

# Nomogram Analysis of Clinical Characteristics and Mortality Risk Factor of Non-Fermentative Gram-Negative Bacteria-Induced Post-Neurosurgical Meningitis

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**Objective:** To explore the clinical characteristics of post-neurosurgical meningitis (PNM) patients infected with nonfermenting Gram-negative bacilli (NFGNB) and to evaluate the related mortality risk factors.

**Methods:** A cohort analysis of PNM patients infected with NFGNB in Beijing Tiantan Hospital and Capital Medical University from 2012.1 to 2020.12. The microbial distribution, antimicrobial sensitivity and genotypes were tested, and potential mortality risk factors were evaluated using Mann-Whitney U or chi-squared tests. Independent risk factors for mortality were established by constructing a logistic model.

**Results:** A total of 2940 PNM patients were enrolled in this study, of whom 207 (17.1%) were infected with NFGNB. Among these patients, 29 died of NFGNB meningitis, with an overall mortality rate of 14.0%. The top three NFGNBs were *Acinetobacter baumannii* (105 cases, 50.7%), *Pseudomonas aeruginosa* (29 cases, 14.0%) and *Acinetobacter lwoffii* (20 cases, 9.7%). Nomogram analysis revealed that hypertension (OR 4.551, 95% CI: 1.464–14.154,  $P = 0.009$ ), external ventricular drainage (EVD) (OR 3.944, 95% CI: 1.286–12.095,  $P = 0.016$ ), and assisted mechanical ventilator (AMV) (OR 6.192, 95% CI: 1.737–22.081,  $P = 0.005$ ) were independent risk factors for mortality. In addition, antibiotic prophylaxis was shown to play a vital role in NFGNB-induced PNM therapy.

**Conclusion:** PNM patients infected with NFGNB have a high mortality rate. Hypertension, EVD and AMV were independent mortality risk factors, and clinical attention should be paid to their prevention and treatment.

**Keywords:** clinical characteristics, nomogram analysis, meningitis, mortality, nonfermenting Gram-negative bacilli

## Introduction

Post-neurosurgical meningitis (PNM) is one of the most common complications in neurosurgical patients, followed by neurosurgery itself, trauma or shunt devices. It is closely related to perioperative morbidity and mortality, with an incidence of 0.7–25%.<sup>1-3</sup> Patients undergoing neurosurgery often have immune deficiency due to complex and lengthy neurosurgical procedures, leading to PNM treatment challenges.<sup>4</sup> Therefore, reducing the incidence of PNM and patient mortality is one of the vital tasks associated with neurosurgery.

The etiology of PNM has shown that a wide spectrum of pathogens can be responsible, including bacteria, fungi, *Cryptococcus*, and so on. Of these, during the last decade, nonfermenting Gram-negative bacilli (NFGNB) have exhibited a trend towards antibiotic resistance and pathogenicity. These pathogens cannot utilize glucose through fermentation, and include *Acinetobacter*, *Pseudomonas*, *Flavobacterium*, *Burkholderia*, etc.<sup>5</sup> With higher antibiotic resistance and

pathogenicity characteristics, NFGNB are perhaps the main types of pathogens causing nosocomial infections.<sup>6,7</sup> In 2017, the World Health Organization (WHO) released a list of bacteria that pose the greatest threat to human health, with carbapenem-resistant *Acinetobacter baumannii* and carbapenem-resistant *Pseudomonas aeruginosa* in the top two rankings (critical), seriously affecting the clinical therapy.<sup>8</sup>

Therefore, we carried out a longitude clinical cohort study in the largest neurosurgery center in China from 2012 to 2020 with the main specific objective of this study is to explore the clinical characteristics of PNM patients infected with NFGNB and to evaluate the related mortality risk factors. Based on the results, neurosurgical surgeons can better accurately prevent or therapeutically target PNM induced by NFGNB. As far as we are aware, this is the first cohort study globally to analyze the clinical characteristics and mortality risk factors for PNM induced by NFGNB.

## Methods and Materials

### Study Design

This study was conducted in Beijing Tiantan Hospital & Capital Medical University. The clinical characteristics of PNM patients from January 2012 to December 2020 were analyzed to evaluate the mortality risk factors. The microbial distribution, drug resistance spectrum, genotypes were also analyzed for epidemiology. Neurosurgery including craniotomy, transsphenoidal surgery and spinal surgery were also investigated. The study was approved by the Ethical Committees of Beijing Tiantan Hospital & Capital Medical University (Approval number: KY-2021-079-02). Each patient enrolled in the study signed an informed consent form.

### Inclusion and Exclusion Criteria

The PNM patients included in this study were followed-up for 90 days, and the inclusion criteria were implement as reported in a previous study,<sup>3</sup> briefly: 1) adult patients (>18 years); 2) patients survived for  $\geq 7$  days; 3) had a NFGNB infection with at least one cerebrospinal fluid (CSF) culture.

Exclusion criteria<sup>9</sup> were patients 1) who had undergone EVD, CSF shunt or stereotaxic surgery; 2) had common clinical contamination bacteria (Coagulase-negative *Staphylococcus*, *Bacillus*, *Propionibacterium*, etc) but without infectious symptoms; 3) who died within 7 days; and 4) without anti-infective treatment in hospital and/or incomplete clinical records.

### Diagnosis Criteria

The definition of NFGNB-induced PNM defined by the US Centers for Disease Control and Prevention (CDC) criteria was followed.<sup>10</sup> Patients with a diagnosis of NFGNB-induced PNM who were included in the study presented with at least one kind of NFGNB proliferation in the cerebrospinal fluid (CSF) or at least one of the signs of meningeal irritation (headache, neck stiffness, or cranial nerve involvement for no other reason). These patients also displayed at least one of the following features: 1) increased protein and/or decreased glucose levels in the CSF; 2) an increased neutrophil count; 3) a positive CSF Gram stain; and 4) a positive blood culture or positive antigen test in the blood or CSF, or an increased antibody titer against the pathogen.

### Microbiology

NFGNBs, isolated from the PNM patients' CSF, underwent bacteria culture, identification, antimicrobial susceptibility tests (ASTs), and genotyping for resistance screening. All the target patient CSF samples underwent a standard procedure for bacterial culture, with 1–3 mL infused into an aerobic automated culture bottle (bioMerieux, Marcy l'Etoile, France). Then, the bottles containing patients' CSF were transferred into a BACT/ALERT<sup>®</sup> 3D automated culture system (bioMerieux, Marcy l'Etoile, France) until the bacteria grew to positive detection levels. Subsequently, 50  $\mu$ L of broth was sub-cultured into a Columbia sheep blood agar for 24 h until obvious colony growth. Bacteria with NFGNB characteristics were identified using VITEK-2 compact and VITEK MS systems (bioMerieux, Marcy l'Etoile, France). All bacteria identified as NFGNB were chosen for AST, and two methods were selected, namely the disk agar diffusion method (Kirby–Bauer method) and the micro-broth dilution method. The breakpoints of each antibiotic followed the Clinical and Laboratory Standards Institute (CLSI) 2021 guidelines (CLSI, 2021). A total of 17 resistances genes from 5 antibiotics resistance phenotypes were selected

for screening. These included meropenem (*bla*<sub>KPC</sub>, *bla*<sub>OXA-66</sub>, *bla*<sub>OXA-23</sub>, *bla*<sub>NDM</sub>, *bla*<sub>VIM</sub>, *bla*<sub>IMP</sub>), ceftriaxone (*bla*<sub>TEM</sub>, *bla*<sub>SHV</sub>, *bla*<sub>CTX-M-9</sub>, *bla*<sub>CMY</sub>, *bla*<sub>CTX-M-1</sub>), amikacin (*aadA1*, *aaaCI*), levofloxacin (*qnrA*, *qnrS*) and polymyxin B (*mcr-1*). The genotypes of target resistance were tested by MNCP-II, a method based by LAMP method which can test 35 different kinds of antibiotic related genes in one panel, and the platform was developed by our research group in 2020.

## Risk Factors

In the present study, all of the NFGNB-induced PNM patients' clinical records were reviewed and the relevant risk factors extracted, including gender, age (years), body temperature (°C), malignant tumor, traumatic brain injury (TBI), diabetes; hypertension, long operation duration (>3 h), reoperation, craniotomy, surgical site (head or spine), type of incision (I or II), ICU admission, CSF leakage, external ventricular drainage (EVD), lumbar drainage (LD), and assisted mechanical ventilation (AMV).

## Therapy

In this study, three categories of antibiotics usage composed the whole therapy regimen, including antibiotic prophylaxis (AP), antibiotic empirical therapy (AET) and antibiotic definitive therapy (ADT). The three categories were defined as follows: 1) PNM patients who had AP, received antibiotics 0.5 h ahead of their neurosurgical operation; 2) PNM patients who had AET received antibiotics before the AST result; 3) patients who had ADT received antibiotic therapy by AST guidance. Three other categories of antibiotic choice were defined as follows: 1) mono-antibiotic: only one type of antibiotic was used in the treatment; 2) dual antibiotic: two different types of antibiotics combination in the treatment; 3) triple or more antibiotic: three or more types of antibiotics combination in the treatment. In addition, the usage rate of high-grade antibiotics (third- or fourth-generation cephalosporin or carbapenem) was evaluated.

## Statistical Analysis

Categorical data are shown given counts and percentages. Numerical data are presented as the mean ± SD or median with interquartile range (IQR), depending on the degree of skewness in the distributions evaluated using the Kolmogorov–Smirnov (K-S) test.  $\chi^2$  or Fisher's exact tests and a *t*-test or Mann–Whitney *U*-test were used for descriptive statistics to evaluate risk factors, when appropriate. To determine the independent risk factors for mortality caused by NFGNB-induced PNM, any variables with a *P*-value <0.05 in the univariate analysis were carried forward for inclusion in the logistic regression model. Statistical significance was defined as *P* < 0.05 and the calibration was analysed with the Hosmer–Lemeshow (H-L) test for goodness-of-fit. Pathogen distribution and AST were performed by WHONET 5.5. Statistical analyses were carried out using SPSS version 22.0 (IBM, New York, USA). Graphs were generated using Prism 7.0 (GraphPad, San Diego, USA).

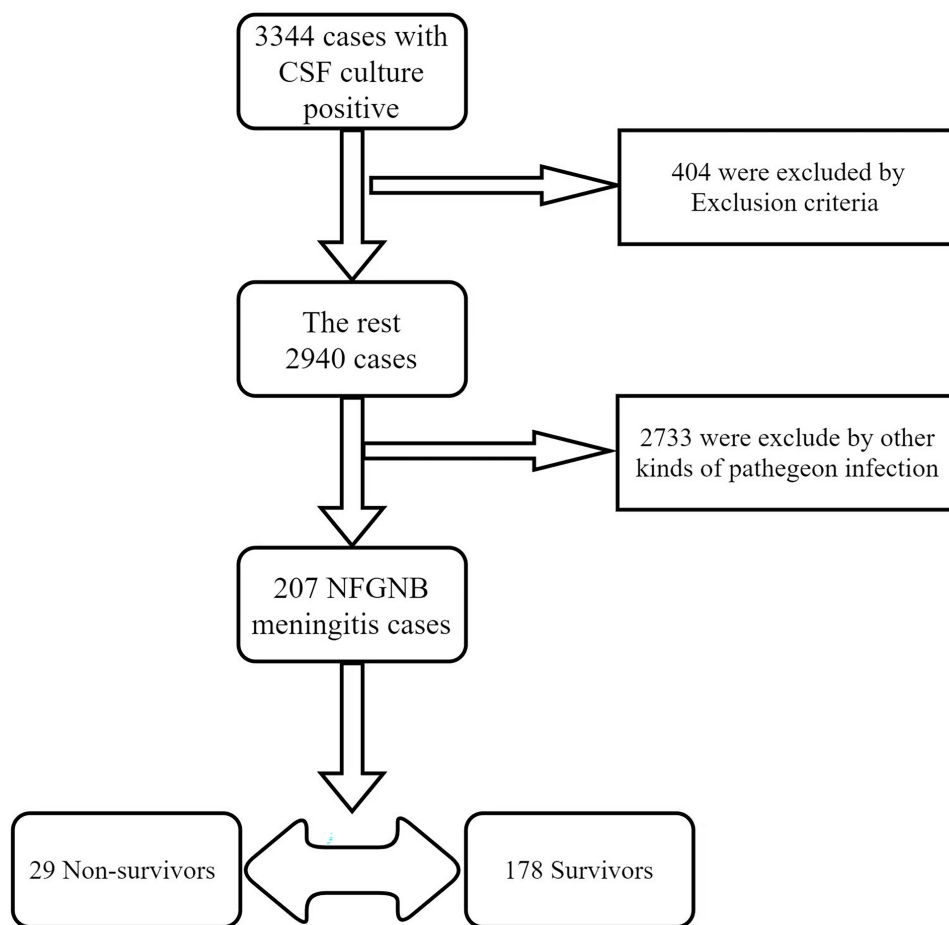
## Results

### Patients

Over the 9-year study period, 3344 episodes of patients with CSF culture positive were recorded in Beijing Tiantan Hospital and Capital Medical University. Among them, 188 patients only underwent a CSF shunt, 73 exhibited cultured *Corynebacterium* or *Micrococcus*, 37 patients survived for <7 days, 64 had incomplete medical records, and 42 patients had multiple infections. A total of 2940 patients were included in the study, 59.4% (1746/2940) were Gram-positive bacterial PNM, and 40.5% (1194/2949) were Gram-negative PNM. In the Gram-negative group, 207 patients (17.3%) were infected by NFGNB, among all NFGNB-induced PNM patients, 29 died from meningitis and 178 survived, with a mortality rate of 14.0%. The flow chart of the study is shown in [Figure 1](#).

### Microbiology Distribution

In 207 NFGNB-induced PNM, *A. baumannii* (105, 50.7%), *P. aeruginosa* (29, 14.0%) and *Acinetobacter lwoffii* (20 cases, 9.7%) occupied the first three classes. The bacterial distribution ratio of patients in the survivors and non-survivor groups was similar; however, the distribution in the non-survivor group was more concentrated, mainly for *A. baumannii* and *P. aeruginosa* ([Figure 2](#)).



**Figure 1** Flow chart of the whole study.

## AST

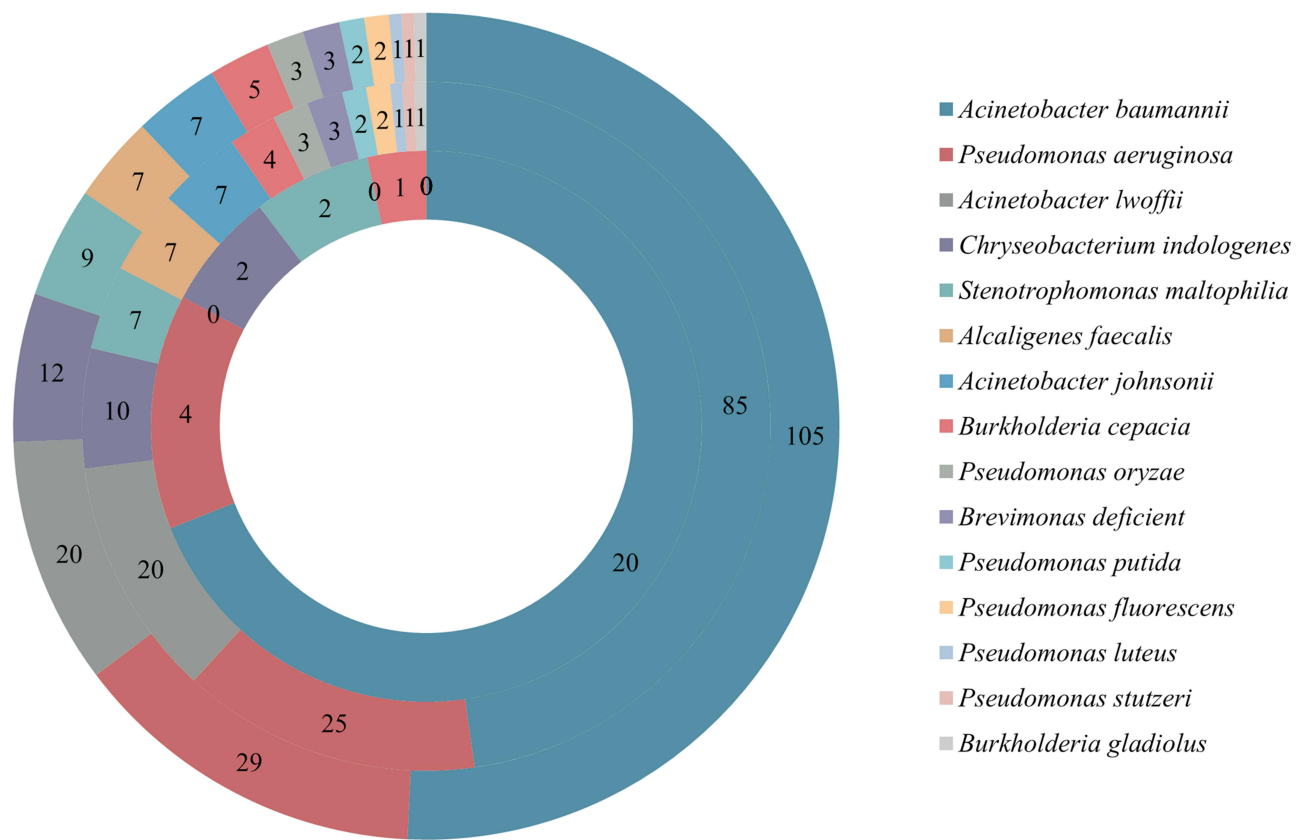
Due to the huge difference in AST of NFGNB, this study conducted statistics on the two types of bacteria (*A. baumannii* and *P. aeruginosa*) with the highest ratio. And, 49.5% of *A. baumannii* were resistance to meropenem, while the resistance rate of *P. aeruginosa* to imipenem and meropenem was different (31.0% vs 24.1%), respectively. For polymyxin B, resistance of *A. baumannii* was slightly lower than that of *P. aeruginosa*, which were 1.0% and 6.9%, respectively. Other AST results are shown in [Figure 3](#).

## Genotypes of *A. baumannii* and *P. aeruginosa*

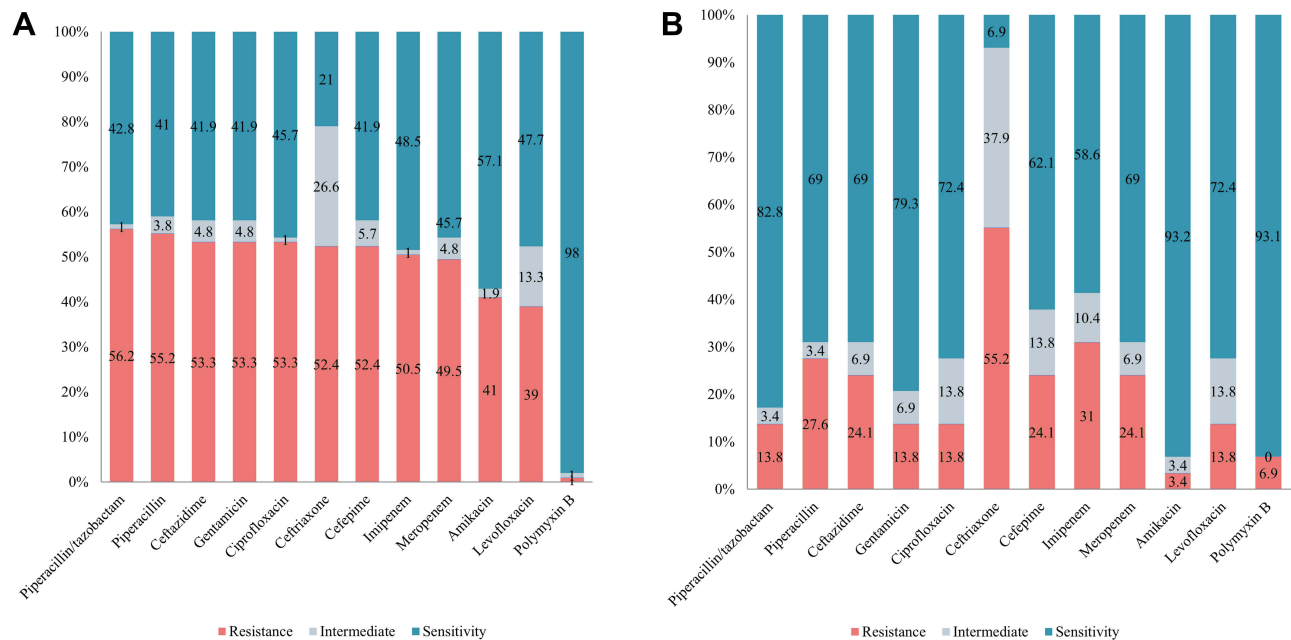
The AST genotypes of *A. baumannii* and *P. aeruginosa* are shown in [Table 1](#). The carbapenemase genotype *bla<sub>OXA</sub>* in *A. baumannii* had the highest proportion, while ESBL-related enzymes exhibited the highest proportion of *bla<sub>TEM</sub>*. For *P. aeruginosa*, the ratio of *bla<sub>KPC</sub>*, ESBL + *OMPK* was greater, and *bla<sub>SHV</sub>* was greater than that of *bla<sub>TEM</sub>*. The aminoglycoside resistance gene *aadA1* accounted for a large proportion. In contrast, the detection rate of *aaaC1* and the quinolone genes *qnrA* and *qnrS* were low. It is noteworthy that *mcr-1* was not detected in *A. baumannii* and *P. aeruginosa* at all.

## Univariate Analysis of NFGNB-Induced PNM

Of the 207 patients included in the final analysis ([Table 2](#)), their mean age was 36.3 years (SD 14.8). Of the patients, 119 (57.4%) were male and 88 (42.6%) were female, with a primary body temperature of 37.8°C (SD 0.9). Univariate analyses revealed that NFGNB-induced PNM patients with TBI ( $P = 0.004$ ), hypertension ( $P = 0.003$ ), ICU admission ( $P = 0.001$ ), EVD ( $P < 0.001$ ) and AMV ( $P < 0.001$ ) were significantly different in the two groups.



**Figure 2** Bacterial distribution of NFGNB-induced PNM (Out ring: Distributions of whole NFGNB enrolled in this study; Middle ring: distribution of NFGNB isolated from the survived PNM patients; Inner ring: distribution of NFGNB isolated from the non-survived PNM patients).



**Figure 3** AST of *A. baumannii* (A) and *P. aeruginosa* (B).

**Table 1** Distribution of Antibiotic Usage and Gene in *A. baumannii* and *P. aeruginosa*

Antibiotic	Gene	<i>A. baumannii</i>	<i>P. aeruginosa</i>
Meropenem	<i>bla<sub>KPC</sub></i>	40.4% (21/52)	28.6% (2/7)
	<i>bla<sub>OXA-66</sub></i>	86.5% (45/52)	14.3% (1/7)
	<i>bla<sub>OXA-23</sub></i>	82.7% (43/52)	14.3% (1/7)
	ESBL + OMPK	42.3% (22/52)	28.6% (2/7)
	<i>bla<sub>NDM</sub></i>	3.8% (2/52)	0.0% (0/7)
	<i>bla<sub>VIM</sub></i>	3.8% (2/52)	14.3% (1/7)
	<i>bla<sub>IMP</sub></i>	1.9% (1/52)	14.3% (1/7)
Ceftriaxone	<i>bla<sub>TEM</sub></i>	69.1% (38/55)	17.6% (3/17)
	<i>bla<sub>SHV</sub></i>	43.6% (24/55)	29.4% (5/17)
	<i>bla<sub>CTX-M-9</sub></i>	18.2% (10/55)	11.8% (2/17)
	<i>bla<sub>CMY</sub></i>	0.0% (0/55)	11.8% (2/17)
	<i>bla<sub>CTX-M-1</sub></i>	3.6% (2/55)	0.0% (0/17)
Amikacin	<i>aadA1</i>	48.8% (21/43)	100.0% (11/11)
	<i>aaaC1</i>	7.0% (3/43)	0.0% (0/1)
Levofloxacin	<i>qnrA</i>	17.1% (7/41)	25.0% (1/4)
	<i>qnrS</i>	2.4% (1/41)	0.0% (0/4)
Polymyxin B	<i>mcr-1</i>	0.0% (0/1)	0.0% (0/2)

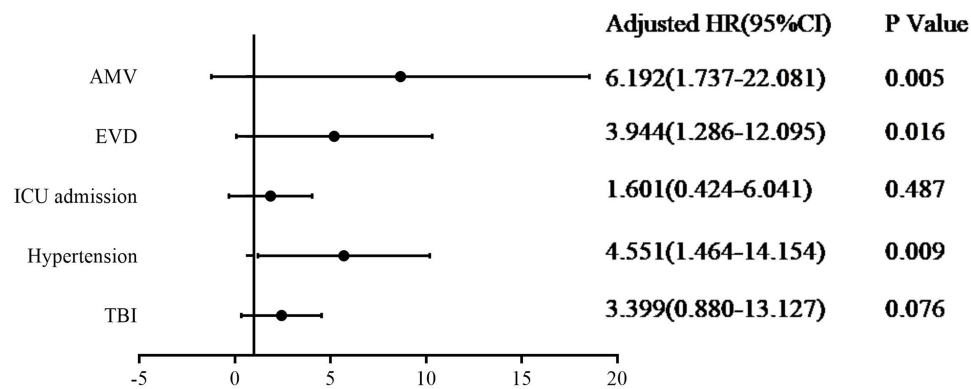
**Table 2** Univariate Analysis of NFGNB-Induced PNM

Characteristics	Total (207)	Survivors (178)	Non-Survivors (29)	P-value
Gender (male)	119 (57.4%)	98 (55.1%)	21 (72.4%)	0.105
Age (years)	36.3 ± 14.8	35.9 ± 15.2	37.9 ± 15.3	0.578
Body temperature (°C)	37.8 ± 0.9	37.8 ± 0.7	37.9 ± 0.8	0.844
Malignant tumor	67 (32.4%)	57 (32.0%)	10 (34.5%)	0.832
TBI	13 (6.3%)	7 (3.9%)	6 (20.7%)	<b>0.004</b>
Diabetes	9 (4.3%)	6 (3.4%)	3 (10.3%)	0.116
Hypertension	25 (12.1%)	16 (9.0%)	9 (31.0%)	<b>0.003</b>
Long operation duration (>3 h)	138 (66.7%)	118 (66.3%)	20 (69.0%)	0.835
Reoperation	63 (30.4%)	50 (28.1%)	13 (44.8%)	0.083
Craniotomy	166 (80.2%)	142 (79.8%)	24 (82.8%)	0.807
Surgical site (head)	202 (97.6%)	173 (97.2%)	29 (100.0%)	0.999
Type of incision (I)	122 (58.9%)	108 (60.7%)	14 (48.3%)	0.224
ICU admission	111 (53.6%)	87 (48.9%)	24 (82.8%)	<b>0.001</b>
CSF leakage	32 (15.5%)	26 (14.6%)	6 (20.7%)	0.409
EVD	100 (48.3%)	77 (43.3%)	23 (79.3%)	<b>&lt;0.001</b>
LD	50 (24.2%)	41 (23%)	9 (31.0%)	0.355
AMV	94 (45.4%)	70 (39.3%)	24 (82.8%)	<b>&lt;0.001</b>

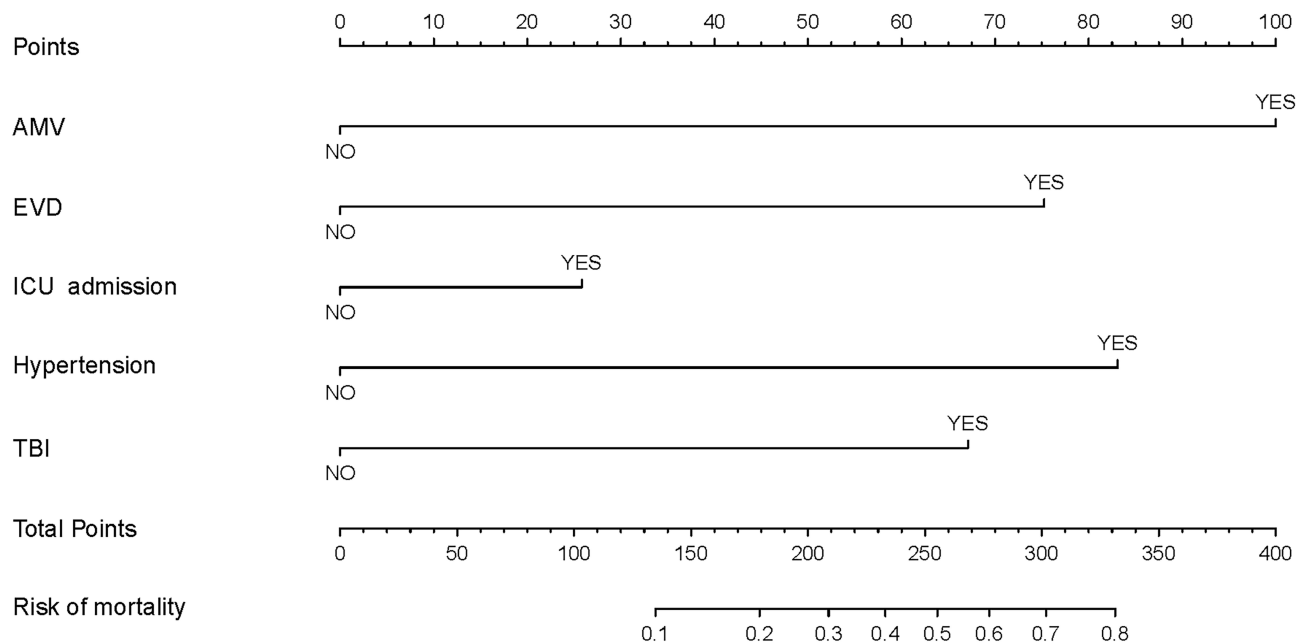
Note: Bold values mean the target parameters is  $p < 0.05$ .

## Multivariate Analysis of NFGNB-Induced PNM

A multivariable logistic model was established for the independent risk factors with statistical significance, as shown in Figure 4. The nomogram prediction model is shown in Figure 5 and [Supplementary Material \(Figures S1 and S2\)](#), either. In NFGNB-induced PNM patients, hypertension (HR 4.551, 95% CI: 1.464–14.154,  $P = 0.009$ ), EVD (HR 3.944, 95% CI: 1.286–12.095,  $P = 0.016$ ) and AMV (HR 6.192, 95% CI: 1.737–22.081,  $P = 0.005$ ) were shown to be independent risk factors for mortality. The verification of the model was carried out as ROC and H-L. The AUC of the model was 0.852, and H-L was 0.193 ( $>0.05$ ) reflecting the fact that the model was of great clinical significance.



**Figure 4** Multivariate analysis of NFGNB-induced PNM.



**Figure 5** Nomogram of the risk of NFGNB-induced PNM mortality.

## Therapy

Of 207 patients, 171(82.6%) received AP, with cefuroxime being the most frequent antibiotic administered. In contrast, 188(90.8%) patients received AET, with high-grade antibiotics usage of 86.7%. A total of 200 patients received ADT, with 90.6% high-grade antibiotics usage. In the present study, between the two groups, a significant difference in the employment of AP was found ( $P = 0.015$ ), but no difference was detected between AET and ADT. Nevertheless, the use rate of ADT and AET in the triple antibiotics regimens was greater in the non-survivors group (Table 3).

## Discussion

Our study highlights the fact that NFGNB-induced PNM patients are at risk of treatment failure and poor outcomes. We first investigated the clinical and microbial epidemiology of NFGNB-induced PNM and explored the factors associated with mortality. In the largest neurosurgery center of China, we conducted this cohort study from 2012 to 2020 and found three factors, which were important predictors of NFGNB-induced PNM, namely hypertension, EVD and AMV. The clinical and microbiological epidemiology of PNM caused by NFGNB was analyzed and the risk factors for death evaluated. The findings have great significance for the diagnosis and treatment of PNM and for preventive interventions

**Table 3** Therapy of NFGNB-Induced PNM

Therapy	Total	Survivors (178)	Non-Survivors (29)	P-value
AP	171 (82.6%)	152 (84.9%)	19 (65.5%)	<b>0.015</b>
AET	188 (90.8%)	161 (90.4%)	27 (93.1%)	0.999
Mono-antibiotic therapy (AET)	49 (26.1%)	44 (27.3%)	5 (18.5%)	0.478
Dual antibiotic therapy (AET)	93 (49.5%)	83 (51.6%)	10 (37.0%)	0.212
Triple or more antibiotic therapy (AET)	46 (24.5%)	34 (21.1%)	12 (44.4%)	<b>0.015</b>
High-grade antibiotic therapy (AET)	163 (86.7%)	140 (87.0%)	23 (85.2%)	0.763
ADT	200 (96.6%)	171 (96.1%)	29 (100.0%)	0.597
Mono-antibiotic therapy (ADT)	33 (16.5%)	31 (18.1%)	2 (6.9%)	0.178
Dual antibiotic therapy (ADT)	88 (44.0%)	78 (45.6%)	10 (34.5%)	0.315
Triple or more antibiotic therapy (ADT)	79 (39.5%)	62 (36.3%)	17 (58.6%)	<b>0.038</b>
High-grade antibiotic therapy (ADT)	155 (90.6%)	29 (100.0%)	184 (92.0%)	0.135

**Note:** Bold values mean the target parameter is  $p < 0.05$ .

for the entire meningitis diseases. In addition, the importance of AP is highlighted in NFGNB-induced PNM, and whether patients experience AP or not can lead to completely different clinical outcomes.

NFGNB are one of the most important groups of pathogens that cause clinical infections, due to their similar metabolism (inability to ferment glucose). In previous reports, *Staphylococcus aureus*, coagulase-negative *Staphylococcus*, and *Enterobacter* accounted for a large proportion of pathogens causing PNM.<sup>11–13</sup> In the past decade, however, PNM caused by NFGNB, especially *A. baumannii* and *P. aeruginosa* has increased year by year with a concomitant increase in drug resistance. According to one report, the carbapenem resistance rate of *A. baumannii* in intensive care units can be as high as 60–70%,<sup>14</sup> and therefore the clinical treatment of patients is extremely difficult. With the widespread use of antibiotics, carbapenem-resistant *A. baumannii* and carbapenem-resistant *P. aeruginosa* have become huge clinical threats. In addition, other NFGNB such as *Burkholderia cepacia* and *Stenotrophomonas maltophilia* possess extensive natural resistance meaning that fewer antibiotics can be used clinically, which also makes treatment problematic.<sup>15,16</sup>

From the perspective of the bacterial distribution, the distribution of the entire NFGNB is relatively concentrated, among which *A. baumannii* and *P. aeruginosa* account for nearly 65%. In the non-survivor group, the two occupied even more of the spectrum, up to 82.8%. It has been reported that the invasiveness of *A. baumannii* and *P. aeruginosa* is higher than that of other NFGNB.<sup>17</sup> *P. aeruginosa* meningitis is much complicated to treat due to the existence of the quorum sensing system and biofilm formation,<sup>18</sup> which can induce drug resistance and lead to an increase in mortality. The carbapenem resistance rate of *P. aeruginosa* can be up to 30%, and the resistance rate of imipenem is even higher than that of meropenem. Resistance to *A. baumannii* in PNM patients is about 50%, therefore carbapenem cannot be widely used for the treatment of NFGNB-induced PNM. In previous studies, polymyxin B, tigecycline and even fosfomycin emerged as effective antibiotics of choice against NFGNB-induced PNM. Nevertheless, these antibiotics have frequently been shown to produce major adverse reactions, such as liver and kidney toxicity, etc, and it has been reported that the employing of loading doses of polymyxin to treat infections caused by NFGNB does not improve clinical outcomes significantly and increases nephrotoxicity.<sup>19,20</sup> Resistance genotypes of NFGNB are different to *Enterobacteriaceae*, where the carbapenem-resistant genotype proportion of *bla*<sub>OXA-66</sub> and *bla*<sub>OXA-23</sub> exhibits the highest proportion in *A. baumannii* and *P. aeruginosa*, and a lower proportion of metalloenzymes (*bla*<sub>VIM</sub>, *bla*<sub>IMP</sub>, *bla*<sub>NDM</sub>). The high proportion of *bla*<sub>TEM</sub> in ESBLs was similar to that reported in previously published literature. However, due to the limited genotypes we used, the genotype coverage for aminoglycosides and quinolones was insufficient.

To better deal with the NFGNB-induced PNM, physicians, pharmacists, and healthcare professionals around the world recognizing the importance of risk factors. As we all know, risk factor assessment is an important measure for predicting the prognosis of a certain type of disease. Physicians and other healthcare professionals can reduce the mortality of meningitis by NFGNB-induced PNM risk factors evaluation, and pharmacists can choose more accurate and effective antibiotics by precious medical procedure, which conducted by risk factor evaluation partly.<sup>21</sup> However, in the treatment of PNM, most previous studies have focused on the risk factors for the occurrence of neurosurgical meningitis, such as EVD, LD, diabetes,

hydrocephalus, Koos grade IV and long operation duration (>3h). An intraoperative blood loss >400 mL is an independent risk factor for PNM for vestibular schwannoma.<sup>22,23</sup> However, few studies have focused on the risk factors for predicting the outcomes of neurosurgical patients, and even fewer have reported mortality risk prediction studies targeting a specific class of pathogens. For example, a study of *Enterobacteriaceae* meningitis predicted patient survival and suggested that a Glasgow Coma Scale (GCS) <8 was a clinical risk factor for death from infection in these patients.<sup>24</sup> To the best of our knowledge, no risk factor evaluation study has been carried out to target NFGNB as a PNM-related pathogen. In the present study, 16 related risk factors for NFGNB-induced PNM were evaluated using multivariate logistic analysis, and it was concluded that hypertension, EVD and AMV were the main risk factors for the poor prognosis of PNM patients. Patients with hypertension usually mean that they have a certain underlying disease and a low immune status, which contributes to poor clinical outcomes.<sup>25</sup> Similar to our study, an Indian study conducted in 2019 showed comorbidities (diabetes,  $p = 0.036$ ; hypertension,  $p = 0.01$ ) were associated with poor outcome of post-neurosurgical *Acinetobacter* meningitis.<sup>26</sup> For patients with EVD, necessary to maintain adequate intracranial pressure or having a CSF shunt, the external persistence exists, making the meningitis difficult to cure.<sup>27</sup> The incidence of EVD infection ranges from 2 to 22%,<sup>28</sup> The best way to reduce infection risk is to remove the EVD as soon as it is no longer needed, thereby decreasing the duration of catheterization which is the most significant risk factor for ventriculitis.<sup>29</sup> The patient needs assisted breathing in the state of PNM, which means that the health of a patient is more serious, and that is one of the risk factors for death.<sup>30</sup> In a Denmark 10 years study target on bacterial meningitis, AMV was calculated as one of the independent mortality risk factors by multivariate analysis.<sup>31</sup> Therefore, for hypertensive patients and those with EVD and AMV, special intensive treatment and prophylaxis care is needed to reduce the mortality rate caused by these insidious infecting pathogens.

Another important discovery in our study was that there was a significant mortality difference in the AP of patients with NFGNB-induced PNM, and the present authors' previous study showed no difference between the two for multidrug-resistant *Enterobacteriaceae* infections. For the non-survivors, the preventive application of antibiotics has great clinical significance. The probable reason may be that, compared with *Enterobacteriaceae*, the prophylactic administration of antibiotics can alleviate the pathogenesis of NFGNB. The pathogenicity of NFGNB is also lower than that of *Enterobacteriaceae* and AP can greatly improve the prognosis of patients. It reports that the combination of antibiotics usage has significant clinical significance in response to NFGNB infections especially in multi-resistance isolates, and when combined with polymyxin and meropenem in treatment of severe carbapenem resistance *A. baumannii* infections,<sup>32</sup> there was a significant decrease in 30-days mortality with no significant difference in nephrotoxicity.<sup>33</sup> Therefore, the combination of polymyxin and meropenem therapy is clinically significant for patients with multi-drug resistant NFGNB infections.<sup>34</sup>

The present study had several limitations. First, it was a retrospective study subject to potential unmeasured confounding factors. Nevertheless to date, it is the largest cohort of patients studied with NFGNB-induced PNM and includes a very extensive list of established prognostic comorbidities, as well as known specific predisposing factors for its development. Second, it was a single-center study, and distributions of antibiotic usage and gene in *A. baumannii* and *P. aeruginosa* of the other hospital were not evaluated. Third, the study did not include all the risk factors related to mortality, such as tumor grading, multiple catheterizations, etc. In our next study, a multicenter cohort will be constructed for mortality assessment, when even more risk factor will be identified.

Overall, this study is the first to analyze the risk factors for mortality after NFGNB infection. NFGNB-induced PNM is a relatively serious clinical challenge worldwide. Hypertension, EVD and AMV were identified as independent mortality risk factors for NFGNB-induced PNM.

## Data Sharing Statement

The original contributions presented in the study are included in the article/[Supplementary Material](#). Further inquiries can be directed to the corresponding authors.

## Ethical Approval and Consent to Participate

The study was approved by the Ethical Committees of Beijing Tiantan Hospital & Capital Medical University (No. KY-2021-079-02) and the study complies with the Declaration of Helsinki. Each patient enrolled in the study signed an informed consent form.

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## Author Contributions

GHZ: design and drafting of the article; data collection and analysis and interpretation. SQW: statistics and genotypes test. HL and GJZ: conduct the whole study.

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## Disclosure

All of the authors declared that no conflict exists in this study.

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