


Bacteriology of community-acquired pneumonia, antimicrobial susceptibility pattern and associated risk factors among HIV patients, Northeast Ethiopia: cross-sectional study

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Abstract

Objective: Pneumonia is an opportunistic infection and it is a major cause of mortality and morbidity among human immunodeficiency virus/acquired immune deficiency syndrome-positive patients. Previous studies have shown the dominant pathogens bacterial isolates were *K. pneumoniae* 27.0%, *S. aureus* 20.8%, *S. pneumoniae* 18.8% and *E. coli* 8.3%. This study aimed to determine bacteriology of community-acquired pneumonia, antimicrobial susceptibility pattern and associated risk factors among human immunodeficiency virus patients in the Northeast Ethiopia: cross-sectional study.

Methods: A health facility-based cross-sectional study was conducted from January to April 2021 at six health facilities in Dessie Town. A total of 378 community-acquired pneumonia patients suspected to be human immunodeficiency virus-positive were recruited using a consecutive sampling technique. Sociodemographic and clinical data were collected using a structured questionnaire. A two-milliliter sputum specimen was collected aseptically from each study participant. Samples were cultivated on blood agar, chocolate agar and MacConkey agar to isolate bacterial pathogens. To identify bacteria pathogens Gram stain, colony morphology and biochemical tests were performed. The Kirby-Bauer Disc Diffusion method was used to perform the antimicrobial susceptibility test. Descriptive statistics, logistic regression analysis was carried out using Statistical package for social science version 25 software. p -value < 0.05 with a corresponding 95% confidence interval (CI) was considered for statistical significance.

Result: The overall prevalence of bacterial pneumonia was 175 (46.3%). Gram-negative bacteria accounted for 119 (68%) and the predominant isolates identified were *Streptococcus pneumoniae* 49 (28%) followed by *Klebsiella pneumoniae* 46 (26.3%), *Pseudomonas aeruginosa* 34 (19.4%). There were 148 (84.6%) multidrug-resistant bacteria overall. Statistically significant factors included viral load, cigarette smoking, cluster of differentiation 4 count, alcohol use, World Health Organization clinical stages III and IV and low white blood cell count.

Conclusion: The study found that both multidrug resistance and bacterial pneumonia were high. Thus, bacterial culture and antimicrobial susceptibility tests should be routinely performed in health facilities in order to prevent and control the spread of bacterial infection and concurrent drug resistance.

Keywords

pneumonia, bacterial infection, human immunodeficiency virus, Ethiopia

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Introduction

Pneumonia is one of the most common pulmonary complications in human immunodeficiency virus (HIV)-positive patients.¹ As HIV influences both the humoral and cellular components of innate immunity; these alterations are specific and are evident even with relatively preserved cluster of differentiation 4 (CD4⁺) T cell counts and undetectable viral loads. Impaired innate immune function, particularly in the setting of CD4⁺ T cell depletion, may contribute to the pathogenesis of opportunistic lung infections.²

Pneumonia is caused by various groups of micro-organisms. In fact, the most common etiological agents are bacteria.³ Bacterial pneumonia is well known to occur at an elevated incidence in individuals living with HIV.⁴ Patients with acquired immune deficiency syndrome (AIDS) and HIV infection at the beginning of the HIV outbreak have a documented rise in the rate of bacterial infection.⁵ HIV patients are more vulnerable to opportunistic and bacterial infections due to alteration of host T-cell function, impaired phagocytic response of neutrophils and macrophages and the inability of B-cells to produce specific and increasing antibodies.⁶ Infection of pulmonary macrophages and lymphocytes with HIV-1 plays a key role in pulmonary disease pathogenesis in AIDS.⁷

Pneumonia is the fourth most common cause of death worldwide, after ischemic heart disease, stroke, and chronic obstructive pulmonary disease, and it is the second most common reason for years of life lost.⁸ It is more frequent in HIV patients, being 5–15 times more common in them than in HIV negative individuals.⁹

Pulmonary complications are a major cause of morbidity and mortality among HIV-infected patients globally.¹⁰ The incidence of bacterial disease among HIV-infected individuals has been substantially reduced by the use of antiretroviral therapy in both developed and developing countries.¹¹ However, even in patients with antiretroviral treatment (ART) and high CD4⁺ counts, bacterial pneumonia remains high.¹² The incidence of bacterial pneumonia among HIV-infected people increased with poor HIV/AIDS control, most commonly manifested by increased viral load and lower CD4 count.¹³

The most common bacterial pathogens that cause bacterial pneumonia are *Streptococcus pneumoniae*, *Haemophilus influenzae*, *Klebsiella pneumoniae*, *Pseudomonas aeruginosa*, *Escherichia coli* and *Staphylococcus aureus*.¹⁴ In Spain, a prospective observational study showed that *S. pneumoniae* was the leading species in causing bacterial pneumonia, accounted for 31.7%.¹¹ Another study revealed that from 161 HIV-infected individuals, bacterial pneumonia was 17%. In addition, *S. Pneumoniae* was the most predominant bacteria (30%), followed by *H. influenzae* and *S. aureus*, accounted for 2%.¹⁵ Moreover, bacterial pneumonia was predominant in HIV-positive patients, accounted for 54%.¹⁶ The

most common bacterial isolates were *K. pneumoniae* 27.0%, *S. aureus* 20.8%, *S. pneumoniae* 18.8%, *E. coli* 8.3%, *H. influenzae*, *Klebsiella oxytoca*, *P. aeruginosa*, 4.2% each, *Enterobacter* spp. 2.1% and unidentified Gram-negative bacteria 10.4%.¹⁷

Inappropriate use of antimicrobials provides selective pressure that favors the emergence of resistant strains in pathogenic bacteria. The majority of bacterial isolates found in people infected with HIV have varying levels of antimicrobial resistance.¹⁸ Multidrug resistance is defined as a bacterium which is non-susceptible to at least one agent in three or more antimicrobial categories.^{19,20}

In the study area, the most commonly prescribed drugs for pneumonia were amoxicillin, amoxicillin-clavulanic acid, erythromycin, gentamicin and ceftriaxone without antimicrobial susceptibility testing (AST). This empirical method of treatment favors the emergence of drug-resistant bacterial strains. The emergence of drug-resistant bacterial strains poses difficulty in the management of pneumonia among HIV-positive patients. Therefore, this study was aimed to assess bacteriology of community-acquired pneumonia (CAP), antimicrobial susceptibility pattern and associated risk factors among HIV patients in the Northeast Ethiopia: cross-sectional study.

Method and materials

Study setting and design

A health facility-based cross-sectional study was conducted from January to April 2021 at the ART clinics of six health facilities in Dessie town, Northeast Ethiopia. These health facilities were Dessie Comprehensive Specialized Hospital, Selam General Hospital, Borumeda General Hospital, Dessie Health Centre, Ethio General Hospital and Bati General Hospital.

Inclusion and exclusion criteria. All HIV-positive patients from selected health facilities who were enrolled in the ART clinic and showed clinical signs and symptoms of pneumonia, such as shortness of breath, chest pain, fever, chills, fatigue, and cough, during data collection were included. However, patients who were critically ill and unable to expectorate sputum were excluded.

Population. All individuals living with HIV and who were enrolled at the ART clinics of health facilities in Dessie town were the source of population. Whereas HIV patients who were enrolled in the ART clinic of selected health facilities developed clinical evidence of CAP and were available in the ART clinic during the study period.

Sample size and sampling technique. The sample size was determined using a single population proportion formula considering (43.7%) taken from a previous study conducted

in patients suspected to have pneumonia in Mekelle, Northern Ethiopia,²¹ marginal error of 5% and 95% confidence interval (CI)=1.96 by using the following formula:

$$n = \frac{(Z_{\alpha/2})^2 * p * (1 - p)}{d^2}$$

where n =the minimum sample size required, $Z_{\alpha/2}$ =the significant value for 95% CI, P =the expected prevalence of pneumonia infection, and d =the margin of error, finally a total of 378 study participants with clinically diagnosed CAP (CD-CAP) were selected using a systematic random sampling. HIV-positive patients who received antibiotics within the past 2 weeks except cotrimoxazole were excluded.

[LE]

Operational definitions

Hand washing habit. Either washing hands with soap and water or killing germs on the hands with an at least 60% alcohol-based hand sanitizer or rub. When you clean your hands, you remove many germs.²²

Smoking habit. Smoker consuming between 11–19 cigarettes and above 20 cigarettes or more per day.²³

ART failure. ART failure is defined as progression of the disease and high risk of mortality after the beginning of ART.²⁴

Co-infection. Co-infection is the simultaneous infection of a host by multiple pathogen species, for instance multi-parasite infections.²⁵

Body mass index. Body fat based on height and weight that applies to adult men and women.²⁶

The normal range for a white blood cell count is typically between 4000 and 11,000/microliters. Factors like age and sex may determine what a normal range type of WBC: normal percentage of overall lymphocyte: 20% to 40%, neutrophil: 55% to 70%, eosinophil: 1% to 4%, monocyte: 2% to 8% and basophil: 0% to 2%.²⁷

Specimen collection and processing

In total, 2 ml sputum specimens were collected using clean, dry, sterile, wide-necked and leak-proof containers from each study participant. Sputum samples were transported immediately to Wollo University Microbiology Laboratory for analysis. Sputum samples were stored at 4°C if a specimen processing delay existed.²⁸ Blood samples were also collected for viral load, CD4⁺ cell and white blood cell count during data collection.

Gene Xpert test was performed for mycobacterium analyses.²⁹ Gram staining with more than 25 polymorphic

nuclear leukocytes and <10 epithelial cells were considered as good and cultivated. However, specimens with more than 10 epithelial cells and less than 25 polymorphic nuclear leukocytes per high power field (100X) were not good and discarded.²⁸

A sterilized loop of specimens were streaked onto Blood agar (HiMedia™), Chocolate agar and MacConkey agar (HiMedia™). The Chocolate agar was incubated in a candle jar at 37°C for 24–48 h. Whereas, Blood Agar and MacConkey agar was aerobically incubated for 24 h at 37°C.^{30,31} Positive growth on Blood agar and MacConkey agar (HiMedia™) was subculture onto Nutrient agar (HiMedia™) for biochemical and antimicrobial susceptibility test.

Bacterial isolation and identification

The bacteria isolate was characterized using colony morphology, haemolysis pattern, Gram staining, and biochemical tests following the standard microbiological procedure. The significant bacterial count was carried out and the significant bacterial count was reported on the basis of observing an excess of 10⁵ CFU/ml.²⁸ Gram-positive cocci were distinguished and recognized based on Gram stain, blood agar haemolysis patterns, colonial characteristics, catalase test, coagulase test, mannitol fermentation test, and optochin (5 µg) susceptibility. Gram-negative bacteria were identified based on Gram reaction, colony morphology and pigmentation, oxidase test, fermentation of carbohydrates, H₂S production, motility, formation of indole, triple sugar iron agar and citrate utilization, lysine decarboxylase or methyl red vogues Proskur utilization, urea hydrolysis and satellite tests.

Antimicrobial susceptibility testing. The isolated organism was tested against antibiotic agents using the Kirby Bauer disc diffusion method on Muller Hilton agar (HMEDIA). Briefly, single pure colonies of isolated species from nutrient agar were picked and transferred to a tube containing 5 ml tryptone-soya broth and mixed to make a homogenous suspension, then incubated at 37°C until the turbidity of the suspension was matched to a 0.5 McFarland standard. Using sterile swab, the inoculum suspension was inoculated over the entire surface of the Mueller Hinton agar plate.³² After application of the selected antimicrobial disks, the plate was incubated overnight at 37°C for 16–18 h. Inhibition zone diameter was measured and the degree of susceptibility was interpreted to each antibiotic according to Clinical and laboratory standards institute (CLSI) guideline.³³ Antibacterial agents were selected based on local prescription habit and CLSI recommendations.

The standard antibiotic discs (Liofilchem-Italy, HARDY Diagnosis-Santa Maria, USA) and its concentrations used as: penicillin (10 µg), chloramphenicol (30 µg), ciprofloxacin (5 µg), clindamycin (30 µg), cefoxitin (30 µg), trimethoprim-sulfamethoxazole (1.25/23.75 µg), cefotaxime (30 µg), ceftriaxone (30 µg), erythromycin (15 µg) and oxacillin (30 µg) are the antimicrobial agents that can be used by Gram-positive

bacteria. Chloramphenicol (30 µg), ciprofloxacin (5 µg), tetracycline (30 µg), gentamicin (10 µg), trimethoprim-sulfamethoxazole (1.25/23.75 µg), ceftriaxone (30 µg), piperacillin-tazobactam (100/10 µg), ceftazidime (30 µg), amikacin (30 µg), ampicillin (10 µg), amoxicillin-clavulanic acid (20/10 µg), meropenem (10 µg) and amoxicillin (10 µg) can be used for Gram-negative bacteria.^{33,34}

Diameters of zones of inhibitions were measured using digital caliper. The interpretation of results of antimicrobial susceptibility tests were based on the standardized table supplied by CLSI³³ as sensitive, intermediate or resistant. Moreover, an isolate was considered Multidrug resistance (MDR) if it is resistant to at least one agent in three or more antimicrobial categories.³⁵

Quality control. Training was given for data collectors; the completeness of the questionnaires was checked by the principal investigator, and about 5% of the questionnaire was pretested at Meseret specialized clinic. Quality control of culture media was verified for sterility testing by overnight incubation of 5% of uninoculated plates/tubes of the prepared media from each batch. Standard reference strains of *S. aureus* American type culture collection (ATCC-25,923), *E. coli* (ATCC-25,922) and *P. aeruginosa* (ATCC-27,853) and for fastidious organisms, *H. influenzae* (ATCC 49,247) and *S. pneumoniae* (ATCC 49,619) were used as control strains.

Statistical analysis. All the data were entered using Epi-data version 4.6.0.4 and exported to Statistical package for social science version 25 for analysis. Descriptive statistics were performed. Bivariate and multivariate analyses were done. Variables with a *p*-value less or equal to 0.25 in bivariable analysis were subjected to multivariable analysis. Adjusted odd ratio with *p* value of <0.05 with 95% CI was statistically significant. Results were presented using graphs and tables.

Ethical considerations. Ethical approval was obtained from the ethical review committee of the College of Medicine and Health Sciences, Wollo University with a protocol number of CMHS/HC/354/13. Written informed consent was obtained from the subjects and the legally authorized representatives of the minor subjects prior to study initiation. Confidentiality and any special data security requirements were maintained and assured. All data and sample collected were kept confidential and used only for the purpose of the study. The positive cases were communicated with their physicians in order to initiate their treatment and management accordingly.

Result

Sociodemographic characteristics

The mean age of the study participants was 40.09 (±12.24) years, ranges from 10 to 70 years. Half (50.5%) of the study

Table 1. Sociodemographic characteristics of pneumonia-suspected HIV-positive study participants (*n* = 378) at Dessie town health facilities, Northeast Ethiopia, 2021.

Demographic and clinical variables	Frequency	Percentage (%)
Sex		
Male	187	49.5
Female	191	50.5
Age in years		
10–16	10	2.6
17–30	77	20.4
31–45	174	46
>45	117	31
Residence		
Rural	89	23.5
Urban	289	76.5
Education status		
Unable to read and write	104	27.5
Only read and write	37	9.8
Primary school completed	100	26.5
Secondary school completed	64	16.9
College and above	73	19.3
Occupation		
Employed	82	21.7
Student	27	6.3
Merchant	97	25.7
House wife	28	7.4
Farmer	16	4.2
Daily laborer	131	34.7
Marital status		
Single	25	6.6
Married	318	84.1
Divorced	18	4.8
Widowed	17	4.5
No of family members		
One	19	5.0
Two	41	10.8
Three	90	23.8
Four	108	28.6
Five	63	16.7
>Six	57	15.1

participants were females and majority of the study participants were urban dwellers (76.5%). In addition, 84% and 35% of participants were married and daily worker, respectively. Two hundred fifty-eight (68.3%) participants were living in a family having less than five members (Table 1).

Clinical and behavioural characteristics

Over 378 participants, 258 (68.4%) were non-smokers, 275 (72.8%) were never drink alcohol and 299 (79.1%) were chewed chat. Among 378 participants, 168 (44.4%), 183 (48.4%) and 137 (36.2%) participants were on World Health Organization (WHO) stage-II, viral load and CD4 count of

Table 2. Behavioural, clinical and health-related characteristics of pneumonia suspected HIV-positive study participants (n = 378) at Dessie town health facilities, Northeast Ethiopia, 2021.

Variables	Category	Frequency	Percentage (%)
Hand washing habit	Yes	366	(96.8)
	No	12	(3.2)
Cigarette smoking	Yes	120	(31.6)
	No	258	(68.4)
Alcohol consumption	Always	23	(6.1)
	Some times	80	(21.2)
	Never	275	(78.7)
Chat chewing	Always	26	(6.9)
	Some times	53	(14)
	Never	299	(79.1)
Current WHO clinical stage	Stage-I	127	(33.6)
	Stage-II	168	(44.4)
	Stage-III	67	(17.7)
	Stage-IV	16	(4.2)
CD4 count	0—200	51	(13.5)
	201—350	137	(36.2)
	351—500	99	(26.2)
	>500	91	(24.1)
White blood cell counts	Low	185	(48.9)
	Normal	170	(45.0)
	High	23	(6.1)
Viral load count	>1000 copies/cell	180	(47.6)
	150–1000 copies/cell	15	(4.0)
	<150 copies/cell	183	(48.4)
History of hospitalization in the last 6 months	Yes	43	(11.4)
	No	335	(88.6)
Prophylaxis TMP.SMX	Yes	193	(51.1)
	No	185	(48.9)
Co-infection	Yes	62	(16.4)
	No	318	(83.6)
ART failure	Yes	49	(13.0)
	No	329	(87.0)
Current BMI	Under weight	108	(28.6)
	Normal	237	(62.7)
	Over weight	33	(8.7)
ART treatment duration	7–12 months	3	(0.8)
	1–5 years	72	(19)
	>5 years	303	(80.2)

WHO=World Health Organization; ART=antiretroviral treatment; BMI=body mass index; CD4=cluster of differentiation 4.

<150 copies/cell and 201–350 cells/mm³, respectively. Moreover, half of the study participants had a white blood cell count of less than 4000 cells/mm³ (Table 2).

Prevalence of bacterial isolates

The overall prevalence of bacterial infection was 46.3% (95% CI: 41.3%–51.3%), and almost equal prevalence rate was observed among male and female participants. Of the total 175 isolates, Gram-negative bacteria accounted for 119 (68%). Overall, the predominant isolates identified were *S.*

pneumoniae 49 (28%) followed by *K. pneumoniae* 46 (26.3%), *P. aeruginosa* 34 (19.4%). Whereas, the least identified bacteria were *Acinetobacter* species and *Proteus vulgaris* 2 (1.1%) each (Figure 1).

Antimicrobial susceptibility profile of bacterial isolates

Gram-positive bacteria showed high levels of resistance to tetracycline 49 (87.5%), penicillin 48 (85.7%), trimethoprim-sulfamethoxazole 34 (78.6%) and chloramphenicol 37

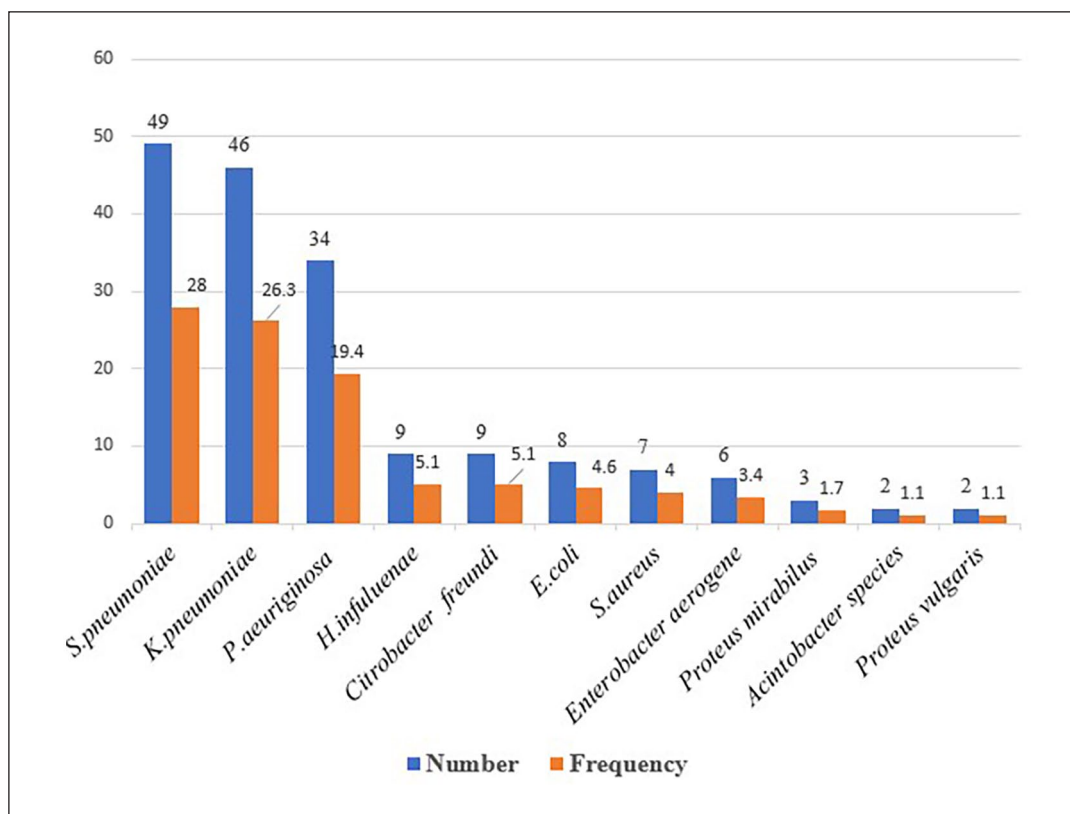


Figure 1. Frequency of bacteria isolated from pneumonia suspected HIV-positive patients at Dessie town health facilities, Northeast Ethiopia, 2021.

Table 3. Antimicrobial resistance and susceptibility pattern of Gram-positive bacteria isolated from pneumonia-suspected HIV-positive patients at Dessie town health facilities, Northeast Ethiopia, 2021.

Gram-positive bacterial isolates	Antimicrobial agents tested	Antimicrobial agents tested										
		E N (%)	FOX N (%)	C N (%)	CXT N (%)	CRO N (%)	CIP N (%)	SXT N (%)	CL N (%)	OX N (%)	TE N (%)	P N (%)
<i>S. pneumoniae</i> (49)	S	21 (42.9)	NT	13 (26.5)	26 (53.1)	26 (53.1)	NT	10 (20.4)	36 (73.5)	37 (75.5)	6 (12.4)	7 (6.3)
	R	28 (57.1)	NT	36 (73.5)	23 (46.9)	23 (46.9)	NT	39 (79.6)	13 (27.3)	12 (24.5)	43 (87.6)	41 (83.7)
<i>S. aureus</i> (7)	S	3 (42.9)	3 (42.9)	3 (42.9)	3 (42.9)	4 (57.1)	4 (51.1)	2 (28.6)	4 (57.1)	5 (71.4)	1 (14.3)	1 (14.3)
	R	4 (57.1)	4 (57.1)	4 (57.1)	4 (57.1)	3 (42.9)	3 (42.9)	5 (71.4)	3 (42.9)	2 (28.6)	6 (85.7)	6 (85.7)
Total (56)	S	24 (42.9)	3 (42.9)	19 (33.9)	29 (51.8)	30 (53.6)	4 (51.1)	12 (21.4)	40 (71.4)	42 (75)	7 (12.5)	8 (14.3)
	R	32 (57.1)	4 (57.1)	37 (66.1)	27 (48.2)	29 (46.3)	3 (42.9)	34 (78.6)	16 (28.6)	12 (25)	49 (87.5)	48 (85.7)

CL=Clindamycin; E=Erythromycin; C=Chloramphenicol; CIP=Ciprofloxacin; TE=Tetracycline; SXT=Trimethoprim-Sulfamethoxazole; CRO=Ceftriaxone; P=penicillin; R=Resistant; OX=oxacillin; FOX=Cefoxitin; S=Sensitive; NT=Not tested.

(66.1%). whereas 40 (71.4%) and 42 (75%) of the Gram-positive isolates were sensitive to clindamycin and oxacillin, respectively. Moreover, 57% of *S. aureus* isolates also showed resistance to methicillin (Table 3).

Majority of the Gram-negative isolates showed resistance rate of 67 (88.2%) for tetracycline, 65 (88.2%) for Ampicillin and 65 (87.8%) for amoxicillin-clavulanic acid. Rates of resistance of Gram-negative bacterial isolates against ceftazidime, tetracycline, trimethoprim-sulfamethoxazole, chloramphenicol, amoxicillin-clavulanic acid, cefotaxime,

amikacin, ceftriaxone, meropenem and gentamicin ranged from 45 (37.8%)–67 (88.2%). However, Gram-negative bacterial isolates showed relatively low resistance against amikacin 45 (37.8) and meropenem 47 (39.5) (Table 4).

MDR patterns of the isolates

Overall, 173 (98.9%) bacterial isolates were resistant to at least one antimicrobial agent and 167 (95.4%) isolates were resistant to ≥ 2 antimicrobials. About 25 (143%) isolates had

Table 4. Antimicrobial resistance and susceptibility pattern of Gram-negative bacteria among pneumonia-suspected HIV-positive patients at Dessie town health facilities ART clinic, Northeast Ethiopia, 2021.

	Antimicrobials tested													
	C	TE	CIP	CXT	SXT	GN	AMP	AMC	CRO	CAZ	AMK	MEM	TZP	
	N (%)	N (%)	N (%)	N (%)	N (%)	N (%)	N (%)	N (%)	N (%)	N (%)	N (%)	N (%)	N (%)	
<i>K. pneumoniae</i> (46)	S 13 (33.3)	4 (8.7)	28 (60.9)	25 (54.3)	9 (19.6)	31 (63.4)	5 (10.9)	6 (13)	25 (54.3)	21 (45.7)	32 (69.6)	32 (69.6)	NT	
	R 36 (66.7)	42 (91.3)	18 (39.1)	21 (45.7)	37 (80.4)	15 (36.6)	41 (89.1)	40 (87)	21 (45.7)	25 (54.3)	13 (30.4)	13 (30.4)	NT	
<i>P. aeruginosa</i> (34)	S NT	NT	13 (38.2)	15 (44.1)	NT	14 (41.2)	NT	NT	15 (44.1)	12 (35.3)	19 (55.9)	20 (58.8)	13 (38.2)	
	R NT	NT	21 (61.8)	19 (55.9)	NT	20 (58.8)	NT	NT	19 (55.9)	22 (64.7)	15 (44.1)	14 (41.2)	21 (61.8)	
<i>C. freundii</i> (9)	S 3 (33.6)	1 (11.1)	6 (66.7)	5 (55.6)	2 (23.3)	5 (55.6)	1 (11.1)	NT	4 (44.4)	3 (33.3)	5 (55.6)	4 (66.7)	NT	
	R 6 (66.4)	8 (89.9)	3 (33.3)	4 (44.4)	7 (77.7)	4 (44.4)	8 (88.9)	NT	5 (55.6)	6 (66.7)	4 (44.4)	3 (33.3)	NT	
<i>H. influenzae</i> (9)	S 1 (11.1)	2 (22.2)	5 (56.6)	6 (66.7)	3 (33.3)	5 (56.6)	1 (11.1)	1 (11.1)	5 (55.6)	4 (44.4)	5 (56.6)	5 (55.6)	NT	
	R 8 (88.9)	7 (77.8)	4 (44.4)	3 (33.3)	6 (66.7)	4 (44.4)	8 (88.9)	8 (88.9)	4 (44.4)	5 (55.6)	4 (44.4)	4 (44.4)	NT	
<i>E. coli</i> (8)	S 3 (37.5)	1 (12.5)	6 (75)	5 (62.5)	3 (37.5)	6 (75)	1 (12.5)	1 (12.5)	5 (62.5)	4 (50)	5 (62.5)	5 (62.5)	NT	
	R 5 (62.5)	7 (87.5)	2 (28.5)	3 (37.5)	5 (62.5)	2 (25)	7 (87.5)	7 (87.5)	3 (37.5)	4 (50)	3 (37.5)	3 (37.5)	NT	
<i>Enterobacter aerogenes</i> (6)	S 1 (16.9)	1 (16.7)	4 (66.7)	3 (50)	2 (33.3)	4 (66.7)	1 (16.9)	1 (16.5)	4 (66.7)	3 (50)	4 (66.7)	3 (50)	NT	
	R 5 (83.1)	5 (83.3)	2 (33.3)	3 (50)	4 (66.7)	2 (33.3)	5 (83.1)	5 (83.5)	2 (33.3)	3 (50)	2 (33.3)	3 (50)	NT	
<i>P. mirabilis</i> (3)	S 1 (33.3)	NT	1 (33.3)	4 (66.7)	0 (0)	0 (0)	0 (0)	0 (0)	2 (66.7)	NT	2 (66.7)	1 (33.3)	NT	
	R 2 (66.7)	NT	2 (66.7)	1 (33.3)	3 (100)	3 (100)	3 (100)	3 (1000)	1 (33.3)	NT	1 (33.3)	2 (66.7)	NT	
<i>P. vulgaris</i> (2)	S 1 (50)	NT	1 (50)	1 (100)	0 (0)	1 (50)	0 (0)	0 (0)	2 (100)	NT	1 (50)	1 (50)	NT	
	R 1 (50)	NT	1 (50)	0 (100)	2 (100)	1 (50)	2 (100)	2 (100)	0 (0)	NT	1 (50)	1 (50)	NT	
<i>Acinetobacter</i> species (2)	S NT	NT	0 (0)	NT	NT	0 (0)	NT	NT	NT	0 (0)	1 (50)	1 (50)	0 (0)	
	R NT	NT	2 (100)	NT	NT	2 (100)	NT	NT	NT	2 (100)	1 (50)	1 (50)	2 (100)	
Total (119)	S 23 (27.7)	9 (11.8)	64 (53.8)	64 (54.7)	19 (22.9)	66 (55.5)	9 (11.8)	9 (12.2)	62 (53)	47 (41.2)	74 (62.2)	72 (60.5)	13 (36.1)	
	R 60 (72.3)	67 (88.2)	55 (46.2)	53 (45.3)	64 (77.1)	53 (44.5)	65 (88.2)	65 (87.8)	55 (47)	67 (58.8)	45 (37.8)	47 (39.5)	23 (63.9)	

NT = Note tested; AMP = Ampicillin; GN = Gentamicin; AMK = Amikacin; CIP = Ciprofloxacin (5 µg); SXT = trimethoprim-sulfamethoxazole; MEM = Meropenem; AMC = Amoxicillin-clavulanic acid; CTX = cefotaxime; CAZ = Cefazidime; CRO = ceftriaxone; TE = Tetracycline (30 µg); C = Chloramphenicol; TZP = Piperacillin tazobactam; R = Resistant; S = Sensitive.

Table 5. Multidrug resistance profile of bacterial isolates (n = 175) from pneumonia-suspected HIV-positive participants at Dessie town health facilities, Northeast Ethiopia, 2021.

Isolated bacteria	Antimicrobial resistance pattern						MDR; n (%)
	R ₀ ; n (%)	R ₁ ; n (%)	R ₂ ; n (%)	R ₃ ; n (%)	R ₄ ; n (%)	≥R ₅ ; n (%)	
Gram positive	0 (0)	2 (3.6)	7 (12.5)	28 (50)	16 (28.6)	3 (5.4)	47 (83.9)
<i>S. pneumoniae</i>	0 (0)	2 (4.8)	5 (30.6)	25 (28.6)	14 (20.4)	3 (6.1)	42 (85.7)
<i>S. aureus</i>	0 (0)	0 (0)	2 (28.5)	3 (43)	2 (28.5)	0 (0)	5 (71.42)
Gram negative	2 (1.7)	2 (3.4)	14 (11.8)	52 (43.7)	28 (23.5)	22 (18.5)	101 (84.9)
<i>K. pneumoniae</i>	0 (0)	2 (4.3)	1 (2.2)	22 (47.8)	11 (23.9)	10 (21.7)	43 (93.4)
<i>P. aeruginosa</i>	1 (2.9)	0 (0)	4 (11.8)	21 (55.9)	5 (14.7)	3 (21.4)	29 (85.3)
<i>E. coli</i>	0 (0)	0 (0)	1 (12.5)	2 (25)	2 (25)	3 (37.5)	7 (87.5)
<i>C. freundii</i>	0 (0)	0 (0)	2 (22.2)	0 (0)	2 (22.2)	5 (55.55)	7 (77.77)
<i>Enterobacter</i> spp.	0 (0)	0 (0)	1 (16.7)	1 (16.7)	3 (33.3)	1 (16.7)	5 (83.3)
<i>H. influenzae</i>	1 (11.1)	0 (0)	1 (22.2)	4 (33.30)	3 (33.3)	0 (0)	7 (77.7)
<i>P. mirabilis</i>	0 (0)	0 (0)	1 (33.3)	0 (0)	2 (66.7)	0 (0)	2 (66.7)
<i>P. vulgaris</i>	0 (0)	0 (0)	1 (50)	1 (50)	0 (0)	0 (0)	1 (50)
<i>Acinetobacter</i> species ^a	0 (0)	0 (0)	2 (100)	2 (100)	0 (0)	0 (0)	2 (100)
Overall total ^b	2 (1.1)	6 (3.4)	21 (12)	80 (45.7)	44 (25.1)	25 (14.3)	148 (84.6)

MDR = multidrug resistance; R₀ = No antibiotic resistance; R₁ = resistance to one; R₂ = resistance to two; R₃ = resistance to three; R₄ = resistance to four; R₅ = resistance to five and more than five antibiotics class.

^aPercent is computed from the total number of each bacteria species.

^bPercent is computed from a total number of isolates.

resistance to five and more antimicrobials. The overall prevalence of MDR bacteria was 148 (84.6%). The MDR rate of Gram-positive and Gram-negative isolates was 47 (83.9%) and 101 (84.9%), respectively. About 43 (93.4%) of *K. pneumoniae*, 7 (87.5%) of *E. coli*, 42 (85.7%) of *S. pneumoniae*, 5 (83.3%) of *Enterobacter* spp. and 67% of *H. influenzae* isolates developed MDR (Table 5).

Factors associated with bacterial infection

All variables with a *p*-value of 0.25 in the bivariate analysis were entered into multivariable logistic regression analysis. Accordingly, alcohol consumption frequently (Adjusted Odds Ratio (AOR)=3.474, 95% CI: 1.07–11.31, *p*=0.039), viral load >1000 copies/ml (AOR=4.88, 95% CI: 1.88–36.54, *p*=0.002), cigarette smoking (AOR=3.87, 95% CI: 1.56–54.87, *p*=0.023), CD4⁺ cell count less than 200 cells/mm³ (AOR=6.5, 95% CI: 1.16–11.21, *p*=0.027), WHO HIV clinical stage IV (AOR=9.51, 95% CI: 1.66–54.45, *p*=0.011), educational status of participants who only read and write (AOR=3.415, 95% CI: 1.48–22.88, *p*=0.006); and most white blood cell count less than 4000 cells/mm³ (AOR=3.5, 95% CI: 1.90–19.57, *p*=0.023) were found to have statistically significant association with bacterial infection (Table 6).

Discussion

Bacterial pneumonia turns into the most common infection in HIV-infected patients and the most joint cause of hospital admission and mortality in high HIV epidemic countries.³⁶

In the present study, the overall prevalence of culture positive sputum among HIV-positive patients was 46.3%. This is comparable with previous studies done in Ethiopia such as Mekelle 43.7%,²¹ Jimma 45%³⁷ and Arbaminch 42.9%,³⁸ and other parts of the world including Nigeria 42.9%³⁹ and Nepal 46.6%.³⁰ The current finding, however, is higher than previous reports from India (16.6%⁹ and 17.1%),⁴³ Malawi 29%,⁴⁰ Nepal 39.7%¹⁷ and Ethiopia (32.1% and 40.3%),^{41,42} but lower than previous prevalence reports from Nigeria (54.07% and 55.6%)^{43,44} and Spain 53.5%.¹⁶ The observed inconsistency might be due to methodological differences, sample size, seasonal variations, or sociodemographic variability of the study participants. Besides, the majority of our study participants had CD4 cell counts of less than 350 cells/mm³ and a higher viral load, which subsequently increases the incidence of bacterial infection and leads to concurrent opportunistic infections.⁷

Similarly, like most previous study reports in our country and elsewhere, our findings showed that the majority of the isolated etiological agents were Gram-negative bacteria (68%).

Table 6. Bivariable and multivariable logistic regression analysis of factors associated with bacterial pneumonia among pneumonia suspected HIV patient's infection at Dessie town health facilities, Northeast Ethiopia, 2021.

Variable of study participant	Category	Bacterial growth N (%)		COR (95% CI)	p-Value	AOR (95% CI)	p-Value		
		Yes	No						
Sex	Male	87 (46.5)	100 (53.5)	1.8 (0.68–12.53)	0.93	NA			
	Female	88 (46)	103 (54)	Ref					
Age of study participant	10–16	4 (40)	6 (60)	Ref					
	17–30	27 (35.1)	50 (64.9)	1.57 (0.15–2.13)	0.405				
	31–45	81 (46.6)	93 (53.4)	1.46 (0.26–0.84)	0.34				
	>45	63 (53.8)	54 (46.2)	1.747 (0.47–1.19)	0.28				
Alcohol drinking habit	Always	13 (56.5)	10 (43.5)	9.46 (1.04–5.83)	0.18	3.47 (1.07–11.31)	0.039*		
	Sometimes	67 (83.8)	13 (16.20)	1.76 (5.13–18.59)	0.29				
	Never	95 (34.5)	180 (65.50)	Ref					
Occupational status	Employed	20 (24.4)	62 (75.6)	Ref					
	Student	9 (37.5)	15 (62.5)	1.86 (0.71–4.90)	0.209				
	Merchant	52 (53.6)	45 (46.4)	1.75 (0.88–16.81)	0.35				
	House wife	17 (60.7)	11 (39.3)	1.9 (0.92–11.91)	0.28				
	Farmer	7 (43.8)	9 (56.2)	1.41 (0.80–17.04)	0.25	1.90 (0.06–15.29)	0.061		
	Daily labor	70 (53.4)	61 (46.4)	3.56 (1.93–6.55)	0.01	2.25 (1.09–6.71)	0.009*		
Educational status	Illiterate	52 (50)	52 (50)	1.43 (0.783–2.623)	0.12	3.45 (1.48–22.88)	0.006*		
	Only read and write	29 (78.4)	8 (21.6)	1.6 (0.89–12.92)	0.26				
	Primary school	38 (38)	62 (62)	0.88 (0.47–1.63)	0.681				
	Secondary-school	26 (40.6)	38 (59.4)	0.981 (0.50–1.94)	0.955				
	College and above	30 (41.1)	43 (58.9)	Ref					
Residence	Rural	57 (64)	32 (36)	2.58 (1.58–4.22)	0.15	2.579 (1.66–18.06)	0.025*		
	Urban	118 (40.8)	171 (59.2)	Ref					
CD ⁺ count	<200 cells/mm ³	35 (68.6)	16 (31.4)	5.51 (8.04–49.39)	0.018	6.50 (1.16–11.21)	0.027*		
	201–350 cells/mm ³	95 (69.3)	42 (30.7)	4.90 (9.46–44.88)	0.21			3.20 (0.72–13.11)	0.128
	351–500 cells/mm ³	36 (36.4)	63 (63.6)	0.56 (0.23–11.60)	0.65				
	>500 cells/mm ³	9 (9.9)	82 (90.1)	Ref					
WBC cells count	Low (<4000)	134 (72.4)	51 (27.6)	11.80 (1.71–19.52)	0.017	3.50 (1.90–19.57)	0.023*		
	High (>11,000)	10 (43.5)	13 (56.5)	0.74 (0.45–13.76)	0.35				
	Normal (4000–11,000)	31 (1)	139 (82.5)	Ref					
WHO clinical stage	Stage I	32 (25.2)	95 (74.8)	Ref					
	Stage II	88 (52.4)	80 (47.8)	0.56 (0.19–11.65)	0.290				
	Stage III	49 (73.1)	18 (26.9)	1.86 (0.65–5.34)	0.251				
	Stage IV	10 (62.5)	6 (37.5)	4.54 (1.44–14.29)	0.010	9.51 (1.66–54.47)	0.011*		
Marital status	Single	5 (20)	20 (80)	Ref					
	Divorced	7 (38.9)	11 (61.1)	1.26 (0.96–0.71)	0.58				
	Widowed	7 (41.2)	10 (58.8)	1.66 (0.25–1.75)	0.404				
	Married	156 (49.1)	162 (50.9)	1.73 (0.27–1.96)	0.528				
Current smoking habit	Yes	88 (73.3)	32 (26.70)	5.45 (1.35–8.73)	0.021	3.87 (1.56–23.82)	0.023*		
	No	87 (33.7)	171 (66.3)	Ref					
Chat chewing habit	Always	20 (76.9)	6 (23.1)	3.48 (2.14–14.07)	0.011	8.48 (1.87–30.07)	0.001*		
	Sometimes	42 (79.2)	11 (20.8)	3.28 (3.11–12.70)	0.056				
	Never	113 (37.8)	186 (62.8)	Ref					
Viral load	>1000 copies/ml	147 (81.7)	33 (18.3)	6.99 (2.8–58.46)	0.021	4.85 (1.88–36.54)	0.002*		
	150–1000 copies/ml	6 (40)	9 (60)	1.88 (0.58–15.03)	0.21				
	<150 copies/ml	22 (12.0)	61 (88)	Ref					
Underlying chronic diseases	Yes	22 (84.6)	4 (15.4)	2.14 (0.45–11.41)	0.45				
	No	153 (43.5)	199 (56.5)	Ref					
Coinfection/comorbidity	Yes	42 (84)	8 (16)	1.13 (0.66–25.29)	0.29				
	No	133 (40.5)	195 (59.5)	Ref					

(Continued)

Table 6. (Continued)

Variable of study participant	Category	Bacterial growth N (%)		COR (95% CI)	p-Value	AOR (95% CI)	p-Value
		Yes	No				
History of ART failure	Yes	16 (61.5)	10 (68.5)	1.52 (0.23–13.17)	0.312		
	No	159 (45.2)	193 (54.80)	Ref			
BMI	Under weight	82 (75.9)	26 (24.1)	1.519 (0.39–12.72)	0.65		
	Over weight	12 (36.4)	21 (63.4)	0.91 (0.43–1.94)			
	Normal weight	81 (36.2)	156 (65.8)	Ref			
ART treatment duration	7–12 months	2 (66.7)	1 (33.3)	Ref	0.340	NA	
	1–5 years	28 (38.9)	44 (61.1)	0.67 (0.28–1.56)			
	>5 years	145 (47.9)	158 (52.1)	0.99 (0.46–2.3)			
Cotrimoxazole prophylaxis	Yes	143 (74.1)	50 (25.9)	Ref	0.35	NA	
	No	32 (17.3)	153 (82.7)	0.65 (0.48–2.10)			

ART = antiretroviral therapy; AOR = adjusted odds ratio; BMI = body mass index; CI = confidence interval; COR = crude odds ratio; Ref = reference.

*Significant at $p < 0.05$.

This finding is compatible with similar studies done in Addis Ababa,⁴¹ Tanzania⁴⁵ and Nepal.¹⁷ However, it is different from the study reports in Bahirdar⁴⁴) and Jimma.⁴³ Etiological differences might be due to environmental contamination and the fact that the majority of Gram-positive bacteria are vulnerable to antibiotics that are widely self-administered, especially the concurrent use of cotrimoxazole drug as a prophylactic agent among HIV patients, which may result in a reduction in the incidence of Gram-positive bacteria.

In the present study, the predominant isolate was *S. pneumoniae* 49 (26.3%) followed by *K. pneumoniae* 46 (47.4%) and *P. aeruginosa* 34 (19.4%). This result is inconsistent with other studies conducted in Ethiopia such as Arbaminch,³⁸ Jimma,⁴³ Bahirdar,⁴⁴ others with different geographical regions of the world like India,^{1,46} United Kingdom⁴⁰ and Spain¹⁵ showed that *S. pneumoniae* was the predominant isolate. On the contrary, previous studies conducted in Mekelle,²¹ Addis Ababa⁴¹ Tanzania⁴⁵ and Bahirdar⁴⁷; and elsewhere in the world including Nepal,¹⁷ Nigeria^{43,44} and India.^{14,26,48} This dominance could be attributed to their capsular nature and the emergence of strains from both species with additional genetic traits. The bacteria's ability to form biofilm, fimbriae and capsular protein to prevent phagocytosis by polymorphonuclear leukocytes and macrophages, as well as the emergence of strains that can acquire additional genetic traits, may explain the bacteria's dominance⁴⁹ and capsule that causes severe pneumonia in immunocompromised people.⁵⁰

According to the international standard for the definition of drug resistance,¹⁶ the overall multidrug resistance was found in 84.6% of the total isolated bacteria. This finding was slightly higher as compared to the studies conducted in Bahirdar Ethiopia 76%,⁴² Nigeria (67.2%)⁵¹ and Cameron 79.4%.⁵² The result was much higher than a study reported from Ethiopia that ranged from 17.9%–56.7%^{18,29,30}; and a systematic review report of 59.7% overall MDR prevalence in Ethiopia.⁵² The reason for the high MDR prevalence might

be due to poor infection control strategies, inappropriate utilization of antimicrobial agents in empirical treatment, extreme antibiotic use, and self-antibiotic prescribing habits.¹⁸ MDR was found in 43 (93.4%) of *K. pneumoniae* isolates, 7 (87.5%) of *E. coli* isolates, 42 (85.7%) of *S. pneumoniae* isolates, 5 (83.3%) of *Enterobacter* spp. isolates and 67% of *H. influenzae* isolates. This is mainly attributed to the ability of those species to express resistance genes, including external and spread on mobile genetic elements, produce extended-spectrum beta-lactamases and carbapenems, aminoglycoside-modifying enzymes, and porin-efflux mechanisms, which are the main contributing factors of MDR.^{53,54}

In this study, Gram negative bacteria showed the highest resistance rate to 67 (88.2%) for tetracycline, 65 (88.2%) for ampicillin and 65 (87.8%) for amoxicillin-clavulanic acid. Similar findings have been reported in previous studies done in Bahirdar,⁴² Jimma,⁴¹ Mekele,²¹ Nigeria,⁴³ Nepal¹⁷ and India.¹⁴ This could be due to the drug's long-term overuse, the expression of extended-spectrum beta-lactamases, which were developed to resist penicillin, cephalosporins, and monobactams, and the expression of carbapenems, which provides resistance to those -lactams, including carbapenems. The capsule, production of biofilm, efflux pumps and production of polysaccharide matrix that coats the cell can limit the penetration of certain agents.^{55,56} On the other hand, lower resistance was observed against gentamicin, amikacin, ciprofloxacin, and meropenem, which is indicative of a possible drug of choice for such infections.

In our study, Gram-positive bacteria were resistant to tetracycline 49 (87.5%), penicillin 48 (85.7%), trimethoprim-sulfamethoxazole 34 (78.6%) and chloramphenicol 37 (66.1%), which is in line with previous studies conducted in Arbaminch,³⁸ Jimma,⁴¹ India¹⁴ and Nepal.⁵⁷ This might be due to Lack of pump inhibition, efflux proteins, structural modifications.⁵⁸ Furthermore, long-term usage, easy availability,

common prophylaxis and indiscriminate use of commonly used drugs such as trimethoprim-sulfamethoxazole and penicillin could lead to an increase in resistance. On the other hand, Gram-positive isolates showed sensitivity of 70.0% to clindamycin, 62.1% to ceftriaxone, and 72.4% to oxacillin, which was also in agreement with other studies conducted in Mekelle,²¹ Bahirdar,⁴⁴ China⁵⁹ and India.⁶⁰

Cotrimoxazole has been used as a prophylactic agent against opportunistic infections in HIV/AIDS patients around the world for over 10 years, and it is now used in all HIV/AIDS patients as a prophylactic agent.⁶¹ In our study, 193 patients were given cotrimoxazole and 69.5% of bacterial isolates were resistant to cotrimoxazole. This finding was supported by a study report from Tanzania (81.3%).⁶¹ This might be attributable to long-term usage of cotrimoxazole as prophylaxis and a common antibacterial agent, which subsequently led to increased development of resistance.⁶¹

Patients with CD4⁺ count less than 200 cells/mm³, viral load 1000 copies/ml and viral load >1000 copies/ml, and white blood cell count less than 4000 cells had a statistically significant association with the occurrence of CAP in the current study, which is comparable to studies from Ethiopia,²¹ Nepal,³⁰ India,¹⁷ Brazil,¹² India¹⁴ and Nigeria.⁴⁴ This might be due to the large number of study participants having a low CD4⁺ count and WHO stages III and IV. When depletion of the CD4⁺ T cells occurs, bacterial infection may occur. Increasing viral numbers weaken the immune status of the patient, causing impairment of their immunological system (alveolar CD4⁺ T-cell function is impaired) and is probably related to the impairment of humoral immunity.^{6,62}

Study participants with a viral load of ≥ 1000 copies/ml were 4.85 times more likely to develop bacterial infection, respectively, as compared to individuals having <150 copies/ml viral load. This is in agreement with studies conducted in Ethiopia,⁴⁴ Spain⁶³ and South Africa.⁶⁴ This might be due to the fact that virus-mediated immunosuppression of the host innate immunity exposes the patient to opportunistic bacteria colonization and viral-induced secondary bacterial infections.⁶²

Cigarette smoking is one of the potential risk factors for the development of pneumonia. In our study, pneumonia patients who smoked frequently were 3.87 times more likely to develop bacterial infection than non-smokers. This was comparable with studies done in Ethiopia,²¹ Brazil¹² and Nepal,¹⁷ Kenya⁶⁵ and Spain^{63,66} that showed a significant association of cigarette smoking with bacterial infection. This is because cigarette smoking produces structural changes in the respiratory tract that cause peribronchial inflammation and fibrosis, increased mucosal permeability, weakens the muco-ciliary clearance, affects pathogen adherence, and disruption of the respiratory epithelium, which in turn exposes one to lower respiratory tract infections, which may increase the cigarette smoke-induced lung inflammation.⁶⁷

Alcohol consumption was also found to have a statistically significant association with bacterial infection. In our study, alcohol consumers were 3.474 times more likely to have bacterial pneumonia than non-consumers. This was in

line with studies done in Ethiopia,²¹ South Africa⁶⁸ and North America.⁶⁹ This might be because of the sedative effects of alcohol, which can reduce the oropharyngeal tone, leading to an increased risk of aspiration of microbes and altering the pathophysiology by decreasing the phagocytic function of the alveolar macrophages, impairing bacterial clearance and diminishing pulmonary defence against infection. Furthermore, high levels of alcohol blunt mental function and suppress cough and gag reflexes. It also decreases mucociliary clearance, impairing both innate and acquired immunity, reduces the production of chemokines, and blunts the chemotaxis of neutrophils.⁶⁹

Limitation of the study

Due to a lack of resources, we did not attempt to isolate atypical bacterial agents such as Chlamydia, mycoplasma, or Legionella species and respiratory infections. Moreover, molecular characterization of the isolated bacterial agents was not done.

Conclusion

In the current study, relatively higher proportions of *K. pneumoniae*, *P. aeruginosa* and *S. pneumoniae* were the most bacterial isolated identified. In this study, majority of isolated bacteria were Gram-negative. Most of the isolates were found susceptible to gentamicin, cefotaxime and ceftriaxone. However, they were resistant to commonly used antimicrobials like amoxicillin, tetracycline, amoxicillin-clavulanate and co-trimoxazole. However, antimicrobial resistance including MDR was observed to a number of commonly used antibiotics, such as trimethoprim-sulfamethoxazole ampicillin, amoxicillin-clavulanic acid and tetracycline. Detectable recent viral load, low CD4⁺ count, late WHO clinical stage, cigarette smoking and alcohol consumption were found significant predictors of bacterial infection. Thus, preventive measures to minimize the risk of the disease including lifestyle factors such as smoking and alcohol consumption; and advocacy of proper antimicrobial usage should be done. ART services should strengthen good adherence of ART and monitoring to decrease patients' viral load and increase CD4⁺ counts. Moreover, culture screening and antimicrobial susceptibility testing should be practiced routinely; and large-scale population-based surveys are needed to assess effectiveness of cotrimoxazole and molecular-based studies are needed to monitor the epidemiology of MDR bacterial isolates and combat antimicrobial resistance.

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Authors' contributions

Conceived and designed the experiments: Mihret Tilahun, Abdurrahman Seid, Daniel Gebretsadik; performed the experiments: Mihret Tilahun, Abdurrahman Seid, Agumas Shibabaw; analyzed the data: Mihret Tilahun, Abdurrahman Seid, Daniel Gebretsadik, Alemu Gedefie, Melaku Ashagrie Belete; and wrote and edited the manuscript: Mihret Tilahun, Abdurrahman Seid, Alemu Gedefie, Agumas Shibabaw, Melaku Ashagrie Belete. All authors read and approved the final manuscript.

Declaration of conflicting interests

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

Ethics approval

The study was done from HIV-positive CAP suspected patients from antiretroviral clinics in the health facilities of the Dessie town. Supportive letter to conduct this study was obtained from college of medicine and health science community service, research and post graduate office, Wollo University with a protocol number of CMHS/HC/354/13. A permission letter was obtained from the Dessie town health office and then in each health institutions.

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Informed consent

Written informed consent was obtained from the subjects and the legally authorized representatives of the minor subjects prior to study initiation.

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Data availability statement

Data supporting the conclusions of this article are within the manuscript and are available on reasonable request from the principal investigators due to ethical reasons.

Supplemental material

Supplemental material for this article is available online.

References

- Shah RR. Microbiological profile of respiratory tract infections among HIV-infected and HIV-non infected patients: a comparative study. *Int Arch Integr Med* 2016; 3(2): 51–59.
- Morris A, Crothers K, Beck J, et al. An official ATS workshop report: emerging issues and current controversies in HIV-associated pulmonary diseases. *Proc Am Thorac Soc* 2011; 8(1): 17–26.
- Tural Önür S, Dalar L, İliaz S, et al. Pneumonia in HIV-infected patients. *Eurasian J Pulmonol* 2016; 18(1): 11–17.
- Ethiopian Federal Ministry of Health. Federal HIV/AIDS prevention and control office federal ministry of health of Ethiopia. Guidelines for management of opportunistic infections and anti retroviral treatment in adolescents and adults in Ethiopia. 2008.
- Gordin FM, Roediger MP, Girard P-M, et al. Pneumonia in HIV-infected persons: increased risk with cigarette smoking and treatment interruption. *Am J Respir Crit Care Med* 2008; 178(6): 630–636.
- Crothers K, Thompson BW, Burkhardt K, et al. HIV-associated lung infections and complications in the era of combination antiretroviral therapy. *Proc Am Thorac Soc* 2011; 8(3): 275–281.
- Lazarous DG and O'Donnell AE. Pulmonary infections in the HIV-infected patient in the era of highly active antiretroviral therapy: an update. *Curr Infect Dis Rep* 2007; 9(3): 228–232.
- Ferkol T and Schraufnagel D. The global burden of respiratory disease. *Ann Am Thorac Soc* 2014; 11(3): 404–406.
- Kandati J, Boorsu SK, Lakshmi Ponugoti M, et al. Bacterial and fungal agents causing lower respiratory tract infections in patients with human immunodeficiency virus infection. *Int J Res Med Sci* 2016; 4(8): 3595–3600.
- Koss CA, Jarlsberg LG, den Boon S, et al. A clinical predictor score for 30-day mortality among HIV-infected adults hospitalized with pneumonia in Uganda. *PLoS One* 2015; 10(5): e0126591.
- Curran A, Falco V, Crespo M, et al. Bacterial pneumonia in HIV-infected patients: use of the pneumonia severity index and impact of current management on incidence, aetiology and outcome. *HIV Med* 2008; 9(8): 609–615.
- Lamas CC, Coelho LE, Grinsztejn BJ, et al. Community-acquired lower respiratory tract infections in HIV-infected patients on antiretroviral therapy: predictors in a contemporary cohort study. *Infection* 2017; 45(6): 801–809.
- Gingo MR and Morris A. Pathogenesis of HIV and the lung. *Curr HIV/AIDS Rep* 2013; 10(1): 42–50.
- Shilpa A, Anuradha K and Venkatesha D. Drug susceptibility pattern of aerobic bacterial isolates from pulmonary infection in HIV seropositives and their correlation with CD4 Count. *IOSR J Dent Med Sci* 2014; 13(3): 37–41.
- Cilloniz C, Torres A, Polverino E, et al. Community-acquired lung respiratory infections in HIV-infected patients: microbial aetiology and outcome. *Eur Respir J* 2014; 43(6): 1698–1708.
- Conklin LM, Bigogo G, Jagero G, et al. High Streptococcus pneumoniae colonization prevalence among HIV-infected Kenyan parents in the year before pneumococcal conjugate vaccine introduction. *BMC Infect Dis* 2015; 16(1): 18.
- Khushbu Y and Satyam P. Bacteriological Profile of Lower Respiratory Tract Infection (LRTI) among HIV Seropositive Cases in Central Terai of Nepal. *Int J Curr Microbiol App Sci* 2015; 4(11): 431–442.
- Mulu W, Yizengaw E, Alemu M, et al. Pharyngeal colonization and drug resistance profiles of *Moraxella catarrhalis*, *Streptococcus pneumoniae*, *Staphylococcus aureus*, and *Haemophilus influenzae* among HIV infected children attending ART Clinic of Felegehiwot Referral Hospital, Ethiopia. *PLoS One* 2018; 13(5): e0196722.

19. Magiorakos A-P, Srinivasan A, Carey R, et al. Multidrug-resistant, extensively drug-resistant and pandrug-resistant bacteria: an international expert proposal for interim standard definitions for acquired resistance. *Clin Microbiol Infect* 2012; 18(3): 268–281.
20. Sweeney MT, Lubbers BV, Schwarz S, et al. Applying definitions for multidrug resistance, extensive drug resistance and pandrug resistance to clinically significant livestock and companion animal bacterial pathogens. *J Antimicrob Chemother* 2018; 73(6): 1460–1463.
21. Adhanom G, Gebreegziabihier D, Weldu Y, et al. Species, risk factors, and antimicrobial susceptibility profiles of bacterial isolates from HIV-infected patients suspected to have pneumonia in Mekelle zone, Tigray, northern Ethiopia. *BioMed Res Int* 2019; 2019: 1–9.
22. Dastider D, Jyoti Sen D, Kumar Mandal S, et al. Hand sanitizers bid farewell to germs on surface area of hands. *Eur J Pharm Med Res* 2020; 7(4): 648–656.
23. Webb MS and Carey MP. Tobacco smoking among low-income Black women: demographic and psychosocial correlates in a community sample. *Nicotine Tob Res* 2008; 10(1): 219–229.
24. Saracino A, Lorenzini P, Caputo SL, et al. Increased risk of virologic failure to the first antiretroviral regimen in HIV-infected migrants compared to natives: data from the ICONA cohort. *Clin Microbiol Infect* 2016; 22(3): 288.e1–288.e8.
25. McShane H. Co-infection with HIV and TB: double trouble. *Int J STD AIDS* 2005; 16(2): 95–101.
26. Sperrin M, Marshall AD, Higgins V, et al. Body mass index relates weight to height differently in women and older adults: serial cross-sectional surveys in England (1992–2011). *J Public Health* 2016; 38(3): 607–613.
27. Enawgaw B, Birhan W, Abebe M, et al. Haematological and immunological reference intervals for adult population in the state of Amhara, Ethiopia. *Trop Med Int Health* 2018; 23(7): 765–773.
28. Walia K, Madhumathi J, Veeraraghavan B, et al. Establishing antimicrobial resistance surveillance & research network in India: journey so far. *Indian J Med Res* 2019; 149(2): 164.
29. Ioannidis P, Papaventsis D, Karabela S, et al. Cepheid GeneXpert MTB/RIF assay for Mycobacterium tuberculosis detection and rifampin resistance identification in patients with substantial clinical indications of tuberculosis and smear-negative microscopy results. *J Clin Microbiol* 2011; 49(8): 3068–3070.
30. Ojha CR, Rijal N, Khagendra K, et al. Lower respiratory tract infections among HIV positive and control group in Nepal. *VirusDisease* 2015; 26(1): 77–81.
31. Murray PR, Rosenthal KS and Pfaller MA. *Medical microbiology E-Book*. Philadelphia, PA: Elsevier Health Sciences, 2020.
32. Zapata A and Ramirez-Arcos S. A comparative study of McFarland turbidity standards and the Densimat photometer to determine bacterial cell density. *Curr Microbiol* 2015; 70(6): 907–909.
33. Weinstein MP and Lewis JS. The clinical and laboratory standards institute subcommittee on antimicrobial susceptibility testing: background, organization, functions, and processes. *J Clin Microbiol* 2020; 58(3): e01864-19.
34. Abbey TC and Deak E. What's new from the CLSI subcommittee on antimicrobial susceptibility testing M100 29th edition. *Clin Microbiol Newsl* 2020; 41(23): 203–209.
35. Magiorakos A-P, Srinivasan A, Carey R, et al. Multidrug-resistant, extensively drug-resistant and pandrug-resistant bacteria: an international expert proposal for interim standard definitions for acquired resistance. *Clin Microbiol Infect* 2012; 18(3): 268–281.
36. Kengne M, Lebogo MBB, Nwobegahay JM, et al. Antibiotics susceptibility pattern of Streptococcus pneumoniae isolated from sputum cultures of human immunodeficiency virus infected patients in Yaoundé, Cameroon. *Pan African Med J* 2018; 31: 16.
37. Regasa B, Yilma D and Sewunet T. *Antimicrobial susceptibility pattern of bacterial isolates from community-acquired pneumonia patients in Jimma University Specialized Hospital, Jimma, Ethiopia*. 2015.
38. Regasa B. Drug resistance patterns of bacterial pathogens from adult patients with pneumonia in Arba Minch hospital, South Ethiopia. *Glob J Med Res* 2014; 14(5): 1–4.
39. Salami A, Oluboyo P, Akambi A, et al. Bacterial pneumonia in the AIDS patients. *West African J Med* 2006; 25(1): 1–5.
40. Hartung TK, Chimbayo D, van Oosterhout JJ, et al. Etiology of suspected pneumonia in adults admitted to a high-dependency unit in Blantyre, Malawi. *Am J Trop Med Hyg* 2011; 85(1): 105–112.
41. Nurahmed N, Kedir S, Fantahun S, et al. Bacterial profile and antimicrobial susceptibility patterns of lower respiratory tract infection among patients attending selected health centers of Addis Ababa, Ethiopia. *Egypt J Chest Dis Tuberc* 2020; 69(2): 399.
42. Temesgen D, Bereded F, Derbie A, et al. Bacteriology of community acquired pneumonia in adult patients at Felege Hiwot Referral Hospital, Northwest Ethiopia: a cross-sectional study. *Antimicrob Resist Infect Control* 2019; 8(1): 1–8.
43. Ojo-Bola O and Oluyeye A. Antibiotics resistance of bacteria associated with pneumonia in HIV/AIDS patients in Nigeria. *Am J Infect Dis Microbiol* 2014; 2(6): 138–144.
44. Urama EU, Enweani IB, Oshim IO, et al. Microbiological profile of respiratory tract infections among HIV sero-positive subjects attending Nnamdi Azikiwe University Teaching Hospital Nnewi, Nigeria. *Amer J Med and Med Sci* 2018; 8: 37–42.
45. Kishimbo P, Sogone NM, Kalokola F, et al. Prevalence of gram negative bacteria causing community acquired pneumonia among adults in Mwanza City, Tanzania. *Pneumonia* 2020; 12(1): 1–9.
46. Mane A, Gujar P, Gaikwad S, et al. Aetiological spectrum of severe community-acquired pneumonia in HIV-positive patients from Pune, India. *Indian J Med Res* 2018; 147(2): 202.
47. Genetu DE and Zenebe Y. *Bacterial profile and their antibiotic resistance pattern among HIV patients diagnosed with pneumonia in Felege-Hiwot referral hospital, Bahir Dar, Northwest Ethiopia*. 2020 [Unpublished manuscript].
48. Hidron AI, Kempker R, Moanna A, et al. Methicillin-resistant Staphylococcus aureus in HIV-infected patients. *Infect Drug Resist* 2010; 3: 73.
49. Effah CY, Sun T, Liu S, et al. Klebsiella pneumoniae: an increasing threat to public health. *Ann Clin Microbiol Antimicrob* 2020; 19(1): 1–9.
50. Lee C-R, Lee JH, Park KS, et al. Antimicrobial resistance of hypervirulent Klebsiella pneumoniae: epidemiology, hypervirulence-associated determinants, and resistance mechanisms. *Front Cell Infect Microbiol* 2017; 7: 483.

51. Adeyemi FM, Adejuyigbe E, Ebhodaghe BI, et al. Molecular characterization and antibiotic resistance profile. *Arch Clin Microbiol* 2018; 6(1): 269–276.
52. Marbou WJT and Kuete V. Bacterial resistance and immunological profile. *J Infect Public Health* 2017; 10(3): 269–276.
53. Muhie OA. Antibiotic use and resistance pattern in Ethiopia: systematic review and meta-analysis. *Int J Microbiol* 2019; 2019: 1–8.
54. Cag Y, Caskurlu H, Fan Y, et al. Resistance mechanisms. *Ann Transl Med* 2016; 4(17): 1–8.
55. Gudata D and Begna F. Antimicrobial resistance. *Int J Res Granthaalayah* 2018; 6(11): 77–93.
56. Liwa AC and Jaka H. Antimicrobial resistance: mechanisms of action of antimicrobial agents In: Méndez-Vilas A (ed.) *The battle against microbial pathogens: basic science, technological advances and educational programs*. Madrid: Formatex, 2015, vol. 5, pp. 876–885.
57. Shrestha S, Acharya A, Gautam A, et al. Lower respiratory tract pathogens and their antimicrobial susceptibility pattern in a medical hospital of central Nepal. *Int J Biomed Adv Res* 2013; 4(5): 335–340.
58. Schindler BD and Kaatz GW. Multidrug efflux pumps of gram-positive bacteria. *Drug Resist Updat* 2016; 27: 1–13.
59. Jinghua M, Gaizhuang L and Qiaoli C. *Pathogens and antibiotic resistance of children with community-acquired pneumonia*. 2017.
60. Menon RU, George AP and Menon UK. Etiology and antimicrobial sensitivity of organisms causing community acquired pneumonia: a single hospital study. *J Family Med Prim Care* 2013; 2(3): 244.
61. Marwa KJ, Mushi MF, Konje E, et al. Resistance to cotrimoxazole and other antimicrobials among isolates from HIV/AIDS and non-HIV/AIDS patients at Bugando Medical Centre, Mwanza, Tanzania. *AIDS Res Treat* 2015; 2015: 1–8.
62. Cribbs SK, Crothers K and Morris A. Pathogenesis of HIV-related lung disease: immunity, infection, and inflammation. *Physiol Rev* 2020; 100(2): 603–632.
63. Benito N, Moreno A, Miro J, et al. Pulmonary infections in HIV-infected patients: an update in the 21st century. *Eur Respir J* 2012; 39(3): 730–745.
64. Feldman C and Anderson R. HIV-associated bacterial pneumonia. *Clin Chest Med* 2013; 34(2): 205–216.
65. Muthumbi E, Lowe BS, Muyodi C, et al. Risk factors for community-acquired pneumonia among adults in Kenya: a case-control study. *Pneumonia* 2017; 9(1): 1–9.
66. Rivero-Calle I, Pardo-Secco J, Aldaz P, et al. Incidence and risk factor prevalence of community-acquired pneumonia in adults in primary care in Spain (NEUMO-ES-RISK project). *BMC Infect Dis* 2016; 16(1): 1–8.
67. Voss M, Wonnemberg B, Honecker A, et al. Cigarette smoke-promoted acquisition of bacterial pathogens in the upper respiratory tract leads to enhanced inflammation in mice. *Respir Res* 2015; 16(1): 1–11.
68. Segal LN, Methé BA, Nolan A, et al. HIV-1 and bacterial pneumonia in the era of antiretroviral therapy. *Proc Am Thorac Soc* 2011; 8(3): 282–287.
69. Simou E, Britton J and Leonardi-Bee J. Alcohol and the risk of pneumonia: a systematic review and meta-analysis. *BMJ Open* 2018; 8(8): e022344.