Systematic Review and Meta-analysis on Extended-Spectrum β-lactamases Producing Klebsiella pneumoniae in Nepal

Manita Shyaula*, Christina Khadka*, Prabin Dawadi and Megha Raj Banjara

Central Department of Microbiology, Tribhuvan University, Kathmandu, Bagmati, Nepal.

ABSTRACT

OBJECTIVE: This systematic review and meta-analysis aimed to assess the pool estimates of extended-spectrum β-lactamases producing K. pneumoniae (ESBL-KP) and study their drug resistance profile by evaluating the studies from Nepal.

METHODS: A literature search was carried out in PubMed, Google Scholar, and NepJOL to screen all articles on ESBL-KP published between 2011 and 2021 from Nepal. This review was conducted following Preferred Reporting Items for Systematic Reviews and Meta-Analysis guidelines. Relevant data were extracted, and R language 4.2.0 software was used for statistical analysis.

RESULTS: The pooled prevalence of K. pneumoniae was 5%, while the pooled prevalence of ESBL and multidrug resistance (MDR) in K. pneumoniae were 23% and 55%, respectively. Imipenem was the drug of choice (in vitro) against ESBL-KP infection.

CONCLUSION: Our analyses showed a high prevalence of ESBL-KP and their high resistance toward commonly used drugs. This study highlights the need for the development of new antibiotics for the management of ESBL-KP infections.

KEYWORDS: K. pneumoniae, meta-analysis, ESBL, multi-drug resistance, Nepal

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Introduction

The increasing incidence of antibiotic-resistant bacteria is a concern both to clinicians and patients due to implications such as therapeutic failures, prolonged hospitalization, and nosocomial infections. Extended-spectrum β -lactamase (ESBL) producing bacteria have been reported to be a common cause of healthcare-associated infections, which can have serious clinical consequences and result in multiple drug resistance.^{1,2} ESBL enzymes hydrolyse penicillin, oxyimino-cephalosporins such as third-generation cephalosporins, and monocyclic amide antibiotics such as aztreonam but are inhibited by sulbactam, tazobactam and clavulanic acid.3-5

Enterobacteriaceae is the major family of bacteria associated with ESBL production.⁶ ESBL-producing Enterobacterales were placed under WHO critical priority pathogen list for research, innovation and new drug development.7 In Enterobacteriaceae family, K. pneumoniae is one of the common ESBL-producing members with the potential to cause infections in both hospital and community settings, resulting in increased morbidity and mortality as well as high medical expenses.8

* These authors contributed equally

The initial occurrences of ESBL-KP were reported in Europe in 1983⁹ and the USA in 1989¹⁰ and since then, there has been a continuous increase of K. pneumoniae-mediated cephalosporin-resistance around the world.¹¹⁻¹³ During the late 1980s, ESBL-KP outbreaks were mostly generated by TEM and SHV producers.^{5,14} Recently, the CTX-M enzymesproducers have increased throughout the world.^{15,16} These ESBL encoding genes are commonly found on mobile genetic components, such as plasmids, which frequently carry genes conferring resistance to other classes of antibiotics such as aminoglycosides, fluoroquinolones, chloramphenicol, and trimethoprim/sulfamethoxazole.^{17,18} As a result, many of these bacterial strains are multidrug-resistant (MDR) and infection by these drug-resistant isolates increases the likelihood of treatment failure and may lead to death.8,19,20

South Asia has been identified as a hotspot for antimicrobial-resistant bacteria.^{21,22} The problem has been frequently accounted in developing countries such as Nepal, where the prevalence of MDR and ESBL-producing bacteria has increased over the past years.^{23,24} There are some scattered studies that reported ESBL-KP in Nepal. However, these are not available in an organized form. To the best of our knowledge, this is the first systematic review and meta-analyses on the

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Figure 1. PRISMA flow diagram summarizing the process of literature search and selection.

burden of ESBL-KP in Nepal. This study aimed to estimate the pooled prevalence of ESBL-KP and their antibiotic resistance profile and the relationship between ESBL production and multidrug resistance by analyzing available studies. This study could aid in designing and implementing hospital infection control strategies to minimize the occurrence and spread of ESBL-KP.

Methodology

Literature review and data sources

A comprehensive literature search was conducted to identify all articles published between January 2011 and December 2021 reporting the prevalence of ESBL-KP from human specimens in Nepal. It was carried out through the electronic databases PubMed, Google Scholar, and NepJOL using the following terms: *Klebsiella pneumoniae*', 'extended-spectrum β-lactamases',

and 'ESBL' together with 'Nepal'. Searches were limited to articles published in English language. Discussions and references of the selected studies were thoroughly scanned for locating additional relevant articles. This study followed the PRISMA (Preferred Reporting Items for Systematic Reviews and Meta-Analyses) guidelines for reporting systematic reviews and meta-analyses (Figure 1).

Study selection procedures and eligibility criteria

Two reviewers (MS and CK) separately examined titles and abstracts of articles for possible inclusion in the review. Eligible studies were chosen in 3 steps: first, based upon the title; second, based on the abstract and third, based on the full-text articles. The selected articles were then reviewed and evaluated by other authors (PD and MRB). To be included in the meta-analysis, studies had to meet the following criteria: prevalence of ESBL-KP in Nepal published between 2011 and 2021, ESBL-KP isolated from human specimens and specify laboratory methods for ESBL detection. Studies were excluded for the following reasons: articles reporting ESBL-KP on nonhuman subjects, duplicate studies, undifferentiated *Klebsiella* spp., review articles, case reports, posters, retrospective studies, articles with incomplete information related to ESBL-KP detection method and not falling within the specified time. Studies reporting ESBL-KP only among MDR isolates, and studies from countries other than Nepal were also discarded.

Data extraction

Data were extracted independently by 2 reviewers (MS and CK) from all eligible studies and entered into MS Excel 2007. The extracted data included first author's surname, year of publication, study area, study setting, sample size, total number of multidrug- resistant *K. pneumoniae* (MDR-KP), ESBL diagnostic method, the prevalence of ESBL-KP, antibiogram of ESBL-KP, sample-wise distribution of ESBL-KP and genes variant encoding ESBL. The data was retrieved and analyzed twice to eliminate any possible errors.

Outcome measurements

The main outcome of the study was evaluating the prevalence of ESBL-producers among *K. pneumoniae* isolates in Nepal. The distribution of ESBL-KP according to sample type and their antibiogram was also recorded. Furthermore, the prevalence of MDR-KP and its correlation with ESBL production were documented.

Quality assessment of studies

The quality of individual studies included in the meta-analysis was evaluated independently by 2 reviewers (MS and CK) using a checklist provided by the Newcastle–Ottawa Scale adapted for cross-sectional studies.²⁵ Any differences were checked and resolved through discussion with the third and the fourth authors (PD and MRB). The checklist consists of 10 questions that each reviewer answered separately for each study. Scores varied from 0 to 10, and studies with \geq 6 points were included in the systematic review and meta-analysis (Supplemental Table S1).

Statistical analysis

The statistical analysis was performed using the R programing language 4.2.0 (Meta package). Data were pooled using a random-effect model and the I^2 statistics (measure of inconsistency) was used to assess heterogeneity across studies with I^2 values of 25%, 50%, and 75% indicating low, moderate, and high heterogeneity, respectively.²⁶ In addition, contour-enhanced funnel plots and Egger's test were utilized for detecting possible publication bias. $^{\rm 27}$

Results

Search results

A total of 1949 potentially relevant articles were initially identified, of which 1834 articles were eliminated after screening the titles and abstracts. After excluding duplicate studies, posters, and case studies, 74 articles remained for further full-text assessment. Based on inclusion and exclusion criteria, 41 articles were further removed, and only 45 articles meeting the inclusion criteria were included in the final meta-analysis. In the total 45 included studies, a total of 1927 *K. pneumoniae* were isolated, of which 676 were MDR and 462 produced ESBL. Based on the study setting, the majority of the studies, 97.77% (44/45) were hospital-based, while only 2.22% (1/45) were communitybased. The flowchart of article selection along with the reasons for exclusion are presented in Figure 1, and the main characteristics of the included studies are provided in Table 1.

Laboratory methods used for ESBL detection

This meta-analysis revealed that the majority of the studies conducted in Nepal used only phenotypic tests for ESBL detection. Altogether, 3 different phenotypic tests were used for ESBL confirmation: combined disk test (CDT), double disk synergy test (DDST), and epsilometric test (E-test). Out of 45 studies included in this meta-analysis, 37 (82.22%) used CDT alone for ESBL-confirmation, 2 (4.44%) used DDST only and one (2.22%) study performed only E-test to estimate ESBL-proportions among *K. pneumoniae*. Similarly, 4 (8.88%) studies used both DDST and CDT, while 1 (2.22%) study performed CDT, DDST and E-test in combination for the confirmation of ESBL-phenotypes. Only 2 (4.44%) studies carried out molecular tests for the detection of ESBL-encoding genes (Table 1).

Province-wise distribution of articles on ESBL-KP in Nepal

The majority of the studies on ESBL-KP were reported from Bagmati Province (93.33%), while there were limited studies from Madhesh Province and Province No. 1 (2.22% in Madhesh Province and 4.44% in Province No. 1). No studies were identified on this subject from other provinces. The number of studies on ESBL-KP was few until 2014. The highest number of articles investigating this topic were published in the year 2017 (Figure 2).

Meta-analysis on the prevalence of K. pneumoniae

Based on the analysis of the eligible studies, the pooled prevalence of *K. pneumoniae* was found 5% (95% confidence interval (CI), 0.03-0.07) among processed samples with significant heterogeneity (P = 96%; P < .01) (Figure 3).

 Table 1. Characteristics of studies included in the meta-analysis.

Sherchan et al ¹⁰ Hospital Bagmati 120 22 5 (0.23) 5 (0.23) CDT	STUDY ID	STUDY SETTING	PROVINCE	TOTAL SAMPLE SIZE	K. PNEUMONIAE	NO. OF MDR (%)	NO. OF ESBL (%)	DIAGNOSTIC METHODS	ESBL GENES DETECTED
Sherchan et al ²⁰ Hospital Bagmati 120 22 5 (0.23) 5 (0.23) CDT	Shrestha et al ²⁸	Hospital	1	_	32	_	8 (0.25)	CDT	_
Shreathara Shreathara Hospital Bagmati 167 42 31 (0.74) 12 (0.29) CDT _ Bhandari et al ¹⁹¹ Hospital Bagmati 224 22 18 (0.82) 6 (0.27) CDT _ Mishra et al ¹⁹² Hospital Bagmati 1120 102 _ 43 (0.42) CDT _ Pathak and Hospital Bagmati 1258 20 _ 7 (0.35) CDT _ Singh et al ¹⁹³ Hospital Bagmati 300 26 _ 4 (0.15) DDST _ Chaudhary Hospital Bagmati 1057 17 7 (0.41) 3 (0.18) CDT _ Parajuli et al ¹⁴¹ Hospital Bagmati 1005 4 _ 1 (0.25) CDT _ Shrestha et al ⁴⁰⁴ Hospital Bagmati 1005 8 _ 2 (0.25) CDT _ Shrestha et al ⁴⁰⁴ Hospital Bagmati 1005 8 QDT </td <td>Sherchan et al²⁹</td> <td>Hospital</td> <td>Bagmati</td> <td></td> <td>22</td> <td></td> <td>5 (0.23)</td> <td>CDT</td> <td>_</td>	Sherchan et al ²⁹	Hospital	Bagmati		22		5 (0.23)	CDT	_
Bhandari et al ¹⁴ Hospital Bagmati 294 22 18 (0.82) 6 (0.27) CDT	Chander and Shrestha ³⁰	Hospital	Bagmati	6308	145	-	24 (0.17)	CDT	-
Mishra et al ¹²³ Hospital Bagmati 1120 102	Khanal et al ³¹	Hospital	Bagmati	187	42	31 (0.74)	12 (0.29)	CDT	_
Pathak and Pathak and Pokharel ^{P41} Hospital Bagmati 1416 25 11 (0.44) 10 (0.40) CDT	Bhandari et al32	Hospital	Bagmati	294	22	18 (0.82)	6 (0.27)	CDT	_
Pokharel ³⁴ Province	Mishra et al ³³	Hospital	Bagmati	1120	102	_	43 (0.42)		_
Tapa et al ¹⁹⁴ Hospital Bagmati 300 26	Pathak and Pokharel ³⁴	Hospital	Bagmati	1416	25	11 (0.44)	10 (0.40)	CDT	_
Chaudhary et al ³⁷ Hospital Bagmati 1986 38 _ 7 (0.18) CDT _ Parajuli et al ⁸⁴ Hospital Bagmati _ 11 9 (0.82) 4 (0.36) CDT _ Shrestha et al ⁹⁹ Hospital Bagmati 1057 17 7 (0.41) 3 (0.18) CDT _ Shrestha et al ⁴⁴⁰ Hospital Bagmati 500 4 _ 2 (0.25) CDT _ Bhandari et al ⁴⁴¹ Hospital Bagmati 5245 109 60 (0.55) 47 (0.43) CDT _ Nepal et al ⁴⁴² Hospital Bagmati 5245 109 60 (0.55) 47 (0.43) CDT _ Nepal et al ⁴⁴⁴ Hospital Bagmati 5245 109 60 (0.55) 47 (0.43) CDT _ Parajuli et al ⁴⁴⁴ Hospital Bagmati 3088 7 5 (0.71) 3 (0.43) CDT _ Rai et al ⁴⁷⁹ Hospital Bagmati 1018 24	Singh et al35	Hospital	Bagmati	1258	20	_	7 (0.35)	CDT	_
et als7 for	Thapa et al ³⁶	Hospital	Bagmati	300	26	-	4 (0.15)	DDST	_
Shrestha et al ¹⁹ Hospital Bagmati 1057 17 7 (0.41) 3 (0.18) CDT	Chaudhary et al ³⁷	Hospital	Bagmati	1986	38	_	7 (0.18)	CDT	_
Shrestha et al ⁴⁰ Hospital Bagmati 500 4	Parajuli et al38	Hospital	Bagmati	_	11	9 (0.82)	4 (0.36)	CDT	_
Bandari et al ⁴⁴ Hospital Bagmati 1003 8	Shrestha et al39	Hospital	Bagmati	1057	17	7 (0.41)	3 (0.18)	CDT	_
Ghimire et al ⁴² Hospital Bagmati 5245 109 60 (0.55) 47 (0.43) CDT	Shrestha et al40	Hospital	Bagmati	500	4	_	1 (0.25)	CDT	_
Nepal et al ⁴³ Hospital Bagmati 1568 39 23 (0.59) 15 (0.38) CDT	Bhandari et al41	Hospital	Bagmati	1003	8	_	2 (0.25)	CDT	_
Nepal et al ⁴⁴ Hospital Bagmati	Ghimire et al42	Hospital	Bagmati	5245	109	60 (0.55)	47 (0.43)	CDT	_
Parajuli et al ⁴⁵ Hospital Bagmati 3088 7 5 (0.71) 3 (0.43) CDT _ Pathak et al ⁴⁶ Hospital Bagmati 977 83 29 (0.35) 13 (0.16) CDT _ Rai et al ⁴⁷ Hospital Bagmati 0.18 26 _ 2 (0.08) CDT _ Rimal et al ⁴⁴ Hospital Bagmati 1018 24 6 (0.25) 3 (0.13) CDT _ Sah et al ⁴⁴ Hospital Bagmati 1018 24 6 (0.25) 3 (0.18) CDT _ Shakya et al ⁵⁰ Hospital Bagmati 2209 17 8 (0.47) 3 (0.18) CDT _ Maharjan et al ⁵¹ Community Bagmati 510 30 2 (0.07) 2 (0.07) CDT, E-test and PCR Pla_TEM Upreti et al ⁵² Hospital Bagmati 182 6 5 (0.83) 2 (0.33) DDST _ Guragain et al ⁵⁴ Hospital Bagmati 180	Nepal et al43	Hospital	Bagmati	1568	39	23 (0.59)	15 (0.38)	CDT	_
Pathak et al ⁴⁶ Hospital Bagmati 977 83 29 (0.35) 13 (0.16) CDT	Nepal et al44	Hospital	Bagmati	-	60	39 (0.65)	26 (0.43)		-
Rai et al ⁴⁷ Hospital Bagmati 26 2 (0.08) CDT Rimal et al ⁴⁸ Hospital Bagmati 1018 24 6 (0.25) 3 (0.13) CDT Sah et al ⁴⁹ Hospital Madhesh 1602 63 58 (0.92) 22 (0.35) CDT Shakya et al ⁵⁰ Hospital Bagmati 2209 17 8 (0.47) 3 (0.18) CDT	Parajuli et al45	Hospital	Bagmati	3088	7	5 (0.71)	3 (0.43)	CDT	_
Rimal et al ⁴⁸ Hospital Bagmati 1018 24 6 (0.25) 3 (0.13) CDT	Pathak et al46	Hospital	Bagmati	977	83	29 (0.35)	13 (0.16)	CDT	_
Sah et al ⁴⁹ Hospital Madhesh 1602 63 58 (0.92) 22 (0.35) CDT	Rai et al47	Hospital	Bagmati	_	26	_	2 (0.08)	CDT	_
Shakya et al ⁵⁰ Hospital Bagmati 2209 17 8 (0.47) 3 (0.18) CDT	Rimal et al48	Hospital	Bagmati	1018	24	6 (0.25)	3 (0.13)	CDT	_
Maharjan et al ⁵¹ Community Bagmati 510 30 2 (0.07) 2 (0.07) CDT, E-test and PCR bla _{TEM} Upreti et al ⁵² Hospital Bagmati 182 6 5 (0.83) 2 (0.33) DDST Mahato et al ⁵³ Hospital 1 3666 27 17 (0.63) 1 (0.04) CDT Guragain et al ⁵⁴ Hospital Bagmati 300 10 4 (0.40) 3 (0.30) CDT Rooku et al ⁵⁵ Hospital Bagmati 100 9 5 (0.56) 4 (0.44) CDT	Sah et al49	Hospital	Madhesh	1602	63	58 (0.92)	22 (0.35)	CDT	_
Upreti et al ⁵² Hospital Bagmati 182 6 5 (0.83) 2 (0.33) DDST	Shakya et al⁵⁰	Hospital	Bagmati	2209	17	8 (0.47)	3 (0.18)	CDT	_
Mahato et al ⁵³ Hospital 1 3666 27 17 (0.63) 1 (0.04) CDT	Maharjan et al ⁵¹	Community	Bagmati	510	30	2 (0.07)	2 (0.07)		bla _{TEM}
Guragain et al ⁵⁴ Hospital Bagmati 300 10 4 (0.40) 3 (0.30) CDT	Upreti et al52	Hospital	Bagmati	182	6	5 (0.83)	2 (0.33)	DDST	_
Rooku et alHospitalBagmati10095 (0.56)4 (0.44)CDTLohani et alHospitalBagmati55041 6 (0.15)CDT and PCR bla_{CTX-M} , bla_{TEM} , $bla_{SH'}$ Maharjan et alHospitalBagmati6047166 (0.93)25 (0.35)CDTMahaseth et alHospitalBagmati556422325 (0.11)CDTAdhikari et alHospitalBagmati10554 (0.802 (0.40)CDTGhimire et alHospitalBagmati3652718 (0.67)3 (0.11)CDT	Mahato et al53	Hospital	1	3666	27	17 (0.63)	1 (0.04)	CDT	_
Lohani et al ⁵⁶ HospitalBagmati55041 $ 6$ (0.15) CDT and PCR bla_{CTX-M} , bla_{TEM} , bla_{SH} Maharjan et al Mahaseth et al Adhikari et al Ghimire et al 600Bagmati6047166 (0.93)25 (0.35)CDT $-$ Mahaseth et al 500HospitalBagmati5564223 $-$ 25 (0.11)CDT $-$ Adhikari et al 500HospitalBagmati10554 (0.802 (0.40)CDT $-$ Ghimire et al 600HospitalBagmati3652718 (0.67)3 (0.11)CDT $-$	Guragain et al54	Hospital	Bagmati	300	10	4 (0.40)	3 (0.30)	CDT	_
Maharjan et al ⁵⁷ Hospital Bagmati 604 71 66 (0.93) 25 (0.35) CDT Mahaseth et al ⁵⁸ Hospital Bagmati 5564 223 25 (0.11) CDT Adhikari et al ⁵⁹ Hospital Bagmati 105 5 4 (0.80 2 (0.40) CDT Ghimire et al ⁶⁰ Hospital Bagmati 365 27 18 (0.67) 3 (0.11) CDT	Rooku et al55	Hospital	Bagmati	100	9	5 (0.56)	4 (0.44)	CDT	_
Mahaseth et al ⁵⁸ Hospital Bagmati 5564 223 _ 25 (0.11) CDT _ Adhikari et al ⁵⁹ Hospital Bagmati 105 5 4 (0.80 2 (0.40) CDT _ Ghimire et al ⁶⁰ Hospital Bagmati 365 27 18 (0.67) 3 (0.11) CDT _	Lohani et al ⁵⁶	Hospital	Bagmati	550	41	_	6 (0.15)		bla _{CTX-M} , bla _{TEM} , bla _{SHV}
Adhikari et al ⁵⁹ Hospital Bagmati 105 5 4 (0.80 2 (0.40) CDT	Maharjan et al57	Hospital	Bagmati	604	71	66 (0.93)	25 (0.35)	CDT	_
Ghimire et al ⁶⁰ Hospital Bagmati 365 27 18 (0.67) 3 (0.11) CDT	Mahaseth et al58	Hospital	Bagmati	5564	223	_	25 (0.11)	CDT	_
	Adhikari et al59	Hospital	Bagmati	105	5	4 (0.80	2 (0.40)	CDT	_
Kayastha et al ⁶¹ Hospital Bagmati 1443 18 16 (0.89) 6 (0.33) CDT _	Ghimire et al60	Hospital	Bagmati	365	27	18 (0.67)	3 (0.11)	CDT	
	Kayastha et al61	Hospital	Bagmati	1443	18	16 (0.89)	6 (0.33)	CDT	_

(Continued)

Table 1. (Continued)

STUDY ID	STUDY SETTING	PROVINCE	TOTAL SAMPLE SIZE	K. PNEUMONIAE	NO. OF MDR (%)	NO. OF ESBL (%)	DIAGNOSTIC METHODS	ESBL GENES DETECTED
Mandal et al62	Hospital	Bagmati	190	50	_	13 (0.26)	CDT	_
Raya et al63	Hospital	Bagmati	5545	30	11 (0.37)	7 (0.23)	CDT	_
Regmi et al64	Hospital	Bagmati	150	13	8 (0.62)	4 (0.31)	CDT	_
Subedi et al65	Hospital	Bagmati	388	18	10 (0.56)	8 (0.44)	DDST and CDT	-
Shilpakar et al66	Hospital	Bagmati	770	35	31 (0.89)	3 (0.09)	E test	_
Adhikari et al67	Hospital	Bagmati	149	36	29 (0.81)	11 (0.31)	CDT	_
Pandey et al68	Hospital	Bagmati	8756	149	48 (0.32)	24 (0.16)	CDT, DDST, E test	-
Karki et al69	Hospital	Bagmati	1155	76	24 (0.32)	2 (0.03)	CDT	_
Kuinkel et al ⁷⁰	Hospital	Bagmati	2197	40	26 (0.65)	7 (0.18)	CDT	_
Karki et al ⁷¹	Hospital	Bagmati	3216	41	20 (0.49)	11 (0.27)	CDT	_
Nepal et al ²³	Hospital	Bagmati	380	30	23 (0.77)	23 (0.77)	CDT	_



Meta-analysis on the prevalence of MDR among K. pneumoniae isolates

The overall pooled prevalence of MDR isolates among *K. pneumoniae* was 55% (95% CI, 0.47-0.64), with a high level of heterogeneity ($I^2 = 81\%$, P < .01) among the studies (Figure 4).

Meta-analysis on the prevalence of ESBL-KP in Nepal

The pooled prevalence of ESBL-KP was 23% (95% CI, 0.19-0.27), with a high heterogeneity (P = 76%, P < .01) among the studies analysed (Figure 5).

Distribution of ESBL-KP according to the type of samples

Sample-wise distribution study revealed blood as the most common sample from which ESBL-phenotypes were recovered. Blood samples accounted for 62.22% of the ESBL-KP isolates, followed by wound aspirates (40%), sputum (34.54%), bronchoalveolar lavage (30.55%), and tracheal aspirates (28.57%). One-fourth of the ESBL-positive isolates were obtained from pus samples. The rate of the isolation from urine and stool specimens were 16.86% and 18.75%, respectively (Table 2).

	Church		<u>е</u> г	Standardised Mean	CMD	05% 01	Maisht.
	Study	g	SE	Difference	SMD	95%-CI	weight
	Parajuli et al 2017	0.00	0.0009		0.00	[0.00; 0.00]	2.6%
	Raya et al 2020		0.0010	· · · · · · · · · · · · · · · · · · ·	0.01	[0.00; 0.01]	2.6%
	Mahato et al 2018	0.01	0.0014	+	0.01	[0.00; 0.01]	2.6%
	Shakya et al 2017		0.0019	+	0.01	[0.00; 0.01]	2.6%
	Bhandari et al 2016		0.0028	+	0.01	[0.00; 0.01]	2.6%
	Shrestha et al 2016b		0.0040	+	0.01	[0.00; 0.02]	2.6%
	Kayastha et al 2020		0.0029	+	0.01	[0.01; 0.02]	2.6%
	Karki et al 2021b		0.0020	+	0.01	[0.01; 0.02]	2.6%
	Singh et al 2015		0.0036	+	0.02	[0.01; 0.02]	2.6%
	Shrestha et al 2016a		0.0039	+	0.02	[0.01; 0.02]	2.6%
	Pandey et al 2021		0.0014	<u>•</u>	0.02	[0.01; 0.02]	2.6%
	Pathak and Pokharel 2015		0.0035	+	0.02	[0.01; 0.02]	2.6%
	Kuinkel et al 2021		0.0029	+	0.02	[0.01; 0.02]	2.6%
	Chaudhary et al 2016		0.0031	+	0.02	[0.01; 0.03]	2.6%
	Ghimire et al 2016		0.0020	*	0.02	[0.02; 0.02]	2.6%
	Chander and Shrestha 2013			*	0.02	[0.02; 0.03]	2.6%
	Rimal et al 2017		0.0048	+	0.02	[0.01; 0.03]	2.6%
	Nepal et al 2017a		0.0040	+	0.02	[0.02; 0.03]	2.6%
	Upreti et al 2018		0.0135		0.03	[0.01; 0.06]	2.4%
	Guragain et al 2019		0.0105		0.03	[0.01; 0.05]	2.5%
	Sah et al 2017		0.0050	<u>+</u>	0.04	[0.03; 0.05]	2.6%
	Mahaseth et al 2019		0.0027		0.04	[0.03; 0.05]	2.6%
	Shilpakar et al 2021		0.0077		0.05	[0.03; 0.06]	2.6%
	Subedi et al 2020		0.0109		0.05	[0.02; 0.07]	2.5%
	Adhikari et al 2020		0.0213		0.05	[0.01; 0.09]	2.2%
	Maharjan et al 2018		0.0107		0.06	[0.04; 0.08]	2.5%
	Karki et al 2021a		0.0075		0.07	[0.05; 0.08]	2.6%
	Ghimire et al 2020		0.0142		0.07	[0.05; 0.10]	2.4%
	Lohani et al 2019 Bhandari et al 2015		0.0116		0.07	[0.05; 0.10]	2.5%
	Nepal et al 2021		0.0160 0.0144		0.07 0.08	[0.04; 0.11] [0.05; 0.11]	2.4% 2.4%
	Pathak et al 2017		0.0093			[0.05, 0.11]	2.4%
	Thapa et al 2015		0.0093		0.08	[0.07; 0.10]	2.3%
	Regmi et al 2020		0.0240		0.09	[0.03; 0.12]	2.3%
	KC et al 2019		0.0240			[0.03; 0.15]	1.9%
	Mishra et al 2015		0.0090			[0.07; 0.11]	2.5%
	Maharjan et al 2019		0.0000		0.12	[0.09; 0.14]	2.4%
	Sherchan et al 2012		0.0391		0.12	[0.11; 0.26]	1.6%
	Khanal et al 2013		0.0347		0.22	[0.16; 0.29]	1.7%
	Adhikari et al 2021		0.0403		0.24	[0.16; 0.32]	1.6%
	Mandal et al 2020		0.0372		0.26	[0.19; 0.34]	1.7%
	Random effects model			\$	0.05	[0.03; 0.07]	100.0%
	Prediction interval					[-0.05; 0.15]	
	Heterogeneity: $I^2 = 96\%$, $p < 0$.	01	_	0.3-0.2-0.1 0 0.1 0.2 0.3			
e 3.	Forest plot depicting the pooled prev	alence					

Correlation between MDR and ESBL production among K. pneumoniae isolates

Figure

A strong positive correlation (Pearson's correlation coefficient of 0.82; 95% CI: .67-.91) was observed between ESBL production and multidrug resistance in *K. pneumoniae* strains (Figure 6 and Supplemental Table S5).

Antimicrobial resistance patterns in ESBL-KP

The antibiotic resistance pattern (in vitro) of ESBL-KP is shown in Table 3. Imipenem was found to be the most effective antibiotic (99.9% sensitive) against ESBL-producing strains of *K. pneumoniae*, followed by amikacin (92.7% sensitive) and nitrofurantoin (68.03% sensitive). The resistance toward

Study	g	SE	Standardised Mean Difference S	SMD	95%-CI	Weight
Maharjan et al 2018	0.07	0.0471		0.07	[-0.03; 0.16]	4.5%
Sherchan et al 2012	0.23	0.1016		0.23	[0.03; 0.43]	3.8%
Rimal et al 2017	0.25	0.1021		0.25	[0.05; 0.45]	3.8%
Karki et al 2021a	0.32	0.0645		0.32	[0.19; 0.44]	4.3%
Pandey et al 2021	0.32	0.0465		0.32	[0.23; 0.41]	4.5%
Pathak et al 2017	0.35	0.0649		0.35	[0.22; 0.48]	4.3%
Raya et al 2020	0.37	0.1106	- +	0.37	[0.15; 0.58]	3.6%
Guragain et al 2019	0.40	0.2000	· · · ·	0.40	[0.01; 0.79]	2.4%
Shrestha et al 2016a	0.41	0.1556		0.41	[0.11; 0.72]	3.0%
Pathak and Pokharel 2015	0.44	0.1327	— <u>—</u>	0.44	[0.18; 0.70]	3.3%
Shakya et al 2017	0.47	0.1664	— <u> </u>	0.47	[0.14; 0.80]	2.8%
Karki et al 2021b	0.49	0.1091		0.49	[0.27; 0.70]	3.6%
Ghimire et al 2016	0.55	0.0711	<u> </u>	0.55	[0.41; 0.69]	4.2%
KC et al 2019	0.56	0.2485		0.56	[0.07; 1.04]	1.9%
Subedi et al 2020	0.56	0.1757		0.56	[0.21; 0.90]	2.7%
Nepal et al 2017a	0.59	0.1230		0.59	[0.35; 0.83]	3.4%
Regmi et al 2020	0.62	0.2176		0.62	[0.19; 1.04]	2.2%
Mahato et al 2018	0.63	0.1527		0.63	[0.33; 0.93]	3.0%
Nepal et al 2017b	0.65	0.1041		0.65	[0.45; 0.85]	3.7%
Kuinkel et al 2021	0.65	0.1275	- <u></u> -	0.65	[0.40; 0.90]	3.4%
Ghimire et al 2020	0.67	0.1571	— • —	0.67	[0.36; 0.97]	2.9%
Parajuli et al 2017	0.71	0.3194		0.71	[0.09; 1.34]	1.4%
Khanal et al 2013	0.74	0.1326		0.74	[0.48; 1.00]	3.3%
Nepal et al 2021	0.77	0.1599		0.77	[0.45; 1.08]	2.9%
Adhikari et al 2020	0.80	0.4000		0.80	[0.02; 1.58]	1.0%
Adhikari et al 2021	0.81	0.1496		0.81	[0.51; 1.10]	3.0%
Bhandari et al 2015	0.82	0.1928		0.82	[0.44; 1.20]	2.5%
Parajuli et al 2016	0.82	0.2727		0.82	[0.28; 1.35]	1.7%
Upreti et al 2018	0.83	0.3727		0.83	[0.10; 1.56]	1.1%
Shilpakar et al 2021	0.89	0.1591		0.89	[0.57; 1.20]	2.9%
Kayastha et al 2020	0.89	0.2222		0.89	[0.45; 1.32]	2.1%
Sah et al 2017	0.92	0.1209			[0.68; 1.16]	3.5%
Maharjan et al 2019	0.93	0.1144		0.93	[0.71; 1.15]	3.6%
Random effects model			↓	0.55	[0.47; 0.64]	100.0%
Prediction interval					[0.13; 0.98]	
Heterogeneity: I^2 = 81%, p <	0.01	_	1.5 -1 -0.5 0 0.5 1 1.5			
re 4. Forest plot of pooled prevalence c	of multid					

Figure 4. Forest plot of pooled prevalence of multidrug resistance among K. pneumoniae in Nepal.

ampicillin was very high (99.9%). Higher resistance was also observed against trimethoprim-sulfamethoxazole (60.22%), ofloxacin (56.69%), and gentamicin (50.9%).

Publication Bias

Quality assessment of all the included studies performed using Newcastle-Ottawa Scale showed the total score ≥ 6 indicating satisfactory and fair selection of articles. The funnel plots for accessing the bias for studies reporting *K. pneumoniae*, ESBL production and MDR profile in *K. pneumoniae* showed the majority of the studies are statistically significant for the inclusion. However, on performing Egger's tests for these parameters, all plots were found asymmetrical (P < .001) indicating publication bias (Supplemental Table S2, S3 and S4). Based on the contour-enhanced funnel plots and quality assessments, the analyses were considered suitable for the publication (Supplemental Figure S1, S2 and S3).

Discussion

Drug-resistant strains of *K. pneumoniae* are becoming a serious concern worldwide. This meta-analysis shows the overall pooled prevalence of *K. pneumoniae* in various processed specimens as 5% in Nepal. However, Odari and Dawadi⁷² reported the prevalence of *K. pneumoniae* in clinical specimens in the context of Nepal as 3%. The variation could be the result of difference in time of the study. Odari and

Study	g	SE	Standardised Mean Difference	SMD	95%-CI	Weig
Karki et al 2021a	0.03	0.0186		0.03	[-0.01; 0.06]	3.7
Mahato et al 2018		0.0370			[-0.04; 0.11]	3.4
Maharjan et al 2018		0.0471			[-0.03; 0.16]	3.2
Rai et al 2017		0.0544			[-0.03; 0.18]	3.1
Shilpakar et al 2021		0.0495			[-0.01; 0.18]	3.2
Ghimire et al 2020		0.0642			[-0.01; 0.10]	2.9
Mahaseth et al 2019		0.0224			[0.07; 0.24]	3.7
Rimal et al 2017		0.0722			[-0.02; 0.27]	2.
Lohani et al 2019		0.0597			[0.02; 0.27]	2. 3.0
Thapa et al 2015		0.0769			[0.00; 0.20]	2.0
Pathak et al 2017		0.0434			[0.07; 0.24]	3.3
Pandey et al 2021		0.0329			[0.10; 0.24]	3.
Chander and Shrestha 2013					[0.10; 0.23]	3.
Kuinkel et al 2021		0.0338			[0.05; 0.30]	2.8
					[-0.02; 0.30]	2.0
Shrestha et al 2016a		0.1019				
Shakya et al 2017		0.1019			[-0.02; 0.38]	2.0
Chaudhary et al 2016		0.0696			[0.05; 0.32]	2.
Sherchan et al 2012		0.1016			[0.03; 0.43]	2.
Raya et al 2020		0.0882			[0.06; 0.41]	2.
Shrestha et al 2011		0.0884			[0.08; 0.42]	2.
Shrestha et al 2016b		0.2500			[-0.24; 0.74]	0.0
Bhandari et al 2016		0.1768			[-0.10; 0.60]	1.
Mandal et al 2020		0.0721			[0.12; 0.40]	2.
Karki et al 2021b		0.0809			[0.11; 0.43]	2.
Bhandari et al 2015		0.1113			[0.05; 0.49]	1.9
Khanal et al 2013		0.0825			[0.12; 0.45]	2.4
Guragain et al 2019		0.1732			[-0.04; 0.64]	1.
Adhikari et al 2021		0.0921			[0.12; 0.49]	2.2
Regmi et al 2020		0.1538	-		[0.01; 0.61]	1.
Upreti et al 2018		0.2357			[-0.13; 0.80]	0.
Kayastha et al 2020		0.1361			[0.07; 0.60]	1.
Sah et al 2017		0.0745			[0.20; 0.50]	2.0
Singh et al 2015		0.1323			[0.09; 0.61]	1.
Maharjan et al 2019		0.0704			[0.21; 0.49]	2.
Parajuli et al 2016		0.1818			[0.01; 0.72]	1.0
Nepal et al 2017a		0.0993			[0.19; 0.58]	2.
Pathak and Pokharel 2015		0.1265			[0.15; 0.65]	1.0
Adhikari et al 2020		0.2828			[-0.15; 0.95]	0.
Mishra et al 2015		0.0643			[0.30; 0.55]	2.8
Parajuli et al 2017		0.2474			[-0.06; 0.91]	0.
Ghimire et al 2016		0.0629			[0.31; 0.55]	2.9
Nepal et al 2017b		0.0850			[0.27; 0.60]	2.4
KC et al 2019		0.2222			[0.01; 0.88]	0.
Subedi et al 2020		0.1571			[0.14; 0.75]	1.:
Nepal et al 2021	0.77	0.1599		0.77	[0.45; 1.08]	1.2
Random effects model				0.23	[0.19; 0.27]	100.
Prediction interval					[0.01; 0.45]	
Heterogeneity: $I^2 = 76\%$, $p < 0$.	.01					

Table 2. Sample-wise distribution of ESBL-KP isolates.

S.N.	SAMPLE TYPES	K. PNEUMONIAE	ESBL K. PNEUMONIAE	OCCURRENCE OF ESBL-KP (%)
1.	Urine	635	107	16.86
2.	Sputum	55	19	34.54
3.	Pus	43	11	25.58
4.	Blood	45	28	62.22
5.	Stool	80	15	18.75
6.	Tracheal aspirates	42	12	28.57
7.	Bronchoalveolar lavage	36	11	30.55
8.	Wound aspirates	5	2	40.00



Dawadi carried out the analyses of studies on *K. pneumoniae* from 2015 to 2021^{72} while our studies included data between 2011 and 2021.

Management of antibiotic resistance in Gram-negative bacteria, mainly MDR-KP, currently represents a significant challenge in the field of infectious diseases.⁷³ In our study, the pooled prevalence of MDR isolates in *K. pneumoniae* was estimated to be 55%, which is in agreement with those of the studies conducted in Bangladesh (55%),⁷⁴ Indonesia (54.49%),⁷⁵ and Iran (58%).⁷⁶ On the contrary, our pooled prevalence is lower than those of the studies from Pakistan (63%),⁷⁷ Brazil (84%)⁷⁸ and India (75%)⁷⁹ but much higher than study from Ethiopia (20%).⁸⁰

Our study showed that the prevalence of MDR-KP was higher in Bagmati Province. The majority of megacities is located in the province and contains dense populations. The high resistance may be linked with antibiotic prescription patterns, environmental exposures, hygiene practices, study periods, poor health, and medical facilities.^{21,72,80} A strong positive correlation was observed between ESBL production and multidrug resistance. This might be due to the fact that ESBL genes in combination with other resistance genes confer such resistance.^{8,81}

K. pneumoniae is a leading cause of opportunistic healthcare-associated infections, which are increasingly complicated by the production of ESBL-enzymes by these bacteria.^{12,82} In the present study, the pooled prevalence of ESBL-KP was 23%. Our result is in accordance with studies conducted in Bangladesh (25.335%)⁸³ and Sri Lanka (25%),⁸⁴ but higher in comparison to the study reported from Pakistan (15.25%).⁸⁵ There are many factors that contribute to a high frequency of ESBLstrains, including acquisition of novel resistance-genes, transfer of resistance contributing genes, lack of effective antibiotic surveillance systems, inadequate infection control strategies, easy availability and misuse of antibiotics, inappropriate dosing schedule, lack of new antibiotics and limited diagnostic facilities.⁸⁰ These factors could have contributed to the higher occurrence of ESBL-KP in Nepal.

ANTIBIOTICS	RESISTANCE (POOLED	95% CI	TEST OF HE	TEROGENEITY	NUMBER OF STUDIES	
	ESTIMATION)		l² (%)	P-VALUE	REVIEWED	
Amikacin	7.30	[–0.0100; 0.1559]	47.9	0.073	7	
Gentamicin	50.90	[0.2796; 0.7383]	0.0	0.587	4	
Ampicillin	99.90	[0.9990; 0.9990]	0.0	1.00	3	
Imipenem	0.10	[0.0010; 0.0010]	0.0	1.00	5	
Nitrofurantoin	31.97	[-0.0004; 0.6398]	40.3	0.170	4	
Ofloxacin	56.69	[0.2249; 0.9090]	48.3	0.102	5	
Trimethoprim-sulfamethoxazole	60.22	[0.4825; 0.7218]	0.0	0.922	6	

Table 3. Antibiotic resistance profile of ESBL-KP.

The study on antibiogram of ESBL-KP revealed a very high level of resistance to ampicillin (99.90%) which is consistent with studies conducted in Bangladesh,74 Saudi Arabia86 and Indonesia.87 The increase in resistance toward these antibiotics may be due to the inappropriate use of antibiotics without medical supervision and with improper administration, as well as insufficient therapy.88 The majority of ESBL-KP were found sensitive to imipenem (99.90%) and amikacin (92.7%). These are in parallel with the findings of studies conducted in Pakistan (imipenem 100%, amikacin 95.2%),89 Indonesia (imipenem 91.5%, amikacin 90.4%)87 and France (imipenem 99.3%, amikacin 93.2%).90 The higher susceptibility of ESBL-KP toward imipenem may be due to the finite use of this antibiotic in Nepal. Besides ESBL production, factors such as biofilm formation and production of other antibiotic hydrolysing enzymes were found to be responsible for emergence in the resistance in Enterobacteriaceae.91,92

In this study, ESBL-KP strains were predominantly recovered from blood (62.22%). Sarojamma and Ramakrishna also reported blood as a major source of ESBL-producers.⁹³ This could be due to the factors like inappropriate use of syringes, poor hand washing practices among healthcare professionals and improper disinfection of skin of prolonged hospital-stayed patients during transfusion.^{94,95} However, reports from other investigations show higher prevalence from other human samples.^{90,96}

Strengths and limitations of our study

This is the first study to calculate the pooled prevalence of ESBL-KP in Nepal. Studies available from all geographical regions of the country, and studies on both hospital and community-based settings were included in the meta-analysis. Our exploration also revealed the pooled estimate of MDR-KP and their correlation with ESBL production, distribution of ESBL-KP in different specimens, and the resistance profile of ESBL-KP to commonly administered antibiotics. The findings from our study could be of use to clinicians and health policymakers. However, this review has some limitations. First, the

study on potential risk factors associated with ESBL-infections such as socioeconomic status, prior antibiotic use, and past medical history of patients was not possible to accumulate due to limited prior research on these factors. Second, the prevalence of ESBL-KP in non-human subjects was not studied due to low number of articles addressing the topic. Third, most studies on ESBL-KP were from hospital-setting, and the precise ratio of nosocomial versus community-acquired bacteria was not determined. Fourth, it was also impossible to determine the mortality rate associated with infections caused by ESBL-KP as no studies were found reporting the consequence. Additionally, investigation on ESBL-genotypes is limited in Nepal. Thus, the pooled prevalence of infections is primarily based on studies on ESBL-phenotype.

The increase in the occurrence of MDR and ESBLproducing strains necessitates an improvement in laboratory facilities as well as laboratory experts' diagnostic skills for assessing antibiotic resistance profiles. The scarcity of studies from provinces other than Bagmati indicates the need for research on this topic from all parts of the country. Our study revealed that the majority of studies on ESBL-KP were from hospital-settings, while studies from community-settings on ESBL-strains were rare. Thus, community-based studies on ESBL strains are recommended. Further research is also required to evaluate the risk factors that might affect the development and implementation of a rational and effective ESBL-KP control strategy. Routine screening of ESBL-strains, molecular approaches for the detection of ESBL-genotypes and proper antibiotic prescription policies are urgently required to combat the burden of ESBL-KP.

Conclusion

Overall, high ESBL production was seen in *K. pneumoniae* which could contribute to multidrug resistance. A decreased susceptibility of ESBL-KP toward the majority of commonly used antibiotics was seen, thereby highlighting the need for rationale use of appropriate antibiotics. The better options for the treatment of ESBL-KP could be imipenem and amikacin.

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Author's Contributions

MS conceived the idea, and MS and CK designed the study. MS and CK reviewed the literature and extracted the data. PD and MRB rechecked the data. PD analysed the data. MS, CK and PD drafted the manuscript. MRB revised and reviewed the manuscript. MRB supervised the overall data analysis and writing. All authors read and approved the final manuscript.

Data Availability

All supplementary files, data generated and analysed will be made available by the corresponding author upon reasonable request.

Supplemental Material

Supplemental material for this article is available online.

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