GenBank and WGS accession numbers for the sequences used to construct the *A. baumannii* OCL database

OC locus	Reference isolate	GenBank or WGS accession number	Base range	Reference
OCL1	A1	CP010781.1	3366405..3375181	(1-3)
OCL2	D36	CP012952.1	646907..655270	(1, 3)
OCL3	A85	CP021782.1	3464825..3473737	(1, 3)
OCL4	A388	CP024418.1	642040..650673	(3)
OCL5	G21	MG231275.1	37120..46166	(3)
OCL6	D46	KF030679.1	28675..37977	(3)
OCL7	OIFC035	AMTB01000038.1 [1]	221522..230586	(3)
OCL8	OIFC111	AMFY01000013.1 [1]	222496..228777	(3)
OCL9	Naval-72	AMFI01000027.1 [1]	34336..40843	(3)
OCL10	AB_TG27343	AMIS01000032.1 [1]	97841..111169	(3)
OCL11	TG22204	ASFV01000009.1 [1]	33588..43841	(3)
OCL12	NIPH 410	ATGJ01000006.1 [1]	47175..57655	(3)

[1] GenBank or WGS sequence not including current nomenclature and naming

References

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2. Kenyon J, Holt K, Pickard D, Dougan G, Hall R. Insertions in the OCL1 locus of *Acinetobacter baumannii* lead to shortened lipooligosaccharides. Research in Microbiology. 2014;165(6):472-5.
3. Kenyon J, Nigro S, Hall R. Variation in the OC locus of *Acinetobacter baumannii* genomes predicts extensive structural diversity in the lipoligosaccharide. PLoS One. 2014;9(9):e107833.