Supplemental Table 1. Serotypes and antimicrobial resistance determinants of global *Pseudomonas aeruginosa* ST111

	A (n=43)	B (n=37)	C1 (n=32)	C2 (n=857)	Total (n=969)
QRDR mutations					
*gyrA*_D87N	1	1	1	14	17
*gyrA*_T83I	7	12	16	843	878
*parC*_S87L	1	2	10	726	739
*parC*_S87W	-	4	1	18	23
Serotypes					
O4	43	37	28	2	110
O12	0	0	0	855	855
O11	0	0	4	0	4
Aminoglycoside modifying enzymes					
aac(2')-I(A267)	-	-	-	46	46
aac(3)-Ia	-	4	-	6	10
aac(6')-29	1	-	-	204	205
aac(6')-29a	-	-	-	409	409
aac(6')-29b	-	-	-	391	391
aac(6')-Ib	-	-	-	116	116
aac(6')-Ib'	-	-	-	120	120
aac(6')-Ib3	-	-	-	54	54
aac(6')-Ib4	-	-	8	50	58
aac(6')-Il	2	4	-	13	19
aadA1	-	1	-	19	20
aadA2	-	-	-	167	167
ant(2'')-Ia	-	3	1	21	25
ant(4')-IIb	-	1	-	11	12
aph(3'')-Ib	6	1	1	25	33
aph(3')-IIb	43	37	32	857	969
aph(3')-VIa	1	2	-	14	17
aph(6)-Id	6	1	1	23	31
β-lactamases					
$bla_{\text{CARB-2}}$	-	-	-	171	171
$bla_{\text{IMP-18}}$	-	-	-	24	24
$bla_{\text{IMP-4}}$	-	-	-	42	42
$bla_{\text{KPC-2}}$	1	-	-	14	15

bla$_{OXA}$[a]	1	-	-	10	11
bla$_{OXA-10}$	-	-	-	15	15
bla$_{OXA-2}$	-	-	4	41	45
bla$_{OXA-395}$	42	37	32	853	964
bla$_{OXA-9}$	-	-	-	65	65
bla$_{PDC}$[a]	-	-	1	13	14
bla$_{PDC-1}$	43	-	-	-	43
bla$_{PDC-3}$	-	37	32	842	910
bla$_{TEM-116}$	-	-	-	69	69
bla$_{VIM}$[a]	-	-	-	18	18
bla$_{VIM-2}$	1	-	4	604	609
Other antimicrobial resistance determinants					
catB7	43	36	31	825	935
cmlB	-	-	-	66	66
crpP	39	32	31	845	947
fosA	43	37	32	857	969
*ftsI*_R504C	-	-	-	23	23
mph(E)	-	-	-	12	12
msr(E)	-	-	-	12	12
*pmrB*_V15I	43	37	32	843	955
qacE	-	-	4	31	35
qacEdelta1	2	9	4	817	832
qacG2	2	-	4	38	44
sul1	2	9	11	827	849
tet(G)	-	-	-	21	21
Large insertion	0	1	0	14	15
Missense	0	0	0	2	2

QRDR; Quinolone Resistant Determining Regions

[a] Subtype was undeterminable.