--GNM--Q-----GI-YN--KA-----
E_42.4	NI	ND	ND	ND	ND	ND	ND	-----D-----I-----V-----T-----	QNE--NM--Q-----GI-YN--KA-----
E_42.6	I	1048	1.86	5836	2924	0.60	0.15	-----D-----I-----V-----T-----	NE--ENM--Q--K-----DI-YN--KA-----
E_42.7	I	1045	0.49	4518	ND	0.31	ND	-----D-----I-----V-----T-----	QDE--NM--Q-----GI-YN--KA-----

^a Clone identifiers are composed of a letter designating the patient (A-E) followed by a number corresponding to the week of sampling relative to enfuvirtide treatment beginning at week 0 (first number preceding the dot) followed by the clone number

^b infectious

^c non infectious

^d Fold Change in 50% Inhibitory Concentration

^e Relative Infectivity

^f HR1 and HR2 domains are underlined and the region between AA 36 to 45, as well as position 138 in HR2 are highlighted in bold letters.

^g not determined