

Table S6. Plasmids of *Mycobacterium abscessus* clinical isolates

Name <sup>1</sup>	Cluster <sup>2</sup>	Length (bp) <sup>3</sup>	ORFs <sup>4</sup>	Copy # <sup>5</sup>	Comments
pGD08	pA	9,547	11	2.9	
pGD42-2	pA	9,547	11	2.6	2 bp different from pGD08
pGD18, pGD62-1, pGD69-1, pGD95-1, pGD108A, pGD108B	pB	25,000	37	1.9	
pGD23	pB	25,002	37	1.8	2 bp insertion vs. pGD18
pGD36-1, pGD47	pB	24,995	37	2.7	
pGD42-1	pB	24,993	38	3.0	
pGD72	pB	24,985	37	1.1	
pGD87	pB	24,994	37	3.0	
pGD22-2, pGD24, pGD34, pGD75, pGD100A, pGD100B	pC	18,117	16	1.9	
pGD39	pC	18,117	16	1.5	
pGD62-2	pC	18,612	17	2.5	
pGD69-2, pGD95-2	pC	18,611	17	1.9	
pGD19	pD	18,605	20	3.1	
pGD45-2	pD	19,406	21	3.1	
pGD85	pD	23,374	26	2.4	
pGD33	pE	25,996	33	1.8	
pGD36-2	pE	24,259	34	5.0	
pGD02	pF	30,963	36	2.2	
pGD25-1, pGD54, pGD102-1	pF	31,413	32	2.5	
pGD86-1	pF	31,343	32	3.1	
pGD25-2	pG	27,424	36	2.5	
pGD45-1	pG	27,427	36	3.9	
pGD86-2	pG	27,424	36	2.3	
pGD102-2	pG	27,425	36	3.9	
pGD58	pH	92,821	122	1.2	
pGD16	pH	Multicontig (C15, 24, 41)			
pGD10	pSin	Multicontig (C16)			
pGD13	pSin	21,881	29	1.7	
pGD21-1	pSin	112,633	150	1.1	
pGD21-2	pSin	155,609	233	1.3	Linear
pGD22-1	pSin	19,694	21	1.6	
pGD25-3	pSin	23,599	26	5.2	
pGD51	pSin	23,656	27	3.6	
pGD52	pSin	22,216	20	3.2	
pGD55	pSin	Multicontig (C15)			
pGD104	pSin	96,413	144	1.3	
pATCC19977 <sup>6</sup>	pSin	23,319	29		

<sup>1</sup>Plasmids are named according to the strains in which they were identified (e.g. pGD22). If there is more than one plasmid in a particular strain, '-1' or '-2', etc. are added. Plasmids identified in different strains, but with identical sequences are shown in the same row.

<sup>2</sup>Cluster designation (pA, pB, etc.) is indicated. Singleton plasmids with no close relatives are indicated as pSin.

<sup>3</sup>Plasmid DNA length is shown in base pairs (bp).

<sup>4</sup>The predicted numbers of open reading frames (ORFs) are listed.

<sup>5</sup>Plasmid copy number is calculated from the fold-difference between the average number of sequenced reads of the plasmid and the corresponding genome. If there is more than one identical plasmid, the average is reported.

<sup>6</sup>Plasmid pATCC19977-1 is the same as the previously reported plasmid in this strain (Ripoli *et al.*, 2009).