

# Microbiological Etiologies of Pneumonia Complicating Stroke

## A Systematic Review

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**Background and Purpose**—Identifying the causal pathogens of pneumonia complicating stroke is challenging, and antibiotics used are often broad spectrum, without recourse to the microbiological cause. We aimed to review existing literature to identify organisms responsible for pneumonia complicating stroke, before developing a consensus-based approach to antibiotic treatment.

**Methods**—A systematic literature review of multiple electronic databases using predefined search criteria was undertaken, in accordance with Cochrane and PRISMA (Preferred Reporting Items for Systematic Reviews and Meta-Analyses) guidance. Published studies of hospitalized adults with ischemic stroke, intracerebral hemorrhage, or both, which identified microbiological etiologies for pneumonia complicating stroke up to January 1, 2017, were considered. Analysis included summary statistics and random-effects meta-analysis where appropriate.

**Results**—Fifteen studies (40% ischemic stroke, 60% ischemic stroke and intracerebral hemorrhage) involving 7968 patients were included. Reported occurrence of pneumonia varied considerably between studies (2%–63%) with a pooled frequency of 23% (95% confidence interval, 14%–34%;  $I^2=99%$ ). Where reported (60%), the majority of pneumonia occurred within 1 week of stroke (78%). Reported frequency of positive culture data (15%–88%) varied widely. When isolated, aerobic Gram-negative bacilli (38%) and Gram-positive cocci (16%) were most frequently cultured; commonly isolated organisms included *Enterobacteriaceae* (21.8%: *Klebsiella pneumoniae*, 12.8% and *Escherichia coli*, 9%), *Staphylococcus aureus* (10.1%), *Pseudomonas aeruginosa* (6%), *Acinetobacter baumannii* (4.6%), and *Streptococcus pneumoniae* (3.5%). Sputum was most commonly used to identify pathogens, in isolation (40%) or in conjunction with tracheal aspirate (15%) or blood culture (20%).

**Conclusions**—Although the analysis was limited by small and heterogeneous study populations, limiting determination of microbiological causality, this review suggests aerobic Gram-negative bacilli and Gram-positive cocci are frequently associated with pneumonia complicating stroke. This supports the need for appropriately designed studies to determine microbial cause and a consensus-based approach in antibiotic usage and further targeted antibiotic treatment trials for enhanced antibiotic stewardship. (*Stroke*. 2018;49:1602-1609. DOI: 10.1161/STROKEAHA.117.020250.)

**Key Words:** bacteria ■ infection ■ pneumonia ■ stroke

Pneumonia complicating stroke occurs frequently, independently increasing mortality 3-fold and increasing hospitalization costs, length of stay, and likelihood of poor outcome

in survivors.<sup>1,2</sup> Although diagnosis remains challenging, the PISCES (Pneumonia in Stroke Consensus) group recommended that Stroke-Associated Pneumonia (SAP) was the preferred

Received November 28, 2017; final revision received May 2, 2018; accepted May 14, 2018.

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Guest Editor for this article was Harold P. Adams, MD.

The online-only Data Supplement is available with this article at <http://stroke.ahajournals.org/lookup/suppl/doi:10.1161/STROKEAHA.117.020250/-/DC1>.

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DOI: 10.1161/STROKEAHA.117.020250

diagnostic terminology covering the spectrum of lower respiratory tract infections complicating stroke within the first week and hospital-acquired pneumonia (HAP) after 1 week.<sup>3</sup> Furthermore, acknowledging the limitations of current biomarkers and accessibility of microbiological samples, modified Centers for Disease Control and Prevention criteria were proposed to aid clinicians and researchers in diagnosing SAP in nonventilated patients.<sup>3</sup>

Once SAP is suspected or diagnosed, however, use of antimicrobials vary and are either clinician dependant or guided by local policy for community-acquired pneumonia (CAP) or HAP.<sup>4</sup> Antibiotics used are often broad spectrum, without recourse to the microbiological cause. The ability to better inform choice of antibiotic therapy in SAP, based on defined or likely microbial cause, might lead to improved outcomes and enhanced antibiotic stewardship. Identifying microbiological cause in nonventilated stroke patients is challenging because of the difficulties in obtaining direct samples from the lower respiratory tract (impaired cough and limited expectoration) and lack of applicable invasive procedures, such as bronchoscopy in conscious stroke patients, in addition to reliance on sputum samples with the inherent risk of contamination from oropharyngeal commensal organisms.<sup>5</sup> Although bacterial colonization of the oropharynx could potentially limit interpretation of positive sputum samples, poor diagnostic sensitivity of microbiological culture methods, such as blood culture specimens (positive in <10%)<sup>6</sup> and pleural fluid aspirate, limit their use. Most importantly, prior use of antibiotics hampers the sensitivity of microbiological techniques, and current stroke guidelines do not recommend early nonselective preventive antibiotic treatment.<sup>7,8</sup> As part of the ongoing PISCES collaboration, we sought to identify microbiological etiologies for pneumonia complicating stroke through a systematic review of available literature to help inform a planned consensus-based approach for antibiotic treatment.

## Methods

A systematic literature review was undertaken in accordance with the PRISMA (Preferred Reporting Items for Systematic Reviews and Meta-Analyses) statement and Cochrane guidance.<sup>9,10</sup> The authors declare that all supporting data are available within the article (and its [online-only Data Supplement](#)).

## Data Sources and Searches

Searches were undertaken in medical databases including Medical Literature Analysis and Retrieval System Online (MEDLINE, National Institute of Health Library Interface, 1946–January 1, 2017), Excerpta Medica database (EMBASE, National Institute of Health Library Interface, 1947–January 1, 2017), Cumulative Index of Nursing and Allied Health Literature (CINAHL, National Institute of Health Library interface, 1937–January 1, 2017) and Cochrane Central Register of Controlled Trials (Wiley interface, current issue) using predefined search criteria and terms (Table I in the [online-only Data Supplement](#)). Hand searching of reference lists for additional eligible articles was also performed, and the members of PISCES group were invited to provide any other potentially eligible articles. Non-English full-text articles were translated and considered for inclusion if eligibility criteria were met.

## Study Selection

Published studies of hospitalized adults with ischemic stroke, intracerebral hemorrhage, or both, which identified any potential pathogen responsible for pneumonia complicating stroke or which had used objective criteria for diagnosing pneumonia complicating stroke (but not reported causative organisms), were independently screened for

eligibility by 2 reviewers (A.K.K. and C.J.S.), using the study title and abstract (Table II in the [online-only Data Supplement](#)). For those studies that used objective criteria for pneumonia but which had not reported causative pathogens, corresponding authors were contacted by e-mail for unpublished information on pathogens responsible for pneumonia if available. Lead or corresponding authors of studies under consideration were also contacted by e-mail to resolve any issues relating to assessment of eligibility or data extraction. Discrepancies relating to eligibility or data extraction were resolved by a consensus discussion between the same 2 study investigators.

## Data Extraction

Data were independently extracted by 2 reviewers (A.K.K. and A.V.) and included study design, sample size, publication status and demographic data (year of study, country of study, clinical environment), stroke type (ischemic, intracerebral hemorrhage, or both), interval from admission to diagnosis of pneumonia, frequency of pneumonia, criteria used for pneumonia diagnosis, mean age, mean National Institutes of Health Stroke Scale score, cardiovascular risk profile, swallow screening, proportion of nil oral/tube-fed, type of culture specimen (sputum, blood culture, pleural or tracheal aspirate, serology), organisms identified, and antibiotic usage (prophylactic or treatment). Reported data in the identified eligible publications were supplemented by contacting corresponding or lead authors where necessary.

## Study Outcomes

The primary outcomes were the frequencies of the most commonly isolated microbial species among the included studies and the proportion of these organisms responsible for pneumonia. Secondary outcomes included (1) the frequency of positive microbiological cultures and proportions of isolated organisms in positive cultures; (2) relationships between microbial species and time interval from stroke onset to pneumonia; and (3) frequency of identified pathogens across different geographical regions.

## Risk of Bias and Quality Assessment

We anticipated inclusion of both randomized controlled trials and non-randomized studies that had pneumonia as an outcome but were not primarily designed to identify microbiological etiologies. Hence, a formal statistical tool for assessing bias or the individual quality of the studies was not used as we were less concerned about the design or the effects of interventions used in the individual studies for this review. However, as cultures for organisms would only be undertaken when pneumonia is suspected or diagnosed, heterogeneity among studies reporting pneumonia was assessed using random-effects model. Heterogeneity was quantified with the  $I^2$  statistic as reported in a previous study.<sup>1</sup> This measures the proportion of variation ascribed to excess heterogeneity beyond that anticipated by chance. Because of anticipated heterogeneity between studies reporting pneumonia, further quantitative meta-analysis on bacterial cause was not undertaken. Summary statistics were instead undertaken to describe primary and secondary outcomes. A post hoc descriptive comparison of the frequency of pathogen species detected in pneumonia complicating stroke with other forms of pneumonia (eg, CAP, HAP) was also undertaken.

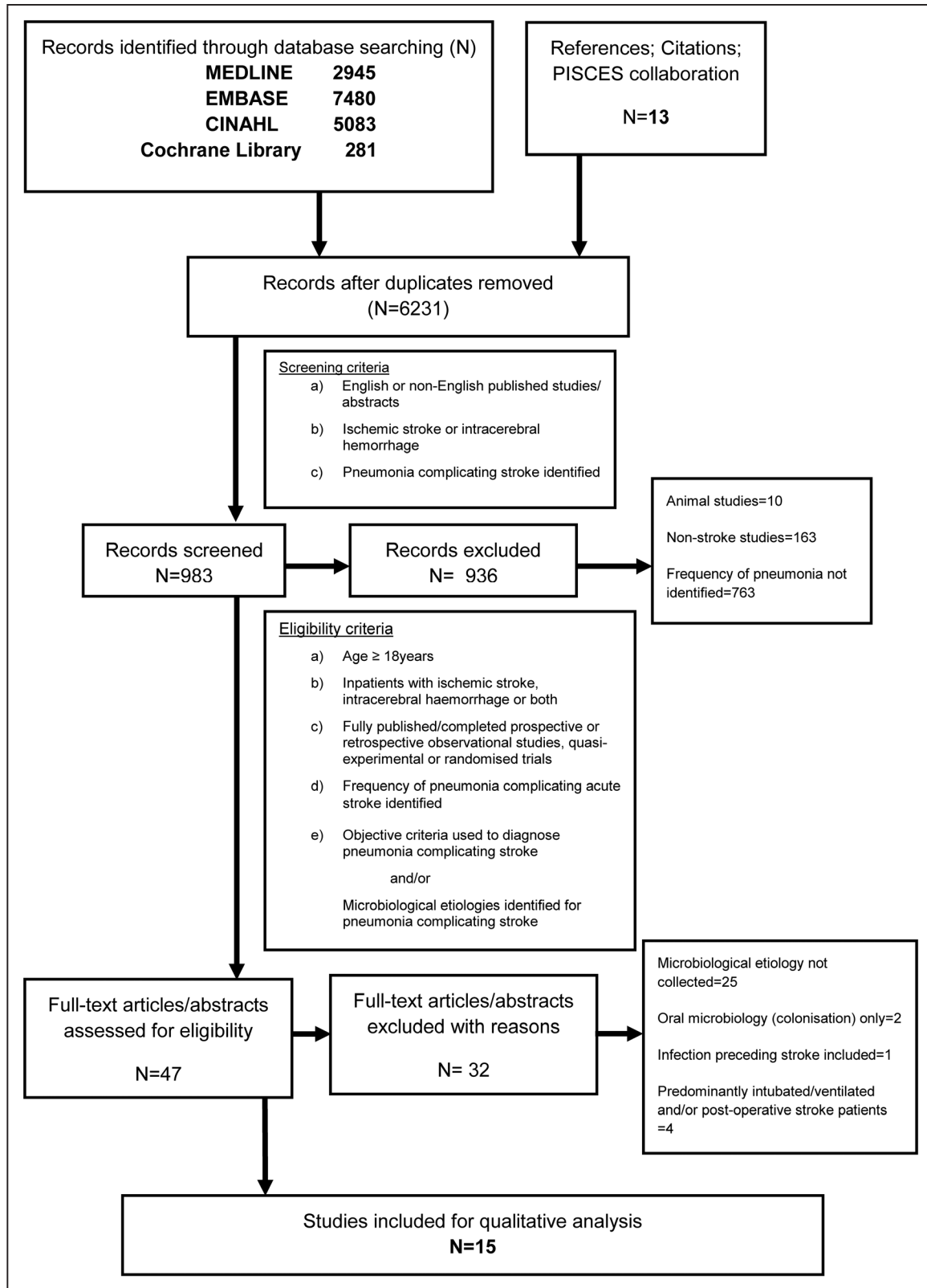
## Results

### Search Results

A total of 6231 unique publications were identified by electronic searches and through the PISCES collaborators (Figure 1). Fifteen fully published studies were finally considered eligible for inclusion.<sup>11–25</sup>

### Study and Patient Characteristics

The studies included retrospective (40%) or prospective (33%) observational studies and randomized trials (27%).



**Figure 1.** Flow diagram of systematic search methodology. CINAHL indicates Cumulative Index of Nursing and Allied Health Literature; and PISCES, Pneumonia in Stroke Consensus.

Fifty-four percent were European or North American studies; 46% were performed in South East Asia or Asia Pacific region. The majority of the studies were conducted on the

acute stroke unit (72%). Other clinical environments included rehabilitation wards (7.5%) and intensive care units (15%). The mean age of the patients in individual studies ranged from

Table 1. Study Characteristics

First Author	Year of Recruitment	Year Published	Country	Design	Participants, n	Stroke Type	Setting	Mean Age, y	Male, %	Mean NIHSS	Time to Pneumonia Diagnosis, d	Culture Specimen	Positive Culture, %	Diagnostic Criteria	Pneumonia Risk Stratification
Chamorro	2002–2004	2005	Spain	1	136	2	1	72.9	50	12	66% <7 d	Sputum/tracheal aspirate	45	2	2
Hassan	1998–2001	2006	Pakistan	3	443	2	1	64	68	NR	67% <2 d	Sputum/tracheal aspirate	38	1	2
Vargas	2001–2002	2006	Spain	2	229	2	1	72.3	51	9	≤7 d	Tracheal aspirate	41	2	2
Ros	1998–1999	2007	Spain	2	258	2	1	74.9	48.8	NR	NR	Blood	22	2	1
Harms	2003–2006	2008	Germany	1	79	1	1	72.5	NR	16	≤11 d	Tracheal aspirate	45	1	3
Sui	2000–2009	2011	China	3	1435	2	2	67.2	55.4	7	63% <3 d	Sputum	38	1	2
Yeh	2006–2007	2011	Taiwan	3	163	2	2	67.4	54	14	NR	Sputum	39	1	3
Fluri	2006–2007	2012	Switzerland	2	383	1	1	71.4	57.7	5	≤5d	Sputum/blood	15	1	1
Chen	2002–2010	2012	Taiwan	3	341	2	3	73.4	48.6	NR	NR	Sputum	NR	3	2
Chen	2006–2011	2013	Taiwan	3	495	1	4	80.3	72.5	15	53% <3 d	Sputum/blood	75	1	2
Satou	NR	2013	Japan	2	16	1	1	83.2	NR	NR	NR	NR	30	2	3
Becker	2005–2009	2014	United States	2	113	1	1	58	66	10	<15 d	Sputum	88	2	1
Warusevitane	2008–2011	2014	United Kingdom	1	60	2	1	77.5	37	19	94% <7 d	Sputum	41	4	3
Li	2009–2011	2014	China	3	1279	1	1	63	NR	NR	NR	Sputum	NR	2	1
Westendorp	2010–2014	2015	The Netherlands	1	2538	2	1	73.5	57	5	NR	Sputum/blood	17	1	1

Design-1 denotes randomized controlled study; 2, prospective observational study; and 3, retrospective study. Stroke type-1 denotes ischemic stroke; 2, ischemic stroke and intracerebral hemorrhage. Setting-1 denotes ASU; 2, Stroke Intensive Care; 3, ASU and rehabilitation unit; and 4, unclear. Criteria-1 denotes Centers for Disease Control and Prevention Criteria; 2, Ad hoc objective criteria; 3, Mann Criteria; and 4, British Thoracic Society Criteria. Pneumonia risk stratification criteria-1 denotes low risk (unselected stroke patients with ≤15% intracerebral hemorrhage, mean NIHSS ≤5); 2, high risk (≥15% intracerebral hemorrhage, NIHSS ≥5); 3, very high risk (exclusively dysphagia studies, exclusively tube fed patients or studies involving patients in intensive care). ASU indicates acute stroke unit; NIHSS, National Institutes of Health Stroke Scale; and NR, not reported.

58 to 83 years. Baseline stroke severity (National Institutes of Health Stroke Scale) was reported in only 73% of studies, with a mean ranging from 5 to 19. Three studies were randomized controlled trials of prophylactic antibiotics.<sup>11,15,25</sup> Two studies were exclusively in nasogastric or percutaneous endoscopic gastrostomy tube-fed participants.<sup>21,23</sup> Thirty-three percent of the studies included mechanically ventilated patients. Entry criteria to the studies varied widely, with only 1 study having the identification of microbiological cause for pneumonia complicating stroke as a primary objective<sup>12</sup>; 2 studies excluded patients on immunosuppressive medication, with prior malignancy, or other forms of immunosuppression before stroke.<sup>11,22</sup> Other comorbidities and cardiovascular risk factor profiles were reported varyingly (Table III in the [online-only Data Supplement](#); Table 1).

### Diagnosis and Frequency of Pneumonia

Reported occurrence of pneumonia varied between studies (2%–63%). Pooled frequency of reported pneumonia was 23% (95% confidence interval, 14%–34%;  $I^2=99%$ ; Figure 2). Substantial heterogeneity was noted even when adjusted to stroke subtype (ischemic stroke,  $I^2=96%$ ; mixed ischemic

stroke and intracerebral hemorrhage,  $I^2=99%$ ) or geographical location (Asian studies,  $I^2=97.4%$  and European or North American studies,  $I^2=97.8%$ ). When reported (60%), the majority of pneumonia occurred within 1 week of stroke (78%). The Centers for Disease Control and Prevention criteria (45%) and ad hoc objective criteria (40%) were the most commonly used objective criteria to diagnose pneumonia.

### Microbiological Cause

Sputum culture was most commonly used to identify pathogens either in isolation (40%) or in conjunction with tracheal aspirate (15%) and blood culture (20%). Reported frequency of positive culture data (15%–88%) varied considerably. Only 3 studies described the culture methods used to identify organisms.<sup>4,13,25</sup> No bacterial growths were reported in 2 studies (15% and 67%).<sup>12,25</sup> Identification of bacterial species in positive cultures varied between studies (Table IV in the [online-only Data Supplement](#)). No pathogen was identified in every study although *Staphylococcus aureus* was identified in positive cultures in 14 of 15 (93%) studies, whereas *Acinetobacter baumannii* was identified in positive cultures in only 6 of 15 (40%) studies (Table IV in the [online-only](#)

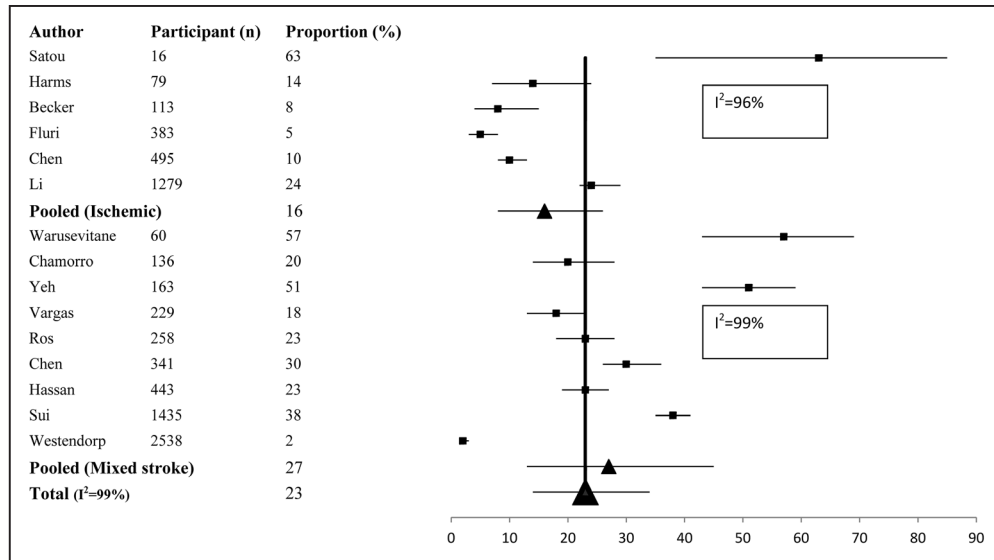


Figure 2. Forest plot of pneumonia frequency according to stroke subtype.

Data Supplement; Figure 3). Antibiotic susceptibility was not reported in the majority of studies; no studies reported any viral or other atypical organisms although it was unclear if these were tested for.

The proportions of microbial species associated with pneumonia also varied between studies. Overall, aerobic Gram-negative bacilli (38%) and Gram-positive cocci (16%) were most frequently responsible for pneumonia; commonly isolated phenotypes (Table 2) included *Enterobacteriaceae* (21.8%: *Klebsiella pneumoniae*, 12.8% and *Escherichia coli*, 9%), *S. aureus* (10.1%), *Pseudomonas aeruginosa* (6%), *A. baumannii* (4.6%), and *Streptococcus pneumoniae* (3.5%). Studies that included patients at relatively higher-risk of pneumonia, that is, exclusively dysphagic patients or intensive care studies,<sup>15,17,19,21</sup> were found to have a high proportion of aerobic Gram-negative bacilli and *S. aureus*, in comparison to lower risk studies (unselected stroke patients with  $\leq 15\%$  intracerebral hemorrhage and mean National Institutes of Health Stroke Scale score  $\leq 5$ ; Table 1). It was not possible to explore relationships between timing of pneumonia or its severity with individual organisms because of insufficient data.

We compared the frequencies of the 8 most commonly identified organisms in pneumonia complicating stroke with

those of hospitalized CAP, ventilator-associated pneumonia (VAP), and HAP (Table 2) from recent reviews of literature (terminologies defined in Table V in the online-only Data Supplement).<sup>26–29</sup> Geographical variations in bacterial cause were observed in our study as seen as in the other reviews. In particular, Gram-negative opportunistic pathogens, such as *P aeruginosa* and *A baumannii*, were more commonly isolated in South Asia or Asia Pacific regions (75% and 100%) as opposed to Western Europe or the United States (28% and 0%). Several organisms were reported with comparable frequency (range) to VAP or HAP (eg, *K pneumoniae*, *E coli*). *S pneumoniae*, the organism most frequently identified in CAP, was detected less often in pneumonia complicating stroke. The organisms most often reported in HAP and VAP (*S aureus* and *P aeruginosa*) were also identified less frequently in pneumonia complicating stroke.

### Antibiotic Usage

Only 4 studies (24%) identified antibiotics used to treat pneumonia complicating stroke.<sup>13,15,20,25</sup> The antibiotic of choice was determined by local hospital policy and commonly included  $\beta$ -lactam (including ureidopenicillin and second/third generation cephalosporins) antibiotics  $\pm$   $\beta$ -lactamase inhibitors and

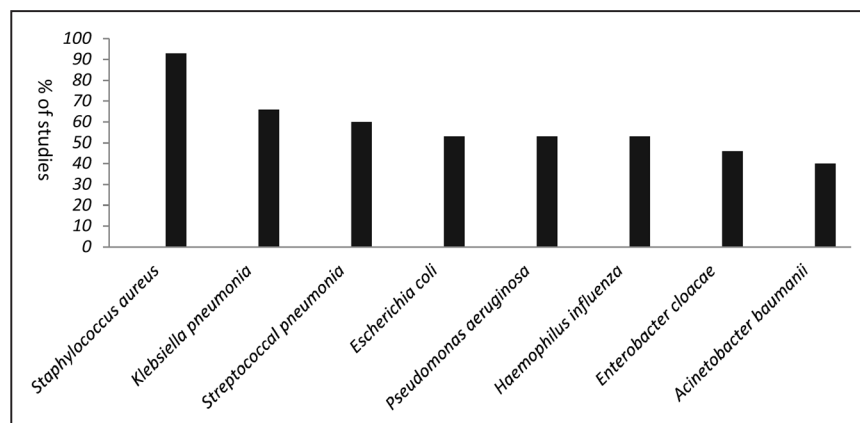


Figure 3. Proportion of studies identifying the 8 most commonly isolated organisms.

Table 2. Frequency of Isolated Organisms in Pneumonia Complicating Stroke in Comparison to Other Forms of Pneumonia

Identified Organisms	HAP (Range, %) <sup>26</sup>	VAP (Range, %) <sup>26</sup>	CAP (Range, %) <sup>26–28</sup>	Pneumonia Complicating Stroke (Weighted Average With Range, %)
Gram-positive cocci				
<i>Staphylococcus aureus</i>	26.6–36.5	19.5–31.9	3–14.1	10.1 (0–36.3)
<i>Streptococcal pneumoniae</i>	1.8–3.2	<3	35–80	3.5 (0–10.7)
Gram-negative bacilli				
<i>Klebsiella pneumoniae</i>	8–10.5	6.6–10.2	3–6	12.8 (0–51)
<i>Escherichia coli</i>	4.6–10.1	3–5	6–12	9 (0–21.7)
<i>Pseudomonas aeruginosa</i>	19–22.4	21.4–26.6	2.8–9	6 (0–11.7)
<i>Acinetobacter baumannii</i>	4.4–13.3	14.3	3–14.3	4.6 (0–22.9)
<i>Haemophilus influenzae</i>	1.3–3.7	NR	5–40	1.9 (0–11.5)
<i>Enterobacter cloacae</i>	6.3–8.5	6–8.8	NR	1.7 (0–10)

CAP indicates community-acquired pneumonia; HAP, hospital-acquired pneumonia; NR, not recorded; and VAP, ventilator-associated pneumonia.

second/third generation fluoroquinolones and was always initiated before obtaining antibiotic sensitivities. Only 1 study reported the proportion of patients with pneumonia receiving antibiotics, the number of pneumonia episodes, and functional outcomes after treatment with antibiotics.<sup>25</sup>

## Discussion

Pneumonia occurs most frequently during the first week after stroke (SAP)<sup>1,3</sup> and may, therefore, include microbiological etiologies associated with hospitalized CAP or HAP. Our study suggests that aerobic Gram-negative bacilli (eg, *K pneumoniae*, *E coli*, and *P aeruginosa*) and Gram-positive cocci (eg, *S aureus* and *S pneumoniae*) were associated with the majority of pneumonia complicating stroke when cultures were sent. A recent review suggested that ≈80% of hospitalized CAP were caused by *S pneumoniae*, *Mycoplasma pneumoniae*, *Chlamydia pneumoniae*, and *Haemophilus influenzae*.<sup>26</sup> The same review found that ≈50% of HAP was caused by *S aureus* and 35% by *Pseudomonas* species, *Klebsiella* species, *Escherichia* species, *Acinetobacter* species, and *Enterobacter* species.<sup>26</sup> The spectrum of identified organisms in our study seems to be more closely related to HAP than VAP or hospitalized CAP although antibiotic susceptibilities were not reported in most of the included studies. Furthermore, none of the included studies reported the results of investigations for viral or atypical pathogens, or even if they were done. Although our study is a comprehensive systematic literature research and collaboration within the PISCES group, our findings need to be interpreted with caution because of several inherent limitations.

First, there was marked heterogeneity between the included studies, which likely contributed to the variation in identified organisms and their relative contributions to pneumonia. For example, studies undertaken in the critical care environment might yield a higher proportion of organisms overall because of access to more direct sampling (eg, bronchoscopy or tracheal aspirate) and also more frequent causes because of organisms typical of HAP or VAP.

We could not identify any single factor that fully explained the high heterogeneity ( $I^2=99\%$ ). Although both stroke type

and our prospective risk categorization (Table 2) showed anticipated differences in pneumonia frequency, even within our very high risk category frequency ranged from 14% to 63% (Figure 2), and heterogeneity remained extremely high ( $I^2>90\%$ ) within each risk category. The asymmetry of the scatterplots seen in the funnel plot also reflects the high heterogeneity among studies, which was not corrected even when subgroup analysis was undertaken with pneumonia risk stratification (Figure I in the [online-only Data Supplement](#)). Sixty-six percent of the included studies in our review were deemed high or very high risk (Table I in the [online-only Data Supplement](#)) for developing pneumonia reflecting the higher frequency of pneumonia seen in this review as opposed to a previous systematic review.<sup>1</sup> Apart from patient selection, the overall high heterogeneity also reflects varying geographical location, different study designs, inclusion criteria, timing from stroke onset to sampling, and differences in standardized outcome definition for SAP. We were unable to differentiate pneumonia and causative pathogens for patients admitted from institutional environments, such as nursing homes, which may have also contributed to heterogeneity. Stroke registries, not routinely expected to collect and maintain data on microbiological cause, were also excluded unless specific mention was made about determining bacterial cause, which could have contributed to selection bias and heterogeneity. The variation in approach to diagnosis of pneumonia complicating stroke is well recognized,<sup>1</sup> and may, therefore, influence the threshold for sending microbiological samples, contributing to verification bias. Second, only one of the studies was primarily designed to identify the microbial cause of pneumonia complicating stroke,<sup>12</sup> whereas the remaining studies collected microbiological data when available within the context of their individual study objectives. It is reasonable to assume that the proportion of positive cultures could be higher if culture samples were sent systematically in all suspected cases of pneumonia. However, compared with other clinical settings (eg, CAP or VAP), consistently obtaining sputum samples in nonventilated stroke patients is challenging and alternative strategies (eg, bronchoscopy for VAP) are limited. Third, it was unclear among most studies as to when culture samples

were sent in relation to onset of stroke and suspicion or diagnosis of pneumonia. Although the majority of pneumonia in our review occurred within a week of stroke symptom onset, it was not possible to further explore microbiological etiologies in relation to timing of pneumonia relative to stroke symptom onset. Although one could hypothesize that organisms commonly associated with CAP are most likely causal in early SAP ( $\leq 72$  hours), and those associated with HAP causal in SAP beyond 72 hours, we were unable to confirm or refute this finding because of limited data in the individual studies. Our observations of an apparent low yield of CAP organisms, and higher yield of HAP organisms could be at least in part because of sampling bias beyond 48 to 72 hours after stroke onset. Fourth, microbiological methods used to collect sputum samples, number of specimens sent when pneumonia was diagnosed, delays in analyzing samples if any, and laboratory techniques used were inadequately reported. None of the studies used modern molecular-based polymerase chain reaction methods or urinary antigen testing (for organisms, such as *S. pneumoniae* and *Legionella pneumophila*). This may also contribute to differences in the observed frequencies of positive culture data and the apparent lack of atypical or viral causes. Interestingly, a recent study reviewing hospitalized patients identified  $\approx 22\%$  of CAP inpatients had viral pathogens (most commonly rhinovirus, 9% and influenza, 6%) implying that a viral cause of SAP in at least some individuals may be possible.<sup>30</sup> Finally, antibiotics preceding index stroke (especially for patients with chronic lung disease) may have influenced microbiological cultures. For example, in the PANTHERIS study (Preventive Antibacterial Therapy in Acute Ischemic Stroke), when sputum samples were analyzed, 36% of samples were positive for organisms in the placebo group as opposed to 9% in the prophylactic antibiotic group.<sup>15</sup> Although the numbers are too small to form further conclusions, prophylactic antibiotics administered to the participating patients with stroke in the 3 randomized controlled trials may have affected frequency of identified organisms.

It is important to emphasize the differences in frequency of pathogens identified in Asian compared with European or North American studies. For example, in a study of hospitalized CAP patients, *S. pneumoniae* appeared to have a lower frequency (13% versus 26%) and *Enterobacteriaceae* appeared to have a higher frequency (9% versus 2.7%) in Asian studies as compared with European studies.<sup>5</sup> Similarly, although limited comparison was possible in our study, the frequency of certain nosocomial pathogens appeared to be higher in Asia or Asia Pacific regions (Table IV in the [online-only Data Supplement](#)) in keeping with higher prevalence of hospital-acquired infection (15.5%) in comparison to Europe (7.1%) or the United States (4%).<sup>31</sup> However, this incongruity may be as a result of selection pressure on clinically relevant bacteria from differing prior antibiotic exposure of patients across continents, as well as possible implementation of pneumococcal vaccination programmes.

The scarce amount of available data on microbiological etiologies of pneumonia complicating stroke might also reflect the clinical routine, as suggested by a recent survey on German stroke units.<sup>4</sup> Treatment guidelines for pneumonia complicating stroke, nevertheless, should take into

account these commonly isolated organisms and also consider local surveillance data, community pathogens, and demographical variations, together with guidance from the World Health Organization global strategy for containment of antimicrobial resistance<sup>32</sup> when recommendations are being made for prescribing empirical antibiotic regimens. Anaerobes, although not identified in our study, are commonly seen in the upper airway mixed with oral flora and in the stomach and are often thought to be responsible for aspiration pneumonia. However, anaerobes are difficult to culture, and if aspiration is suspected, then broader spectrum antibiotics may be required.<sup>28</sup>

The risk of contamination with oral flora, low diagnostic yield (30%–40% sensitivity) with current diagnostic methods,<sup>31</sup> and delay in producing a positive result (at least 24–48 hours) often predisposes to initial broad-spectrum antibiotic prescriptions. An ideal diagnostic method would be more timely and sensitive to identifying pathogens. Polymerase chain reaction assays involving comprehensive molecular testing platforms significantly improve pathogen detection (87% versus 39%) in comparison to sputum culture (including viral pathogens) and provide results within 24 hours, which may help in initiation of pathogen-directed microbial therapy or a rapid de-escalation of broad-spectrum antibiotic therapy.<sup>29</sup> However, validating such technology still depends on a reliable microbiological reference standard (sputum analysis), which may limit its potential utility in nonventilated stroke patients.

## Conclusions

Our study demonstrates an evidence gap in appropriately designed studies that robustly identify microbiological etiologies in pneumonia complicating stroke. Although limited by small and heterogeneous study samples, this review suggests aerobic Gram-negative bacilli and Gram-positive cocci species are frequently associated with pneumonia complicating stroke. Difficulties in obtaining suitable sputum samples among nonventilated stroke patients and poor sensitivity of current diagnostic methods often result in broad-spectrum antibiotic prescriptions for pneumonia. Our study, however, supports the need for a consensus-based approach to antibiotic initiation and further targeted antibiotic treatment trials for enhanced antibiotic stewardship.

## Acknowledgments

We acknowledge Dr Ryan Keh (Speciality Trainee in Neurology, Salford Royal NHS Foundation Trust) and Dr Luciana Miguel Alhambra (Clinical Fellow in Stroke Medicine, Salford Royal NHS Foundation Trust) for their assistance in translating non-English language studies for this review.

## Sources of Funding

Dr Meisel is supported by the German Research Foundation (EXC257 and SFB-TR84).

## Disclosures

Dr Meisel received project funding by Thermo Fisher Scientific BRAHMS GmbH, Germany for a stroke trial. The other authors report no conflicts.

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**Microbiological Etiologies of Pneumonia Complicating Stroke: A Systematic Review**  
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*Stroke*. 2018;49:1602-1609; originally published online June 18, 2018;

doi: 10.1161/STROKEAHA.117.020250

*Stroke* is published by the American Heart Association, 7272 Greenville Avenue, Dallas, TX 75231

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Print ISSN: 0039-2499. Online ISSN: 1524-4628

The online version of this article, along with updated information and services, is located on the World Wide Web at:

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## **SUPPLEMENTAL MATERIAL**

### **Microbiological etiologies of pneumonia complicating stroke: A systematic review**

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Cover title: Bacterial etiology of pneumonia complicating stroke

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**Online only Table I: Search Terms**

<b>Search Areas</b>	<b>Thesaurus terms</b>	<b>Free Text Terms</b>
<b>MEDLINE</b>	Subject Search in MESH: exp *Cerebrovascular disorders/; exp *Pneumonia/;  [Limit to: Publication Year 1946-2016]	Stroke*,Pneumonia*, “respiratory tract infection*”, “chest infection*”  [Limit to: Publication Year 1946-2016]
<b>EMBASE</b>	Subject Search on Emtree: exp *Cerebrovascular disease/ exp *Pneumonia/;  [Limit to: Publication Year 1901-2016]	Stroke*, Pneumonia*, “respiratory tract infection*”, “chest infection*”  [Limit to: Publication Year 1901-2016]
<b>CINAHL</b>	Subject Search: exp *Cerebrovascular disorders/; exp *Pneumonia/;  [Limit to: Publication Year 1946-2016]	Stroke*,Pneumonia*, “respiratory tract infection*”, “chest infection*”  [Limit to: Publication Year 1901 to 2016]
<b>Cochrane Cochrane Central Register of Controlled Trials</b>	[Limit to: Publication Year 1980-2016]	All text: “stroke” “pneumonia”  Publication Year from 1980 to 2016, in Cochrane Reviews (Protocols only), Trials and Methods Studies (Word variations have been searched)

**Online only Table II: Eligibility criteria**

**Inclusion criteria:**

Age  $\geq$  18years

Fully published studies or abstracts

English or non-English language

Inpatients with ischemic stroke, intracerebral haemorrhage, or both

Randomized (RCT's) and other controlled trials, including cluster RCTs, controlled (non-randomized) clinical trials (CCTs) or cluster trials, prospective comparative cohort studies, retrospective observational studies, case-control or nested case-control studies.

Incidence or prevalence of pneumonia following admission with stroke reported

Possible microbiological etiologies identified as responsible for pneumonia complicating stroke OR objective criteria used for diagnosing pneumonia

**Exclusion criteria:**

Age < 18years

Exclusively intubated and mechanically ventilated patients

Exclusively pneumonia preceding index stroke

Case reports

**Online only Table III: Patient characteristics**

Author	NG or PEG feeding (%)	Prophylactic antibiotics	Co-morbidities (%)								MV (%)	
			COPD/other lung pathology	Immunosuppression	Previous Stroke	Smoking history	Heart disease	AF	HTN	DM		
<b>Chamorro<sup>1</sup></b>	48	Yes	9.5	Excluded	17.6	16.9	12.5	NR	63.2	22.8	0	
<b>Hassan<sup>2</sup></b>	NR	No	NR	NR	NR	NR	NR	NR	NR	NR	NR	0
<b>Vargas<sup>3</sup></b>	42	NR	10	4.3	16.1	16	NR	NR	60	19	0	
<b>Ros<sup>4</sup></b>	NR	NR	NR	NR	12.4	NR	13.2	12.8	63.2	35.3	0	
<b>Harms<sup>5</sup></b>	66	Yes	7.5	8.8	11.4	11.4	37.2	34.1	64.5	31.6	8.8	
<b>Sui<sup>6</sup></b>	47	NR	NR	NR	NR	49	32	NR	NR	44	2.2	
<b>Yeh<sup>7</sup></b>	73	No	NR	NR	NR	NR	NR	NR	NR	NR	12.2	
<b>Fluri<sup>8</sup></b>	NR	NR	NR	NR	NR	NR	34	19.4	80	19.3	0	
<b>Chen<sup>9</sup></b>	58	NR	NR	NR	NR	NR	NR	NR	82	39	0.3	
<b>Chen<sup>10</sup></b>	NR	NR	25.5*	14*	31.4*	43*	27.5*	NR	70.6*	29.4*	23.5*	
<b>Satou<sup>11</sup></b>	100	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
<b>Becker<sup>12</sup></b>	NR	No	NR	Excluded	35.3	37	24	NR	53.1	24	NR	
<b>Warusevitane<sup>13</sup></b>	100	No	16	NR	NR	NR	NR	60	66	28	0	
<b>Li<sup>14</sup></b>	NR	NR	NR	NR	NR	40	NR	NR	74	20	NR	
<b>Westendorp<sup>15</sup></b>	NR	Yes	8.2	17.3	32.5	24.4	13	15.4	55.2	19.7	0	

NR=Not Reported, COPD=Chronic Obstructive Pulmonary Disease, AF=Atrial fibrillation, HTN=hypertension, DM=diabetes mellitus, MV=mechanical ventilation, \* with pneumonia

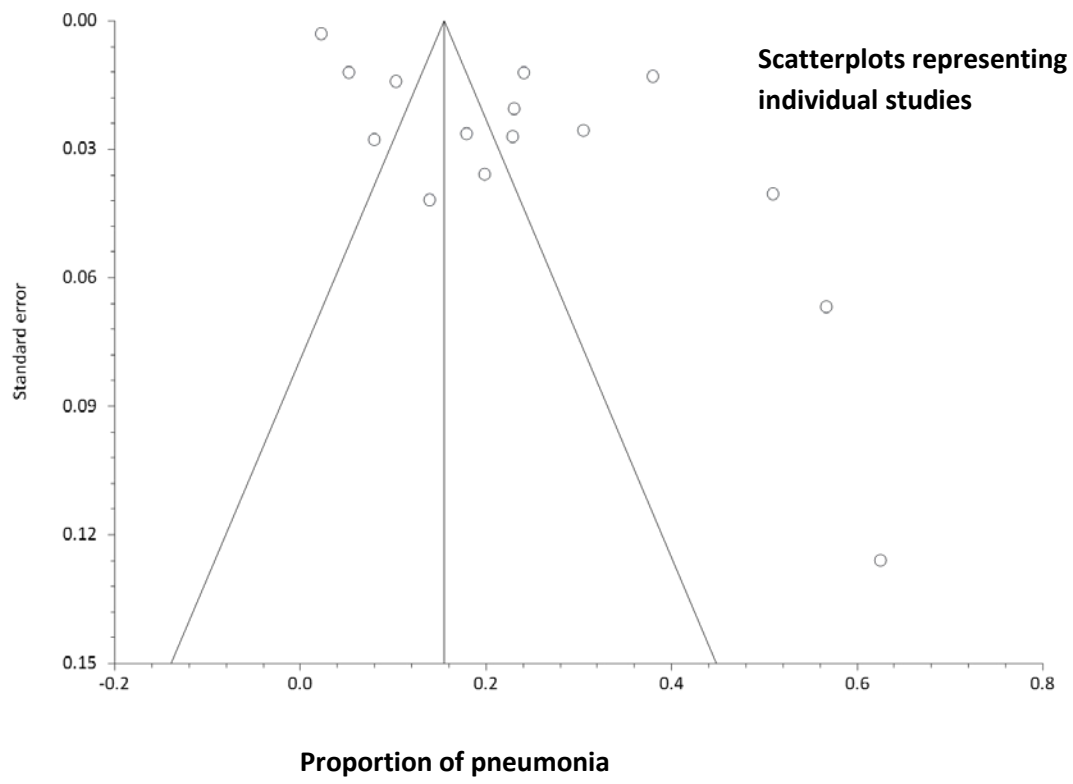
**Online only Table IV:** The eight most commonly isolated organisms in positive cultures (%) in the included studies

	European or North American Studies								Asian Studies						
	Chamorro	Vargas	Ros	Harms	Fluri	Becker	Warusevitane	Westendorp	Sui	Yeh	Li	Satou	Hassan	Chen	Chen
<b>GRAM POSITIVE COCCI</b>															
<i>Staphylococcus aureus</i>	11	×	5	36.3	5	22.2	11.7	1.7	9.3	19.2	6.8	20	11.7	24.1	1.9
<i>Streptococcal pneumoniae</i>	3.7	2.4	×	9	×	×	×	1.7	0.5	1.2	10.7	×	3.9	2.8	1
<b>GRAM NEGATIVE BACILLI</b>															
<i>Klebsiella pneumoniae</i>	×	×	5	×	×	11.1	11.7	×	8.8	51	18.2	10	3.9	15.4	23.5
<i>Escherichia coli</i>	×	7.3	×	×	×	×	5.8	×	7.7	9	21.7	×	1.9	2.8	10
<i>Enterobacter cloacae</i>	×	7.3	1.7	×	×	×	5.8	×	×	9.6	×	×	1.9	3.8	10
<i>Pseudomonas aeruginosa</i>	×	2.4	×	×	×	11.1	×	×	6.6	4.8	8.2	×	11.7	6.7	3.9
<i>Haemophilus influenzae</i>	3.7	21.9	×	×	5	11.1	×	×	6.6	×	×	×	3.9	11.5	7.8
<i>Acinetobacter baumannii</i>	×	×	×	×	×	×	×	×	1.4	22.9	8.4	×	3.9	6.7	7.8

For each study, data represent % of positive cultures due to the microbial species listed. X indicates not identified

**Online only Table V: Definition of pneumonia terminologies**

<b>Types of pneumonia</b>	<b>Definition</b>
<b>Hospitalized community acquired pneumonia (CAP)<sup>16</sup></b>	Pneumonia i.e. symptoms and signs consistent with an acute lower respiratory tract infection in the community, associated with new radiographic shadowing for which there is no other explanation and needing hospital admission
<b>Hospital acquired pneumonia (HAP)<sup>17</sup></b>	Pneumonia in a hospitalised patient > 48-72 hours after admission, without any incubation period
<b>Ventilator-associated pneumonia (VAP)<sup>17</sup></b>	Pneumonia in a patient occurring > 48-72 hours after intubation
<b>Stroke associated pneumonia (SAP)<sup>18</sup></b>	Pneumonia in a patient $\leq 7$ days from stroke symptom onset (HAP after this period); Early SAP < 72hrs, late SAP > 72hrs



**Online only Figure I:** Funnel Plot



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