

ORIGINAL RESEARCH

Capsule Thickness, Not Biofilm Formation, Gives Rise to Mucoid *Acinetobacter baumannii* Phenotypes That are More Prevalent in Long-Term Infections: A Study of Clinical Isolates from a Hospital in China

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Xiaoting Hua Tel/Fax +86-571-86006142 Email xiaotinghua@zju.edu.cn **Background:** *Acinetobacter baumannii* is a nosocomial pathogen of critical importance due to the increasing numbers of antibiotic-resistant isolates. Colonies can have a smooth or matt appearance, but also exhibit slimy, mucoid growth, with the latter being increasingly isolated in patients in recent years.

Methods: We isolated 60 *A. baumannii* strains from altogether 56 patients and found that all patients were infected by mucoid strains, with four patients having also matt phenotypes in addition to the mucoid ones. The morphology of the colonies and capsules was observed. The antibiotics susceptibilities were tested, and the biofilm formation ability was determined by crystal violet staining. The whole-genome sequencing (WGS) was performed on all the strains, and then the core genome multilocus sequence typing (cgMLST) and drug resistance gene analysis were performed. Finally, a part of isolates were selected to test virulence in a *Galleria mellonella* model.

Results: We observed much larger capsules in the mucoid strains compared to the matt isolates. But the mucoid phenotype did not correlate with the amount of biofilm produced by the strain. Almost all mucus-type *A. baumannii* were multi-drug resistant isolates, containing various antibiotic resistance genes. The main ST types of mucoid-type *A.baumannii* were ST191 and ST195, of which ST191 isolates were more virulence, while ST195 isolates were weaker

Conclusion: The mucoid *A. baumannii* had resistance to most antibiotics and some strains had high virulence, which should be paid attention in clinical.

Keywords: Acinetobacter baumannii, capsule, biofilm formation, mucoid phenotype

Introduction

Acinetobacter baumannii has become an important nosocomial pathogen causing a range of infections, including skin and soft tissue infections, infections of the urinary tract, endocarditis and meningitis. 1,2 Due to its ability to acquire various antibiotic resistance, more than 80% of Acinetobacter species are defined as multidrug-resistant (MDR), and without treatment options result in high mortality rates, while presenting a huge healthcare burden in general. 3,4

Acinetobacter baumannii isolates have been found to exhibit both, mucoid and non-mucoid (matt) phenotypes. In case of *Pseudomonas aeruginosa*, during the course of an acute early infection establishing itself as chronic infection,

a previously matt phenotype can "switch" to a mucoid one which shows reduced growth rate concurrent with a changed metabolic pathway and over-expression of alginate.5 The mucoid Ps. aeruginosa isolates secrete a variety of substances including various polysaccharides to form a biofilm, which protects it from antibiotics and contributes to persistence in the host as well as to virulence in general.^{6,7} In *Klebsiella pneumonia*, hypermucoviscous K.pneumonia isolates are considered to be hypervirulent.⁸ In A. baumannii, the K locus cluster is responsible for capsular biosynthesis, 9 with several genes involved in the regulation process, including itrA and wzc.¹⁰ When these genes are non-functional, the nonencapsulated mutants are quickly inactivated and killed by serum. 11 We previously reported that A. baumannii are able to evolve to a mucoid phenotype in vivo. 12 In this work, we aimed to define the molecular basis of mucoid A. baumannii formation, but also analysed antibiotic resistance and virulence of mucoid isolates.

Materials and Methods

Ethics Statement

The study was carried out in accordance with the principles of the Declaration of Helsinki and was approved by the Research Ethics Committee of Hangzhou General Hospital of Chinese People's Armed Police which waived the need for consent, as the patient information was anonymized and deidentified prior to analysis.

Patients and Bacteria Strains

60 *A. baumannii* isolates from 56 hospitalized patients in Chinese People's Armed Police Forces, Hangzhou during 2014–2018, were collected. After obtaining single colonies, all isolates were identified as *A. baumannii* by 16S rRNA analysis (primers: forward, 5'-ACATCGTTTACGG CATGGACT-3', and reverse, 5'-TGCCGCGTGTGTGAA GAA-3'). ATCC19606 and MDR-ZJ06¹³ were taken as reference isolate in this study.

Capsule Thickness Measuring

Bacterial capsules were detected using the India ink method. ¹⁰ Briefly, bacterial colonies and India ink were mixed in 0.85% NaCl solution. And the bacteria were observed and photographed under light microscopy (1000×). Ten capsules were measured for each isolate by image J software (https://imagej.nih.gov/ij/).

WGS and Core Genome Multilocus Sequence Typing (cgMLST)

The genomic DNA of 60 isolates were extracted using a QIAamp DNA Mini kit (Qiagen, Valencia, CA, USA) following the manufacturer's recommendations, and were sequenced at Zhejiang Tianke (Hangzhou, China) on an Illumina HiSeq2000 platform (Illumina, San Diego, CA, United States) following a pair-end 2×150 bp protocol. *De novo* assembled using CLC Genomic Workbench 8.5.1 (Qiagen, Aarhus, Denmark). Phylogenetic analyses were performed based on validated *A. baumannii* cgMLST version 1.0 (2390 target genes), using Ridom SeqSphere+software (Ridom GmbH, Munster, Germany). 14

Antimicrobial Susceptibility Testing

The Minimal Inhibitory Concentrations (MICs) of colistin, tigecycline and eravacycline were determined by the broth microdilution method with cation-adjusted Mueller-Hinton broth (CAMHB), and MICs of the other antibiotics, including ceftazidime, cefepime, cefoperazone/sulbactam, ciprofloxacin, amikacin, gentamicin, imipenem, meropenem, were detected by the agar dilution method according to the Clinical and Laboratory Standards Institute (CLSI) guidelines (CLSI-M100-S26). The results of tigecycline and eravacycline were interpreted according to EUCAST breakpoints, and the breakpoint of eravacycline was referred to tigecycline.

Growth Rate Determination

Four independent cultures per strain were grown overnight and diluted to 1:100 in MH broth before aliquots were placed into a flat-bottom 100-well plate with three replicates each. The plate was incubated at 37°C with agitation while the OD₆₀₀ of each culture was determined every 5 min for 20hrs using a Bioscreen C MBR machine (Oy Growth Curves Ab Ltd., Finland). The growth rate was determined by applying a linear regression to the exponential growth rate data using an R script.¹⁵

Biofilm Assay

Biofilm assays of each isolate were performed in triplicates as described previously, ¹⁶ with slight modifications. Overnight bacterial cultures were diluted by fresh Luria-Bertani (LB) medium until the bacterial suspension reached an OD₆₀₀ of 0.5. Two hundred milliliters of the cell number-adjusted bacterial cultures were transferred to 96-well polystyrene plates, and incubated for 72 h at 37°C.

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Cells were washed three times with phosphate-buffered saline (PBS), and stained with 0.1% crystal violet for 20 min at room temperature. Wells were washed again and dried naturally. The bound dye was solubilized with 200 µL of 95% ethanol and quantified by measuring the absorbance at 540 nm. Three wells containing LB without bacteria served as a negative control. Strong biofilmforming ability was defined that OD540 of the isolate was more than ATCC19606. Weak biofilm-forming ability was defined with OD₅₄₀ between ATCC19606 and negative control. If OD₅₄₀ of the isolate was less than the negative control, it was defined as no biofilm-forming ability.

In vivo Virulence Study Using G. mellonella

Virulence was evaluated in the insect model G. mellonella as described previously.¹⁷ Briefly, groups of 10 larvae (~200 mg; Yuejiayin, Tianjin, China) were stored in the dark at 4°C prior to use. To determine the virulence of isolates, every larva was infected with 1×10^6 CFU bacteria (n = 10), and 10 larvae injected with 10 μL PBS as a negative control. Survival was monitored up to 72hrs post-infection at 37°C. Experiments were performed in triplicate and re-tested when the difference between the two experiments was >15%. Data from all experiments were combined to calculate the mean values of percent survival values.

Statistical Analysis

Statistical analyses were performed using GraphPad Prism 6 software (GraphPad Software Inc., La Jolla CA, USA). Independent samples Student's t-test was used to determine differences between mucoid and matt A. baumannii in capsule thickness, growth rate and biofilm-forming ability. Survival rates were evaluated using Kaplan Meier survival curves and analyzed with the log-rank (Mantel-Cox) method. Chi-squared or Fisher's exact tests were used to determine differences in the number or percentage of isolates between isolates. Differences were considered significant at P < 0.05.

Results

Clinical Parameters of Patients Infected with A. baumannii

The clinical data of the 56 patients are shown in Table 1. The median age of the patients was 61.68 ± 14.29 years, with 60.7% male patients. The average hospitalization time was 90.4 \pm 82.5 days with 41.1% having been

Table I Baseline Characteristics of Patients and Antimicrobial **Treatments**

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· · · · ·	Invalid	19 (33.9%)
Death 3 (5.4%)	Deterioration	10 (17.9%)
	Death	3 (5.4%)

Notes: alnvasive procedure: including endotracheal intubation or incision, central venous catheterization, arterial catheterization, and hemodialysis tube, angiography or embolization, PICCO, \geq 2 punctures (thoracic puncture, abdominal puncture, lumbar puncture, bone marrow puncture). bOperation: including operations during hospitalization and I month before admission. Antibiotic treatment: uses of antibiotics within half a year prior to isolation. dOthers: linezolid, sulfonamides.4

admitted to an Intensive Care Unit (ICU). The most prevalent comorbidities were cerebrovascular diseases followed by hypertension and diabetes. A 89.3% of the patients underwent invasive procedures, with 30.4% undergoing an operation.

The previous uses of antibiotics half a year prior to isolation of the microbial strains were recorded. The antibiotics were classified into 11 categories, with more than 50% of the patients having used β-lactam, β-lactamase inhibitor combination or carbapenem antibiotics. The clinical outcome of the patients during the duration of the study found only 42.9% of the patients fully cured, while the medical condition of approx. 18% deteriorated, and 3 died during the duration of the study, one of the bacterial infections.

The Majority of the A. baumannii Isolates Display a Mucoid Phenotype

From the 56 patients, we isolated 60 *A. baumannii* strains and found that all patients were infected by mucoid strains, with four patients having also matt phenotypes in addition to the mucoid ones. The colonies of mucoid *A. baumannii* appeared moist, irregular round, with an elevated surface. Adjacent colonies often formed fused associations on agar plates (Figure 1). In contrast to this, matt *A. baumannii* colonies had a smaller, round appearance with distinct edges.

Mucoid A. baumannii Isolates Display Thicker Capsules

After determining the matt and mucoid phenotypes of the isolates, we then analysed the capsular types of the 60 strains. Here, we found that the capsular types of 29 strains were so-far unknown, and the remaining 31 strains belonged to six capsular types; 17 strains belong to the type KL3, five strains to KL2, four strains to KL9, two strains to KL14, two strains to KL7, and one strain to KL77. We then visualized the capsules of mucoid and matt *A. baumannii* isolates by India ink staining, and observed by light microscopy that matt *A. baumannii* had much thinner capsules compared to mucoid isolates (Figure 2A–D). The capsule thickness was determined by





mucoid (403)

matt(404)

Figure I Colony morphology of the A. baumannii isolates on MH plates. Isolate 403 was mucoid with moist, irregular round colonies, and isolate 404 was matt with round and neat edge colonies.

measuring the diameter of the cell halos (including the cells), with the capsule thickness of the reference strain MDR-ZJ06 (matt phenotype) as a comparison. The average capsule thickness of mucoid A. baumannii isolates was determined to be $3.24~\rm um~\pm~0.60$, more than $2.5~\rm times$ that of the matt phenotype (p < 0.001). Interestingly, we found that the capsule thickness of KL7 isolates was significantly higher than that of other capsular types (p < 0.05). In contrast to this observation, no differences among the rest of the capsular types were observed (Figure 2E). In the four patients where we isolated both, matt and mucoid phenotypes, the capsule thickness of mucoid-type A. baumannii was also significantly higher than that of the matt isolate (p < 0.05) (Figure 2F).

Antibiotic Resistance Did Not Correlate with Capsule Thickness

The MICs of all isolates for 11 antibiotics were determined (Table 2). The resistance rate to cephalosporins, quinolones and carbapenems antibiotics was up to 96.7%, followed by aminoglycoside antibiotics (amikacin of 73.3% and gentamicin of 83.3%). The susceptibility to cefoperazone/sulbactam, tigecycline and colistin was lower, with resistance rates of 15.0%, 35.0% and 3.3%, respectively. The novel antibiotic eravacycline, which belongs to the tetracycline class, had the highest antibacterial activity to mucoid *A. baumannii* with a resistance rate of only 1.7%. And the 4 matt isolates (isolate 402, 404, 406 and 408) had similar resistances, which were only sensitive to colistin and eravacycline.

Genomic Data, Sequence Types and AMR Genes of the Isolates

According to the analysis by cgMLST, the 60 isolates belonged to eight different sequence types (ST), including five new types (Figure 3). The main ST of mucoid *A. baumannii* were ST191 and ST19, followed by ST 368, 208, 136, 547, 784 and 938. Having obtained the wholegenome data, we were able to determine the occurrence of resistance genes and the mutations that confer resistance to the isolates (Figure 4). There were only two resistance genes in the isolate N35, which was susceptible to all 11 antibiotics (Table 2). The isolate N32 and N37 contained four resistance genes, *bla*_{ADC-25}, *bla*_{OXA-66}, *bla*_{OXA-23}, and *sul2*, which confer resistance to β-lactam and carbapenem antibiotics. The remaining 57 isolates all contained more than five resistance genes. The copy number of resistance genes in all isolates

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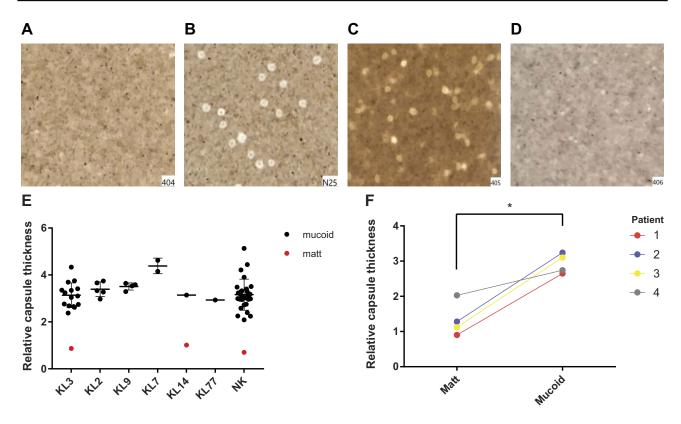


Figure 2 Capsule thickness. (A-D) (A) Matt isolate 404; (B) mucoid isolate N25; (C, D) mucoid isolate 405 and matt isolate 406, isolated from the same patient. (E) The average capsule thickness of different capsular types strains. Mucoid A baumannii isolate was more than 2.5 times that of the matt phenotype (p < 0.001); (F) Capsule thickness of 8 isolates, isolated from 4 patients, one matt isolate and one mucoid isolate were isolated from each patient. *p < 0.05; ***p < 0.0001.

was 1, except that isolate N7 contained double copy of the *sul2* gene. The mutation in *adeN* or *adeR* appeared in 30 isolates. The gene *adeR* is involved in the regulation of efflux pump AdeABC and *adeN* regulates efflux pump AdeIJK, which both contribute to tigecycline resistances. The isolates contained the mutation all not susceptible to tigecycline (Table 3).

Table 2 Antimicrobial Susceptibility in A. Baumannii

Antibiotics	Number of Isolates			Resistance
	Susceptible	Intermediate	Resistant	Rate (%)
CAZ	1	1	58	96.7
FEP	1	1	58	96.7
CIP	1	1	58	96.7
IMP	1	1	58	96.7
MEN	1	1	58	96.7
GEN	4	6	50	83.3
AK	16	0	44	73.3
TGC	4	35	21	35.0
SCF	12	39	9	15.0
CST	58	0	2	3.3
ERA	46	13	1	1.7

Abbreviations: CAZ, ceftazidime; FEP, cefepime; CIP, ciprofloxacin; IPM, imipenem; MEM, meropenem; GEN, gentamicin; AK, amikacin; TGC, tigecycline; SCF, cefoperazone/sulbactam; CST, colistin; ERA, eravacycline.

Single Nucleotide Polymorphism (SNP) Analysis

To investigate possible molecular mechanisms for the appearance of mucoid or matt phenotypes, we analysed the SNPs of the double isolates that were obtained from one patient. The isolates 405 and 406 were obtained from one patient, and 407 and 408 were obtained from a second patient. All four isolates belonged to ST195. According to the sequence comparison of isolate 405 (mucoid), isolate 406 (matt) revealed two mutations in gene *ptk* (Table 4). The other two isolates, 407 (mucoid) and 408 (matt), also showed one mutation in the same gene, *ptk* (Table 5). This result indicated that gene *ptk* might be a key determinant for matt versus mucoid phenotypes during capsule biosynthesis, an observation that was consistent with previous studies. ^{10,12}

No Correlation Between Biofilm Formation and Mucoid/Matt Phenotypes of A. baumannii Isolates

Previously, it was reported that matt *A. baumannii* isolates produced more biofilms than mucoid ones. Therefore, we determined the biofilm-forming ability

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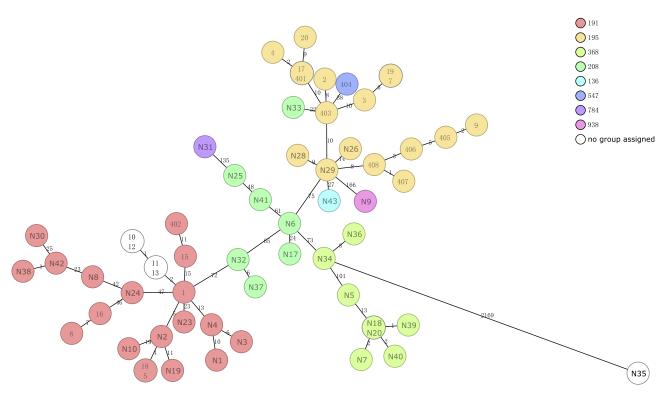


Figure 3 Minimum spanning tree based on cgMLST analysis. Each circle represents an allelic profile, i.e. genotype, based on sequence analysis of up to 2390 target genes. The numbers on the connecting lines illustrate the numbers of target genes with different alleles.

of all strains by crystal violet staining, which can be quantified by measuring the optical density at 540 nm, OD₅₄₀. As reference served the type strain ATCC19606. While the reference strain was set to 1 ± 0.18 , the negative control with media only, gave a value of 0.25 ±0.07. Three isolates formed no biofilms with OD₅₄₀ values lower than the negative control. Fifty (83.3%) isolates showed weak biofilm-forming ability with OD₅₄₀ values between the negative control and the reference strain (0.63 \pm 0.22). The remaining seven isolates (11.7%) formed extensive biofilms with OD₅₄₀ values significantly higher than the control strain ATCC19606 (1.23 \pm 0.30). While a significant difference between the groups with a different biofilmforming ability (p < 0.05) was observed, no significant difference was observed between the mucoid A. baumannii and the matt isolates (0.68 \pm 0.11, 0.66 ± 0.13 , p > 0.05), and also no differences between different capsular types (Figure 5A).

Mucoid and Matt A. baumannii Isolates Show Similar Growth Rates

Previously as the study reported that mucoid A. baumannii isolates display a slower growth rate

than their matt counterparts.¹⁸ We, therefore, measured the growth rates of all isolates. Taking the growth rate of isolate 1 (a mucoid strain) as a reference, the relative growth rates of all isolates were calculated. Our results showed that there was no significant difference between mucoid *A. baumannii* and matt *A. baumannii* (p > 0.05) in the relative growth rate, and no differences between different capsular types (Figure 5B).

No Correlation in Virulence with Matt versus Mucoid A. baumannii in the G. mellonella Model

According to ST and resistance genes of all isolates, we selected the isolate 13, N9, N10, N20, N25, N29, N31, N35, N43, and the two double isolates (matt and mucoid together; 401–408), a total of 17 isolates for virulence test in *G. mellonella* (Figure 6A–D). The virulence of 17 clinical isolates varied widely, with an average 72 h survival rate ranging from 3.33% to 73.33%. The percent survival of the isolate 402, 405, 13, N9, N10, N25, N31 and N35 was less than 50%, among which, the percent survival of 402, 13 and N35 was lower than 10%. The isolates 401, 403, 404, 406, 407, 408, N20, N29 and N43 displayed a lower virulence level, with survival rates at 72 hrs of higher than

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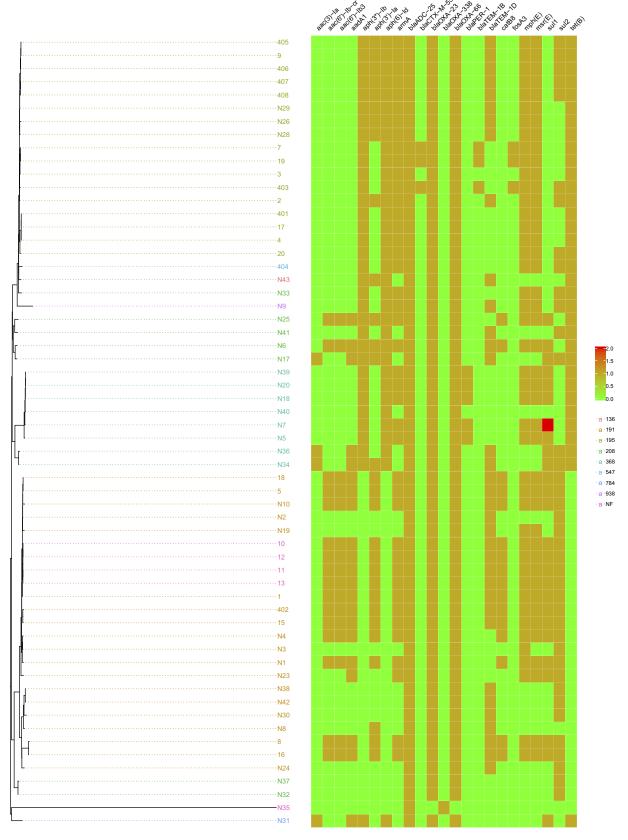


Figure 4 Resistance genes. The isolate N35 only had two resistance genes, which was susceptible to all 11 antibiotics (Table 2). The isolate N32 and N37 contained four resistance genes, and the remaining 57 isolates all contained more than five resistance genes.

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Table 3 adeN and adeR Mutation Conferred Tigecycline Resistance

Isolates	TGC(mg/L)	Mutation
1	8	adeN ISAba1 299
5	8	adeN ISAba1 299
10	8	adeN ISAba1 299
11	4	adeN ISAba1 299
12	8	adeN ISAba1 299
13	8	adeN ISAba1 299
15	4	adeN ISAba1 299
18	8	adeN ISAba1 299
402	8	adeN ISAba1 299
NI	4	adeN ISAba1 299
N2	4	adeN ISAba1 299
N3	4	adeN ISAba1 299
N4	4	adeN ISAba1 299
N5	4	adeN IS26 277
N6	4	adeN ISAba1 299
N7	8	adeN IS26 277
NI0	4	adeN ISAba1 299
NI7	4	adeN ISAba1 299
NI8	4	adeN IS26 277
NI9	>32	adeN ISAba1 299
N20	8	adeN IS26 277
N23	16	adeN ISAba1 299
N25	8	adeN ISAba1 288
N31	16	adeR D23N N115T
N32	8	adeN ISAba1 299
N34	8	adeN ISAba1 299
N36	8	adeN ISAba1 299
N37	8	adeN ISAba1 299
N39	8	adeN IS26 277
N40	4	adeN IS26 277
N41	4	adeN ISAba1 288

50%. Interestingly, when testing the double isolate that was obtained from one patient, the virulence of the matt isolate (402) was significantly higher than the mucoid one (401, p < 0.0001). In must be mentioned that 401 belongs to ST191, while 402 belongs to ST195 which makes it difficult to conclude whether virulence is influenced by the matt or mucoid phenotype of *A. baumannii*. An indication that it might the phenotypes might not correlate with virulence is the data we obtained from the other two double isolates, obtained from a single patient each (403, 404 and 407, 408). When comparing these isolates, no difference in virulence could be observed. We also were able to determine that the isolates of the capsule-type KL3 generally displayed a lower virulence level, while isolates N31 (KL14) and N25 (KL2) had a higher virulence level. Interestingly, we

Table 4 SNPs Analysis of Isolate 405 (Mucoid) and 406 (Matt)

Gene	Туре	Product	Effect
sppA	SNP	Putative signal peptide peptidase SppA	missense_variant c.415A>G p.Lys139Glu
ftsI	SNP	Peptidoglycan D, D-transpeptidase Ftsl	missense_variant c.1589G>A p.Arg530His
tadA	SNP	tRNA-specific adenosine deaminase	missense_variant c.109T>C p.Ser37Pro
ptk	SNP	Tyrosine-protein kinase	missense_variant c.2000A>G p.Asp667Gly
			missense_variant c.1628C>T p.Pro543Leu
рþх	SNP	Exopolyphosphatase	missense_variant c.1441A>G p.Thr481Ala
rapZ	del	RNase adapter protein RapZ	frameshift_variant c.688delA p.Met230fs
ibaG	SNP	Acid stress protein IbaG	missense_variant c.65A>G p.Asn22Ser
_	SNP	Hypothetical protein	missense_variant c.443C>T p.Pro148Leu
_	del	Putative TonB- dependent receptor	frameshift_variant c.1297delT p.Trp433fs

Table 5 SNPs Analysis of Isolate 407 (Mucoid) and 408 (Matt)

Gene	Туре	Product	Effect
nimR_I	SNP	HTH-type transcriptional regulator NimR	missense_variant c.578A>G p.His193Arg
rpsA	SNP	30S ribosomal protein	missense_variant c.410G>A p.Gly137Asp
ptk	SNP	Tyrosine-protein kinase	missense_variant c.833T>A p.Leu278GIn
fhuA_4	SNP	Ferrichrome-iron receptor	synonymous_variant c.273A>G p.Leu91Leu
_	del	Hypothetical protein	frameshift_variant c.3000delA p.Val1001fs

found highly virulent isolates with so-far unknown capsular types, such as isolate 13 and N35.

Discussion

In 1998 the first mucoid *A. baumannii* strains in China were reported, which were isolated from the sputum of

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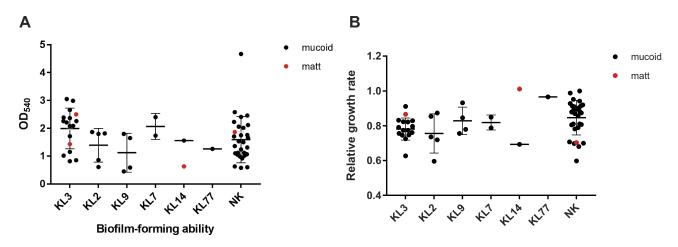


Figure 5 Biofilm-formating ability and relative growth rate. (A) Biofilm-forming ability of different capsular types. (B) Relative growth rate of different capsular type strains, taking the isolate I as reference. Both biofilm-forming ability and relative growth rate had no significant differences between matt and mucoid isolates.

Abbreviation: NK, unknown capsular types.

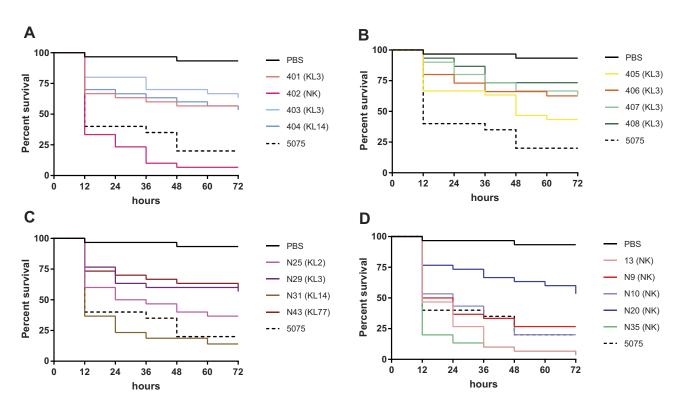


Figure 6 Percent survival of *G. mellonella* infected with *A. baumannii*. (**A**) 401 (mucoid, ST195, KL3) and 402 (matt, ST191, NK) isolated from one patient, and (mucoid, ST195, KL3) and 404 (matt, ST547, KL14) isolated from one patient; (**B**) 403 (mucoid) and 404 (matt) isolated from one patient, and (mucoid) and 404 (matt) isolated from one patient, all 4 isolates belong to ST195, KL3; (**C**, **D**) all isolates were mucoid. **Abbreviation:** NK, unknown capsular types.

patients with chronic bronchitis. ¹⁹ As the isolation of mucoid *A. baumannii* strains is comparably rare in the clinical setting, reports about their isolation rate are lacking to date. We observed that the quantity of mucoid *A. baumannii* isolates is increasing in clinical care. Mucoid bacteria are often associated with chronic

infections, with two major factors possibly contributing to the long persistence in the host: higher levels of antibiotic resistance and stronger attachment due to the mucoid nature of the bacterial colonies. ^{20,21} In our study, we could demonstrate that the capsules of the mucoid *A. baumannii* isolates were significantly thicker than that

of matt *A. baumannii* isolates. In contrast to previous reports, ¹⁸ we could not conclude that matt cells form more biofilm than mucoid ones, as we observed no significant differences in biofilm-forming ability between the phenotypes. In addition, our data showed that the growth rate between the different phenotypes was highly similar.

Our data showed that up to 77% of patients had used more than three categories of antibiotics prior to isolation of the bacteria, which may have led to the observed high resistance rates to major antibiotics in mucoid *A. baumannii*. Notably, the resistance rate of tigecycline had reached up to 35.0%. Wholegenome sequencing revealed that 30 isolates harbored tigecycline resistance mutations in *adeN* and one isolate had a mutation in *adeR*. Other mutations were observed which might confer resistance, although more experiments are required to fully understand the resistance mechanism.

The major sequence types (ST) in this study were ST191 and ST195, which were previously reported as the major STs carrying carbapenem-resistance gene $bla_{\rm OXA-23}$ in the carbapenem-resistance A. baumannii in China.²² Consistently, all ST191 and ST195 isolates in our study carrying a single copy of $bla_{\rm OXA-23}$ were shown to be resistant to carbapenem antibiotics. Interestingly, the survival rate of two ST191 isolates (402 and N10) was similar to AB5057, a model highly virulent A. baumannii strain, 23 but the survival rate of six ST195 isolates (401, 405-408, N29) was significantly higher than AB5075 (p < 0.05). These results indicate that ST191 might belong to the more virulent isolates whereas ST195 might be regarded as isolates with comparably lower virulence. It is noteworthy that the six ST195 isolates all belonged to KL3. Infections with these isolates and another KL3 isolate (403) had significantly higher survival rates compared to AB5075 (p < 0.05), which indicates that the capsule type KL3 might be less virulent. For the other types, more data are required to be able to correlate capsule type with the virulence of the strain.

When performing the virulence test in *G. mellonella*, we could not see any correlation between virulence and the matt or mucoid phenotype. While this test is usually a good indicator of virulence, the larva model might not be the best test of the human pathogen. Therefore, more data and possibly other tests in higher organisms are required to understand virulence levels of the isolates and a possible correlation with the mucoid or matt phenotype.

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Disclosure

The authors report no conflicts of interest in this work.

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