

CidB positions (aa)

CidB_IV variants	2 2	2 2	2 2	2 2	2	2 2	2 2	2 2	4 4	4	4 4	4	4 4	4 4
	4 4	5 5	5 5	5 5	6	6 - 7	7	7	5 5	5	6 - 6	7	7 7	7 - 8
	2 3	0 1	3 4	7 9	2	8 0	6	8	2 3	9	2 6	2	5 6	9 1

(a/1)	DR	RI	RE	P	I	I	QLV	N N	EY	G	RVRVR	Y	GD	RHT
(a/2)	DR	RI	RE	P	I	I	QLV	N N	EY	*	EAPIT	H	EG	**C
(a/3)	DR	RI	RE	P	I	I	QLV	N N	HP	*	EAPIT	H	EG	**C
(b/1)	NQ	SL	GD	E	V	T	AQI	G S	EY	G	RVRVR	Y	GD	RHT
(b/2)	NQ	SL	GD	E	V	T	AQI	G S	EY	*	EAPIT	H	EG	**C
(b/3)	NQ	SL	GD	E	V	T	AQI	G S	HP	*	EAPIT	H	EG	**C
(c/1)†	NQ	SL	GD	E	V	T	AQI	N N	EY	G	RVRVR	Y	GD	RHT
(c/3)†	NQ	SL	GD	E	V	T	AQI	N N	HP	*	EAPIT	H	EG	**C

Upstream polymorphic region

Downstream polymorphic region