

Diverse Genetic Background of Multidrug-Resistant *Pseudomonas aeruginosa* from Mainland China, and Emergence of an Extensively Drug-Resistant ST292 Clone in Kunming

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Supplementary Table S1. New sequence types (STs) and allelic profiles found in this study.

Isolate ID no.	ST	<i>acsA</i>	<i>aroE</i>	<i>guaA</i>	<i>mutL</i>	<i>nuoD</i>	<i>ppsA</i>	<i>trpE</i>	Gender^a	Age	Collection year	Collection month	Comments
GPMDP018	1950	40	22	11	14	4	4	7	M	65	2012	April	
HBMDP006	1956	1	4	1	11	4	10	10	F	63	2011	November	
HBMDP020	1960	16	5	77	3	1	4	7	M	88	2012	April	
HNMDP002	1963	18	5	5	3	1	17	13	F	71	2011	July	
HNMDP015	1964	16	66	11	3	4	29	7	M	40	2011	October	
HZMDP011	1965	158	4	1	10	3	6	3	M	77	2011	September	
SCMDP001	1966	6	5	11	3	4	3	141	M	87	2011	August	
SCMDP004	1967	39	166	41	5	4	6	7	M	61	2011	September	IMP-9-producer
SDMDP037	1968	17	166	6	157	4	4	13	M	74	2012	June	
SJMDP015	1969	111	30	31	141	30	24	55	M	57	2012	March	
SZMDP006	1970	11	10	36	3	4	38	3	M	90	2011	November	
TJMDP022	1971	32	190	3	62	8	7	26	M	63	2012	March	
XYMDP001	1972	17	134	1	3	4	15	7	F	61	2012	June	

XYMDP012	1973	1	5	91	7	4	12	7	M	12	2011	October	IMP-9-producer
XYMDP029	1974	39	5	6	11	4	7	193	M	45	2012	August	VIM-2-producer
ZSMDP007	1975	25	76	36	11	44	4	10	F	53	2011	October	
ZZMDP021	1976	17	76	11	72	3	10	3	M	36	2012	June	IMP-10-producer

^a M, male; F, female.