





1 **Original Full Paper**

2
3 **Assessment of pulmonary lesions associated with Porcine Respiratory Disease Complex in**
4 **Colombian swine herds using slaughterhouse lung scoring systems**

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25
26 **Abstract**

27 Porcine Respiratory Disease Complex (PRDC) causes substantial economic losses in swine
28 production due to decreased efficiency and increased reliance on pharmaceuticals for disease control.
29 Diagnosis of PRDC can be achieved by evaluating lung lesions at slaughterhouses, which enables
30 assessment of herd health status and the formulation of targeted control strategies. In this study, the
31 MADEC and SPES scoring systems were employed to characterize lung lesions associated with
32 PRDC in pigs slaughtered in Colombia. A total of 16,223 red viscera (lungs, heart, and liver) from
33 15 commercially advanced farms were examined, utilizing the MADEC system for cranioventral
34 consolidations and the SPES system for dorsocaudal pleuritis. Additionally, histopathological
35 analyses and multiplex PCR tests were performed on randomly selected samples to identify
36 microscopic lesions and respiratory pathogens. Approximately 70% of diaphragmatic lobes on both
37 sides exhibited mild lesions (MADEC1), while severe lesions (MADEC3–4) were predominantly
38 observed in the cardiac and accessory lobes. The prevalence of pleuritis was 17.7%, with SPES types
39 1 and 2 being most common. *Glaserella parasuis* was the only pathogen detected in the analyzed
40 samples. Furthermore, concurrent use of up to five antimicrobial groups was documented across the
41 farms. In conclusion, the observed respiratory lesions likely reflect the widespread circulation of
42 PRDC agents, underscoring the need to enhance biosecurity, vaccination programs, and
43 epidemiological surveillance in Colombian swine production. The adoption of standardized systems,
44 such as MADEC and SPES, facilitates effective monitoring and control of respiratory diseases
45 throughout the production chain.

46 **Keywords:** Porcine Respiratory Disease complex, lung pathology, abattoirs, antimicrobial use,
47 epidemiological surveillance.

48

49 **Introduction**

50

51 One of the most critical aspects of analyzing animal health plans in animal production systems
52 is determining lesion findings in slaughterhouses, which can assess the suitability of carcasses for

53 human consumption and provide feedback for epidemiological investigations of infectious agents
54 present on the farm, as well as the quality of animal health plans. In swine production systems, it is
55 essential to evaluate lesions associated with Porcine Respiratory Disease Complex (PRDC), a
56 multifactorial disease that primarily affects swine farms worldwide. PRDC causes significant
57 economic losses for the industry by adversely affecting production parameters. Additionally, it
58 increases the cost due to the use of antimicrobial drugs for controlling bacterial infections, as well as
59 vaccines and other biosecurity measures. Only a small number of pigs raised under commercial
60 conditions are slaughtered without pulmonary lesions (11, 17, 28).

61 To assess pulmonary lesions and other organs, as well as their association with various
62 pathogens, including viral and bacterial infections, in pigs at the slaughterhouse, various approaches
63 have been developed. These approaches are known as systems for evaluating and monitoring
64 pulmonary lesions, some of which were proposed by various authors (10, 12, 13, 15, 33, 35). These
65 systems help to determine the incidence and severity of injuries caused by respiratory diseases in
66 pigs, such as enzootic pneumonia, atrophic rhinitis, pleurisy, and pleuropneumonia (27).

67 The most commonly used classification systems are MADEC and SPES. The Slaughterhouse
68 Pleurisy Evaluation System (SPES) assigns a score for dorsocaudal pleurisy based on the extension
69 and localization of pleural adherences associated primarily with *Actinobacillus pleuropneumoniae*
70 infections (7). Otherwise, the MADEC system scores the lesion as cranioventral consolidation,
71 characterized by gray-to-purple coloration and associated with *Mycoplasma hyopneumoniae*
72 infection (20). One of the most common lesions observed in the analysis of meat carcasses in
73 slaughterhouses is pneumonia and pleurisy, with prevalences oscillating between 19% and 79% and
74 between 3.8% and 62%, respectively, according to the country and the system used (25).

75 Although these systems are valuable, it is essential to consider their limitations, as they may
76 overlook subclinical lesions or localized lesions in deep organs. Additionally, the sensitivity to
77 punctuation can vary among veterinary professionals, depending on their level of expertise. However,
78 the early detection and characterization of lesions in the slaughterhouse contribute to a better

79 understanding of the epidemiology of PRDC, enabling the implementation of more effective control
80 measures, reducing economic losses, and improving animal welfare (17, 20).

81 In Colombia, there are no prior studies analyzing pulmonary lesions associated with PRDC in
82 pig slaughterhouses from commercial pig farms. Therefore, the objective of this study was to evaluate
83 pulmonary lesions in pigs at slaughter in Colombia using the MADEC and SPES scoring systems.

84

85 **Material and Methods**

86

87 *Farms' characteristics and inspection of viscera at the slaughterhouse*

88 A total of 16,223 red viscera (lungs, heart, and liver) were inspected over a six-month period
89 (January to July 2019) in a private slaughterhouse in Bogotá, Colombia. The animals originated from
90 15 farrow-to-finish farms located in the “central region” of the country, comprising the departments
91 of Boyacá, Cundinamarca, Meta, and Tolima. All farms used antimicrobials and vaccination
92 programs against bacterial pathogens, including *Mycoplasma hyopneumoniae*, *Actinobacillus*
93 *pleuropneumoniae*, and *Glaserella parasuis*, as well as viral agents such as Porcine Circovirus Type
94 2 (PCV2) and Porcine Parvovirus Type 1 (PPV1). Data on antimicrobial use were collected from the
95 participating farms through structured records provided by farm managers. Information included the
96 types of antimicrobial groups used to control respiratory diseases during the production cycle.
97 Antibiotics were classified into pharmacological groups for subsequent descriptive and co-occurrence
98 analyses. Post-mortem inspection was conducted along the slaughter line in the processing area using
99 standard procedures, including visual examination as an initial screening step to identify gross
100 pathological alterations, an approach associated with a reduced risk of carcass contamination (11).

101

102 *Macroscopic inspection*

103 The lungs inspected were evaluated using the MADEC system for cranioventral consolidation
104 (used for Enzootic Pneumonia), which estimated the proportion (%) of lung parenchyma affected and

105 its extension. The classification was MADEC1: <25%, MADEC2: >26-50%, MADEC3: 51-75% and
106 MADEC4:>75% (10). The SPES system, as described by Dottori et al. (7), involves assigning a score
107 to each lung based on the presence, extent, and location of adhesions (33). Lesions can be observed
108 in the diaphragmatic lobes and require at least three months to resolve; therefore, their disappearance
109 before arrival at the slaughterhouse is doubtful (8). This classification system includes five different
110 scores, determined as follows: 0, absence of chronic pleuritis; 1, cranioventral lesion—pleural
111 adhesion between lobes or on the ventral edge of the medial and diaphragmatic lobes; 2, focal
112 unilateral dorsocaudal lesion; 3, bilateral lesion or extensive unilateral lesion affecting at least one-
113 third of a diaphragmatic lobe; and 4, severely extensive bilateral lesion affecting more than one-third
114 of both diaphragmatic lobes (20). Pleuritis observed at the slaughterhouse with scores between 2 and
115 4 is a lesion indicative of infections associated with *Actinobacillus pleuropneumoniae* (21).

116

117 *Microscopic evaluation*

118 For histopathological analysis, eleven samples were selected based on the presence and severity
119 of macroscopic lesions observed during slaughterhouse inspection, aiming to include representative
120 cases of different lesion patterns. The number of samples analyzed was influenced by logistical and
121 budgetary constraints associated with laboratory processing. Fragments of tissue of 2 cm³, with both
122 healthy and injured tissue present, were fixed in 10% buffered formalin and sent to the SERVIPAT®
123 pathology laboratory in Bogotá, D.C., Colombia, where they were processed using routine protocols,
124 with sections of 4 µm that were stained with Hematoxylin and Eosin (HE) and analyzed
125 microscopically according to the method described by Karabasil et al. (14).

126

127 *Molecular Detection of Infectious Agents*

128 From 11 samples, three samples of 10 grams from three randomly selected samples with severe
129 MADEC lesions (MADEC4) were selected for molecular analysis using quadruplex PCR to assess
130 the presence of selected bacterial pathogens associated with PRDC (*Actinobacillus*

131 *pleuropneumoniae, Actinobacillus suis, Glaserella parasuis, Streptococcus suis*). The limited number
132 of samples processed was determined by logistical and budgetary constraints. Samples were packed
133 in a sterilized vacuum bag and frozen at -80°C until they were sent at 4°C in a cooler to the Molecular
134 Biology Laboratory from the Diagnostic Unit of the Faculty of Agricultural Sciences at the University
135 of Antioquia. A commercial kit, BIONEER AccuPower Porcine Respiratory 4-Plex PCR®, was used.
136 A total of 4 µL of DNA was added to the PCR mix and submitted to the following protocol: initial
137 denaturation at 95°C for 5 minutes(min), followed by a second denaturation at 95°C for 30 seconds,
138 annealing at 65°C for 1 min, and extension at 72°C for 5 min, for 35 cycles. Finally, three µL of the
139 amplified product were loaded onto a 2% TBE agarose gel (4).

140

141 *Statistical analysis*

142 Descriptive statistics were used to present the absolute and relative frequencies of lesions and
143 their distribution across sides and lobes within each system. Correlational analyses were also
144 conducted for the MADEC system to assess whether the distribution of lesion categories (MADEC1–
145 MADEC4) was independent of side (Left vs. Right). A 2×4 contingency table (side × MADEC
146 category) was created, and Pearson's chi-squared test of independence was applied. Similarly, for
147 each side, a contingency table (Lobe × MADEC category) was generated (3 × 4 for the Left; 4 × 4
148 for the Right), and Pearson's chi-squared test was applied using Fisher's Monte Carlo method.

149 To identify differences between pairs of lobes, paired post-hoc comparisons with Bonferroni
150 correction and Monte Carlo simulation were conducted. By grouping the flank and lobe into a single
151 variable, “Complete Lobe” (7 levels), its independence from the MADEC category was assessed
152 using Pearson’s chi-squared test, again employing Monte Carlo simulation for low cell counts. To
153 further explore which lobes differed significantly in the distribution of MADEC categories, paired
154 post-hoc comparisons between lobes (2×4 tables) with Bonferroni correction were performed. To
155 estimate the probability of each MADEC category according to anatomical location (flank and lobe),
156 a multinomial logistic regression model was fitted, which is appropriate for response variables with

157 more than two categories and no strict numerical order. This regression estimates the probabilities of
158 belonging to each MADEC category relative to the base category (MADEC1).

159

160 **Results**

161

162 A total of 16,223 red viscera (comprising lungs, heart, and liver) were inspected over six months
163 in 2019. The primary lesions found were bronchopneumonia (9,381), pulmonary pleurisy (2,781),
164 pulmonary abscess (207), and interstitial pneumonia (133). Otherwise, 554 red viscera had
165 polyserositis (pleurisy, pericarditis, and hepatic adherences); additionally, 787 hearts had pericarditis,
166 and 146 livers had adherence, and 2,234 inspected viscera had no apparent pathological changes.

167 Due to the severity of the lesions, and in accordance with sanitary normativity (22, 23), a total
168 of 1,326 red viscera were condemned by national sanitary authorities during the study. The most
169 common causes of condemnation were pulmonary pleurisy, pericarditis, and hepatic adhesions, and
170 they originated from Cundinamarca, Tolima, and Boyacá.

171

172 *MADEC System*

173 The MADEC system was employed across 15,727 pulmonary lobes, and the overall results are
174 presented in Table 1. The most common distribution pattern was <25% (MADEC 1) in 70% of
175 pulmonary lobes bilaterally, followed by MADEC 2 (22.5% in the right lung and 21.1% in the left
176 lung). More serious scores (MADEC3 and MADEC4) are present at low proportions (MADEC3:
177 Right 5.1%, Left 8.0%; MADEC4: Right 2.4%, Left 0.8%). Figure 1 shows the percentage
178 distribution of pulmonary lesions by the flank and lobe evaluated. MADEC1 lesions show a higher
179 prevalence, primarily in the right (95.1%) and left (98.7%) diaphragmatic lobes. Additionally,
180 MADEC2 showed similar prevalence across cardiac lobes bilaterally (27.1% on the right and 25.5%
181 on the left).

182 This predominance indicates significant heterogeneity in the distribution of severity scores
183 between lobes for each flank. There were statistically significant differences between the lobes on the
184 same side ($p = 0.0099$ for the left side and $p = 0.0216$ for the right side). For the left lung, the severity
185 of lesions evaluated is as follows: Apical/Cardiac, $p = 0.00423$; Apical/Diaphragmatic, $p = 0.003$;
186 and Cardiac/Diaphragmatic, $p = 0.003$. For its part, on the right side, the results were as follows:
187 Accessory/Apical: $p = 0.00104$, Accessory/Cardiac: $p = 0.00259$, Accessory/Diaphragmatic: $p =$
188 0.00905 , Apical/Cardiac: $p = 0.00135$, Apical/Diaphragmatic: $p = 0.00980$, and
189 Cardiac/Diaphragmatic: $p = 0.00198$.

190 The Heatmap (Fig. 2) indicates that MADEC1 lesions were most probably in all lobes,
191 especially in the left (96.5%) and right (96.3%) diaphragmatic lobes. MADEC2 and MADEC3 were
192 most likely in the right accessory lobe (32.6% and 4.5%, respectively) and the left cardiac (25.3%
193 and 10.9%). MADEC4 was most likely in the right apical lobe (9.3%). Multiplex PCR analysis was
194 performed on three lung samples with severe MADEC lesions (MADEC4); among these samples,
195 *Glaserella parasuis* was detected, whereas the other targeted pathogens in the assay were not
196 identified. This pattern suggests that diaphragmatic lobes are more likely to experience mild injuries,
197 whereas the right accessory and left cardiac lobes have a greater likelihood of severe injuries. These
198 findings align with the observed anatomical distribution and previous statistical analyses.

199

200 *SPES System*

201 In contrast, the SPES system enabled analysis of the distribution of pleuritis types, which are
202 typically associated with infections caused by *Actinobacillus pleuropneumoniae* infection (Fig. 3).
203 Type 2 pleuritis was the most prevalent, affecting 35% of cases (978 out of 2.781 lungs), followed
204 by types 1, 3, and 4.

205

206 *Antibiotic therapy*

207 Antibiotic use patterns for the control of respiratory diseases on farms are summarized in
208 Figures 4 and 5. Overall, usage frequencies were similar across most groups, with
209 Fluoroquinolones/Quinolones used on 11 out of 15 farms. Beta-lactams/Aminopenicillins and
210 Pleuromutilins were each used on seven farms, while Phenicol and Macrolides were each used on
211 six farms. The least commonly used antibiotics were Tetracyclines (4/15), Miscellaneous antibiotics
212 (2/15), and combined Sulfonamides (1/15) (Fig. 4). Usage patterns per farm, which indicate the
213 simultaneous use of the eight antimicrobial groups, are shown in Figure 4. Only two farms relied on
214 a single pharmacological group for respiratory disease control, while the remaining farms use
215 between two and five different antimicrobials concurrently. Based on these results, a co-occurrence
216 analysis was conducted to identify common drug combinations across all farms (Fig. 5). Notably, the
217 most frequent combinations involved Fluoroquinolones/Quinolones, particularly their use with
218 Macrolides, as well as combinations with Tetracyclines, Phenicol, Pleuromutilins, and Beta-
219 lactams/Aminopenicillins (four farms). Three farms combined Fluoroquinolones/Quinolones with
220 Sulfonamides and miscellaneous antimicrobials (Fig. 5).

221

222 *Histopathologic results*

223 Of the 11 tissue samples submitted for histopathologic examination, three results are shown.
224 Macroscopic and microscopic (histopathologic) images from SPES4 (Fig. 6A), SPES2 (Fig. 7A), and
225 MADEC2 (Fig. 8A) lung samples collected during the study are shown. Histologic sections stained
226 with Hematoxylin & Eosin reveal bronchiolar epithelium exfoliation (SPES4, 40×) (Fig. 6B), and
227 pulmonary edema (SPES4, 100×) (Fig. 6C), moderate activation of BALT (SPES2, 40×) (Fig. 7B),
228 and mild pulmonary congestion (MADEC2, 40×) (Fig. 8B). These findings are consistent with
229 nonspecific inflammatory and circulatory alterations and are not pathognomonic of specific
230 etiological agents such as *Mycoplasma hyopneumoniae* or *Actinobacillus pleuropneumoniae*. Among
231 the remaining samples, three exhibited severe, multifocal suppurative bronchointerstitial
232 pleuropneumonia, accompanied by marked activation of bronchus-associated lymphoid tissue

233 (BALT). Four samples showed incipient or mild multifocal mixed bronchopneumonia with moderate
234 BALT activation, while one sample presented chronic polyserositis. Overall, these findings are
235 consistent with inflammatory processes affecting the respiratory system; however, they are not
236 pathognomonic and do not allow definitive attribution to specific etiological agents.

237

238 **Discussion**

239

240 Studies investigating lung lesions in slaughtered pigs are diverse, underscoring the value of
241 such analyses for evaluating the effectiveness of health programs on pig farms. To the authors'
242 knowledge, this is the first study of its kind conducted in Colombia. While global research has
243 documented the prevalence of MADEC-type lesions in slaughtered pigs, no studies have
244 characterized the anatomical distribution of these lesions within individual lobes or assessed whether
245 specific lobes are more closely associated with certain degrees of injury. Our study, therefore,
246 provides the first such patterns described in Colombia and is among the earliest worldwide. Predicted
247 probabilities for MADEC1 and MADEC2 lesions were similar in the apical and cardiac lobes on both
248 sides. For more severe lesions (MADEC 3 and 4), this pattern persisted bilaterally in the cardiac lobes
249 and in the accessory lobe. These findings confirm that lung involvement is not uniform across lobes
250 and suggest an increased susceptibility to severe lesions in certain lobes, possibly due to host factors
251 (anatomical or physiological) or pathogen factors (including pathogenicity, virulence, and
252 mechanisms of transmission).

253 Although molecular analysis was limited in this study, the observed cranioventral pattern of
254 lung lesions is consistent with those commonly associated with airborne transmission of *M.*
255 *hyopneumoniae* (32). However, this interpretation should be made with caution, as direct etiological
256 confirmation was not established. Among the small number of samples analyzed by PCR, *Glaserella*
257 *parasuis* was detected; this finding is exploratory and cannot be used to infer the absence of other
258 PRDC-associated pathogens, but may suggest an opportunistic infection pattern, as previously

259 described by Zhang et al. (36). It is important to note, however, that the inability to detect certain
260 microorganisms (as occurred in this study) may reflect the multifactorial nature of these lesions,
261 influenced by a combination of biological, immunological, environmental, and management factors
262 that affect pathogen epidemiology, pathogenesis, virulence, and ultimately, their detectability. Also,
263 it may be due to sampling limitations or to potential nucleic acid degradation in slaughterhouse-
264 derived tissues, rather than to their true absence in the evaluated farms (1, 5). Therefore, further
265 investigation is required to better understand the role of coinfections and pathogen interactions in
266 lesion development. In some instances, pigs may recover from acute disease but retain anatomical
267 lesions detectable at slaughter, even in the absence of identifiable microbial agents (20).

268 De Conti et al. (6), using the MADEC methodology in Brazil, found that the central pulmonary
269 lesion is cranioventral consolidation, with an average MADEC score ranging from 1.58 to 2.83 (mild
270 to moderate lesions), like the results obtained in this study; however, *Pasteurella multocida* was
271 identified as the predominant causative agent of lesions, in coinfection with Influenza A virus and *M.*
272 *hyopneumoniae*. Globally, studies such as Pallarés et al. (24) have found that the predominant
273 pulmonary lesions in Portugal and Spain, in descending order, were bronchointerstitial pneumonia,
274 suppurative bronchopneumonia, fibrinous bronchopneumonia, interstitial pneumonia, and pleuritis,
275 findings consistent with those of the present study. Similarly, Przyborowska-Zhalniarovich et al. (29)
276 reported bronchopneumonia as the most common lesion (57.8%), with an average MADEC score of
277 1.74, which aligns with the results observed in this study.

278 Despite these similarities, notable differences exist. In this study, all lungs examined exhibited
279 some degree of MADEC-type lesion, with mild lesions (MADEC1: 70%) being the most common.
280 Conversely, Przyborowska-Zhalniarovich et al. (30) found that only 45.1% of pig lungs in Poland
281 showed lesions, with the majority being severe (MADEC 2, 3, and 4), collectively accounting for
282 94.57%. Kuberka et al. (16) also noted that lung lesions at slaughter were common (73.5% with mild
283 to moderate lesions) and primarily due to multifactorial microbial causes. Differences between studies
284 may stem from varying management practices, environmental factors, and seasonality. Notably, the

285 fact that all evaluated pulmonary lobes showed some degree of lesion highlights systemic challenges
286 in Colombian pig production. This widespread occurrence may reflect deficiencies in farm
287 management, biosecurity, and disease prevention, increasing the risk of PRDC. These findings
288 underscore the need for targeted research to identify local risk factors and implement effective
289 preventive strategies. Expanding research on this topic within Colombia is crucial for understanding
290 the dynamics of respiratory disease in national production systems and for developing effective
291 prevention and control strategies for PRDC. Additionally, it is important to acknowledge that lesion
292 assessment at slaughterhouses involve a degree of subjectivity, which may introduce observer-related
293 variability, particularly in the classification of mild lesions; this should be considered in future
294 studies.

295 Otherwise, the overall occurrence of pleuritis-type lesions, as assessed using the SPES system,
296 was 17.68%, with SPES types 1 and 2 lesions being the most prevalent. This prevalence exceeds that
297 reported in studies from Brazil—Arruda et al. (2) 11.3%, Galdeano et al. (9) 13.1%, Baraldi et al. (3)
298 9%, and Petri et al. (26) 12.3%—and Poland, where Kuberka et al. (16) observed an overall
299 prevalence of 12.1% and an average score of 1.96. Regarding the SPES system’s utility for lesion
300 analysis and clinical correlation, Sipos et al. (34) found that pulmonary scores may not accurately
301 reflect field clinical situations in porcine pleuropneumonia. Their research indicated that SPES results
302 were higher during the subacute/chronic phases than during the acute/peracute phases. This finding
303 also underscores the importance of effective vaccination protocols for reducing lung lesions at the
304 herd level.

305 Alawneh et al. (1) found that, according to the SPES system, the overall prevalence of pleuritis
306 in pigs in the Philippines was 22%, a result slightly higher than that reported in this study. They also
307 identified associated risk factors, including the lack of swine cholera vaccination programs, pigs
308 originating from commercial farms rather than small-scale farms, and the presence of another pig
309 farm within 500 meters. All of this highlights the need to establish longitudinal epidemiological
310 studies on pig farms across different departments to determine risk factors, disease presentation

311 dynamics, and their direct impact on national productivity, and to combine these with analyses at the
312 slaughterhouse level. Economic analyses of the effect of pleuritis indices in pigs at slaughter revealed
313 that pigs with a higher degree of lesion incurred an additional cost of USD 1.29 per kilogram,
314 compared to USD 1.32 for milder cases. Furthermore, total revenue decreased by 1.36%, and the
315 return on investment dropped from 5.33% to 3.90%, primarily due to reductions in average daily
316 weight gain (18).

317 Finally, regarding antibiotic use during the finishing stage on the surveyed pig farms, up to five
318 different drugs were used within the production systems, a trend that appears to be common globally.
319 Mallioris et al. (19) reported that the primary reasons for antimicrobial use varied according to
320 production stage, with musculoskeletal/neurological diseases (62.9% of farms) and respiratory
321 diseases (52.3%) being the main drivers. The most frequently used pharmacological groups were
322 tetracyclines, macrolides, lincosamides, and trimethoprim-sulfonamides, consistent with the findings
323 of this study. Patterns of use among different subpopulations are crucial for understanding these
324 dynamics. Additionally, the frequent use of these drugs may reflect the need for more frequent
325 treatment, longer treatment durations, or higher doses (19).

326 In conclusion, lung lesions in pigs at slaughter were persistent. Most lobes showed some degree
327 of involvement according to the MADEC scale, with mild lesions predominating (MADEC1: 70%).
328 Pleuritis (SPES) had a significant prevalence (17.7%), with SPES categories 1 and 2 being the most
329 common. A cranioventral pattern was identified for mild/moderate lesions, and a concentration of
330 severe lesions (MADEC3–4) in the cardiac and accessory lobes. The predominant detection of
331 *Glaserella parasuis* in the submitted samples suggests its possible involvement in severe lesions,
332 although the small number of molecular samples limits etiological conclusions. The frequent and
333 concomitant use of several antimicrobials on the evaluated farms poses risks to therapeutic efficacy
334 and the selection of resistance. Longitudinal studies are recommended.

335

336 **Data Availability**

337 All the original contributions presented in this study are included in the article/supplementary
338 material. Further inquiries can be directed to the corresponding author.

339

340 **Author Contributions**

341 **Paula Andrea Heredia Beltrán:** Investigation, Data curation, Formal analysis, Writing
342 original draft. **Miguel Andrés Soler Uribe:** Conceptualization, Methodology, Formal analysis,
343 Writing – original draft preparation. **Victor Manuel Acero Plazas:** Conceptualization, Formal
344 analysis, Supervision, Writing – review and editing. **Luis Edgar Tarazona Manrique:** Formal
345 analysis, Visualization, Writing – review and editing. All authors have read and approved the final
346 version of the manuscript.

347

348 **Conflict of Interest**

349 The authors declare no competing interests.

350

351 **Generative AI Use Statement**

352 The authors did not use generative artificial intelligence tools or technologies in creating or
353 editing any part of this manuscript.

354

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359

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- 479

480 **Tables**

481

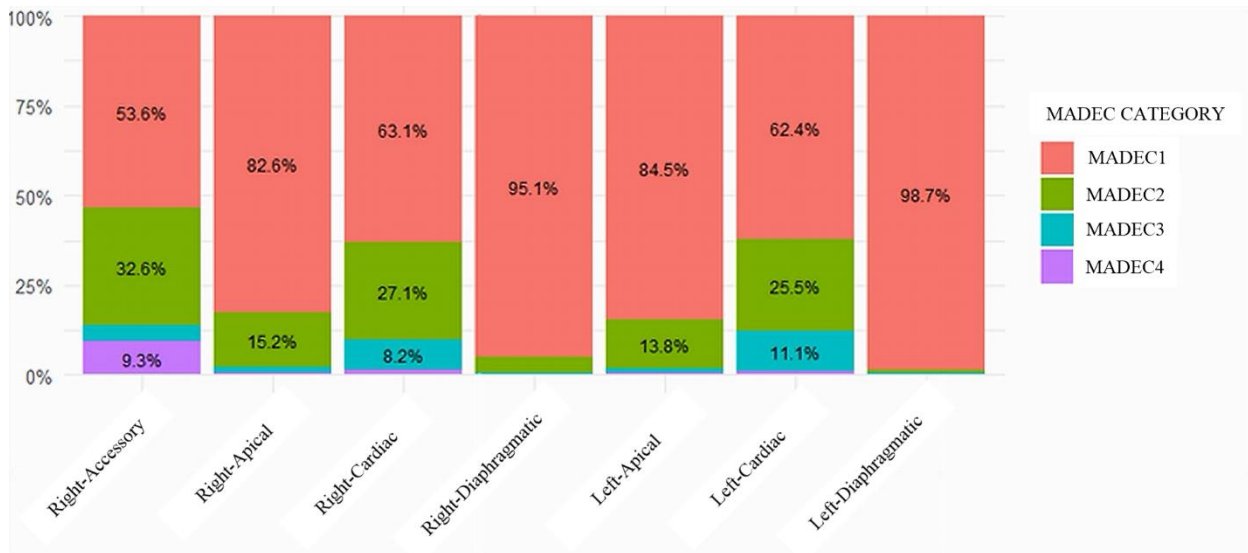
482 **Table 1.** Macroscopic lesions distribution during inspection in a slaughterhouse using the MADEC

483 system.

Lobe	Side	Number of samples	Number of simples according to the MADEC system result			
			1	2	3	4
			<25%	26-50%	51%-75%	>75%
Apical	Left	1,548	1,308	214	17	9
Cardiac	Left	4,606	2,875	1,176	510	45
Diaphragmatic	Left	448	442	3	2	1
Apical	Right	2,541	2,099	385	38	19
Cardiac	Right	4,319	2,727	1,169	356	67
Diaphragmatic	Right	843	802	35	4	2
Accessory	Right	1,423	763	464	64	132
Total (n)		15,728	11,016	3,446	991	275
(%)			(70%)	(22%)	(6%)	(2%)

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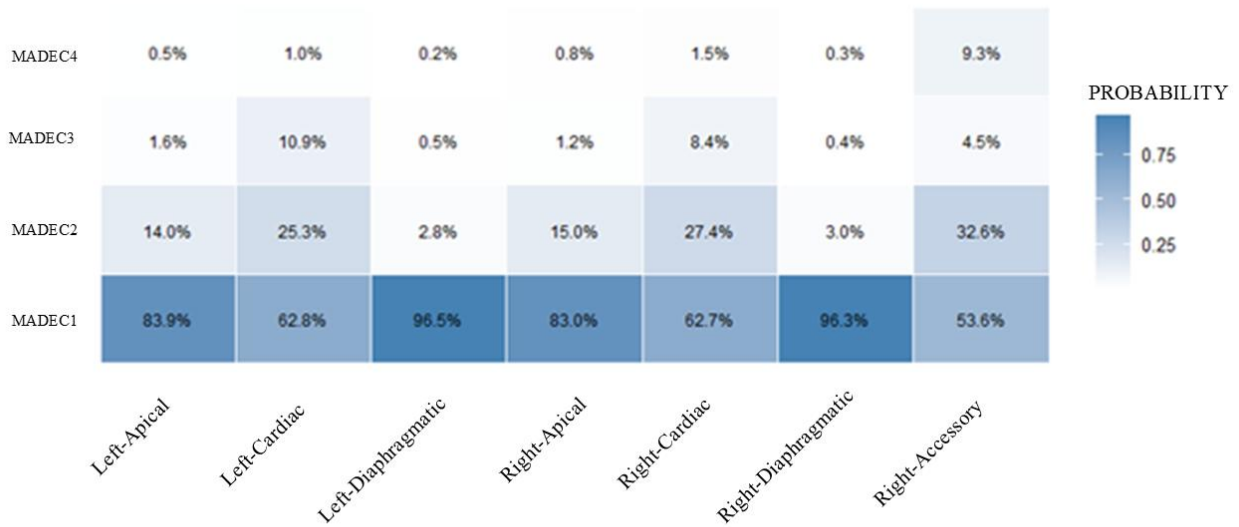
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487 **Figure 1.** Percentage distribution of MADEC lesions for each pulmonary lobe according to the side

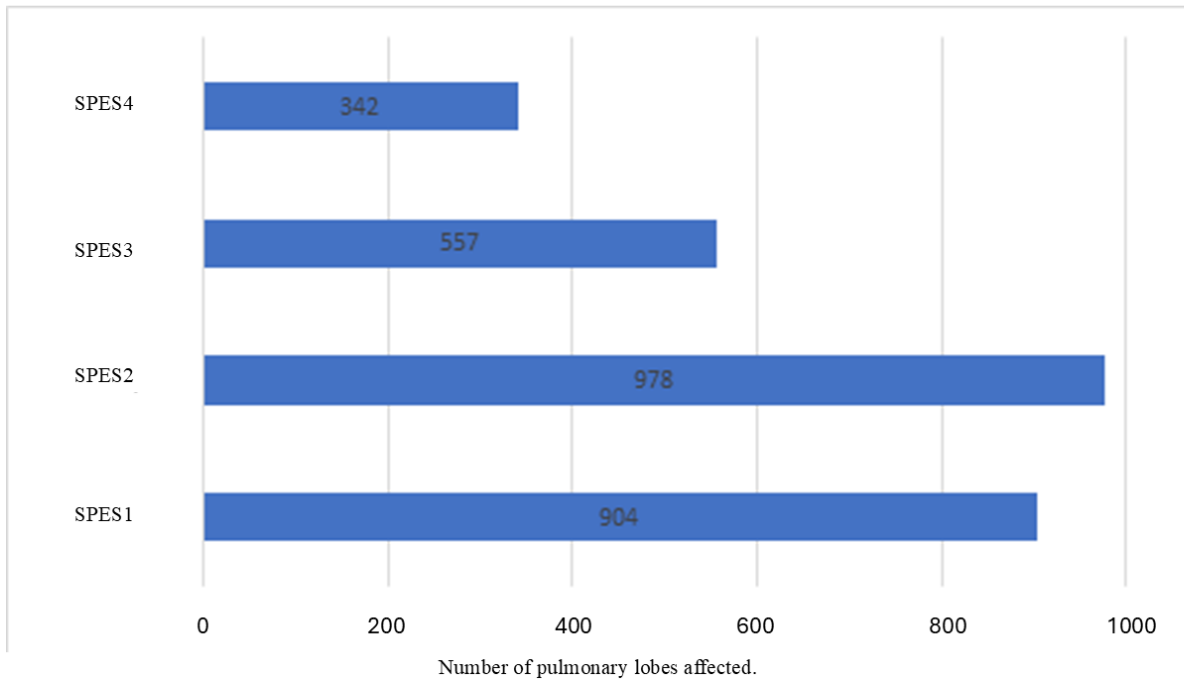
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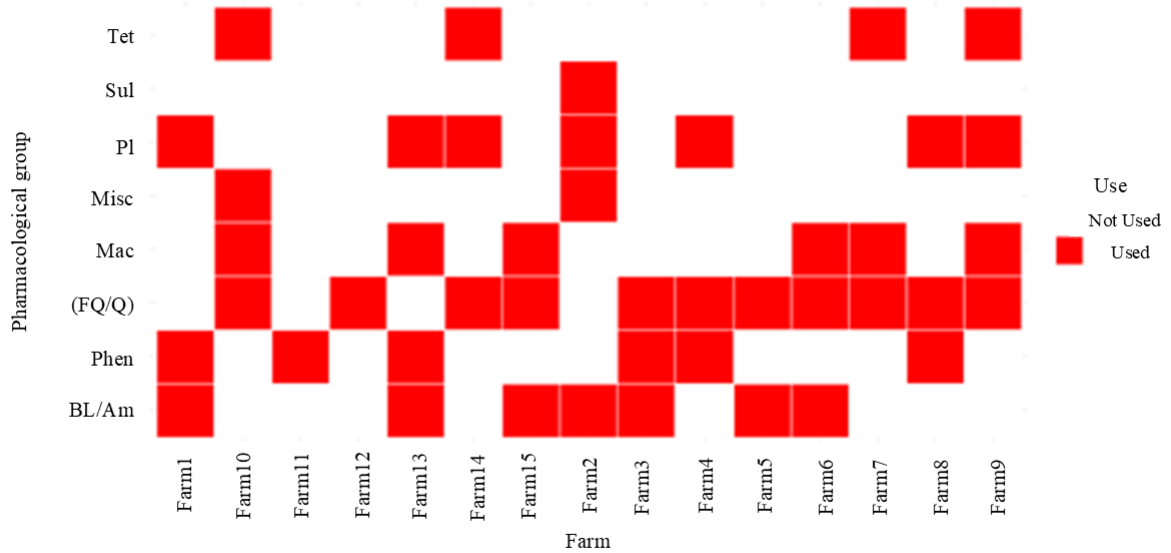
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490 **Figure 2.** Probability distribution for the presence of MADEC lesions for each lobe evaluated.

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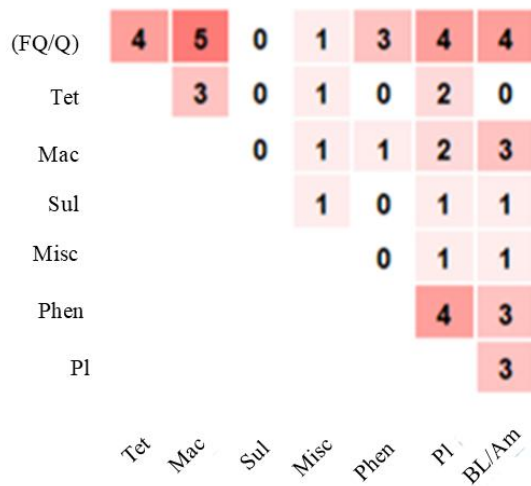


492
 493 **Figure 3.** Lesion distribution according to the type of SPES lesion found
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495
 496 **Figure 4.** Distribution patterns of antibiotic usage on commercial pig farms included. Note:
 497 Fluoroquinolones/Quinolones: (FQ/Q); Beta-lactams/Aminopenicillins: (BL/Am); Tetracyclines
 498 (Tet); Miscellaneous (Misc); Sulfonamides (Sul); Macrolides: (Mac); Phenicols: (Phe);
 499 Pleuromutilins (Pl).

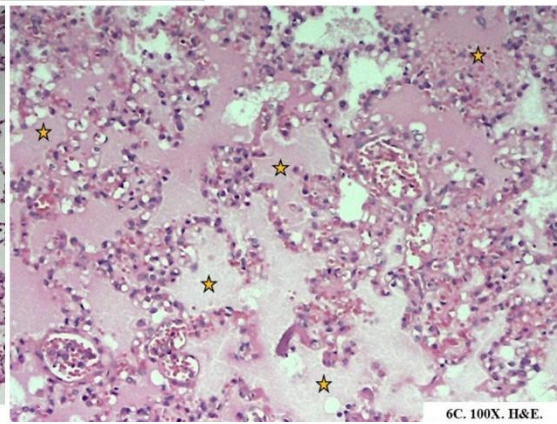
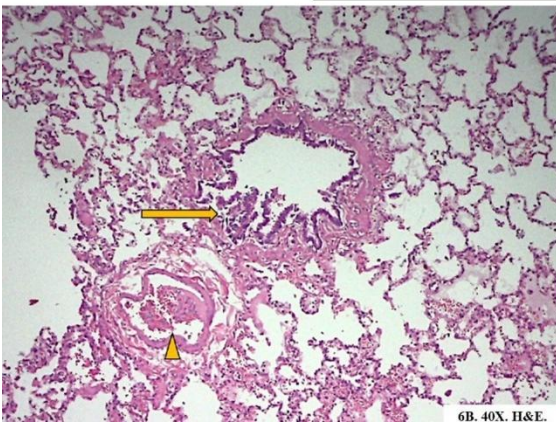
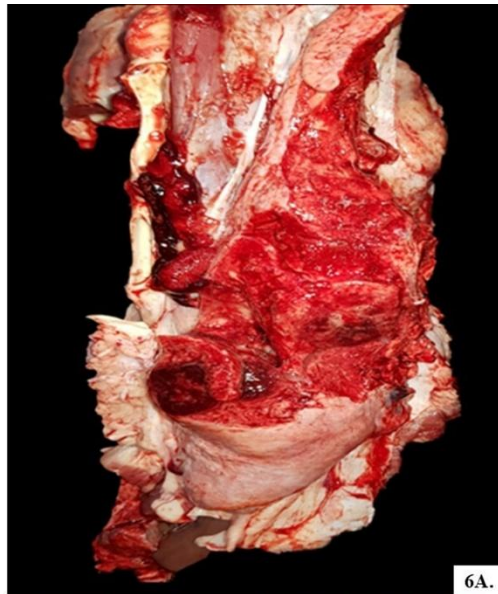
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502 **Figure 5.** Heatmap for the co-occurrence of pharmacological groups in pig farms. Note:
503 Fluoroquinolones/Quinolones: (FQ/Q); Beta-lactams/Aminopenicillins: (BL/Am); Tetracyclines
504 (Tet); Miscellaneous (Misc); Sulfonamides (Sul); Macrolides: (Mac); Phenicols: (Phe);
505 Pleuromutilins (Pl).

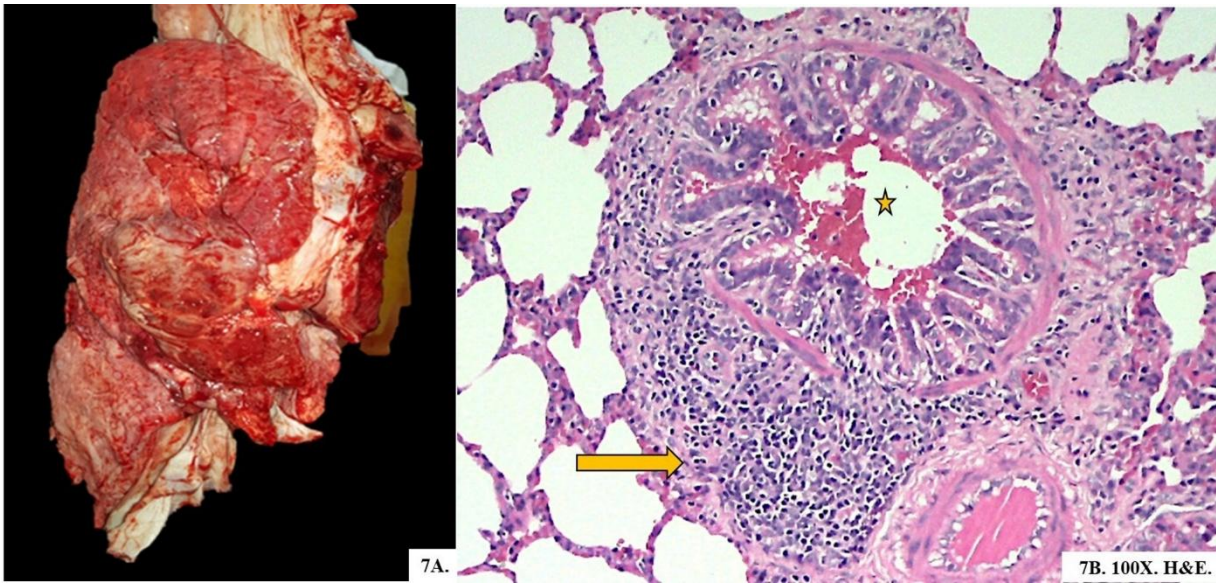
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508 **Figure 6.** Macroscopic and microscopic (histopathologic) images from a SPES4 sample collected
509 during the study. 6A. Gross image of the lung showing lesions corresponding to a SPES4 score. 6B.
510 Lung tissue section at 40× magnification (Hematoxylin & Eosin stain). The arrow indicates
511 exfoliation of the bronchiolar epithelium, and the arrowhead indicates vascular congestion. 6C. Lung
512 tissue section at 100× magnification (H&E stain). The star indicates pulmonary edema.

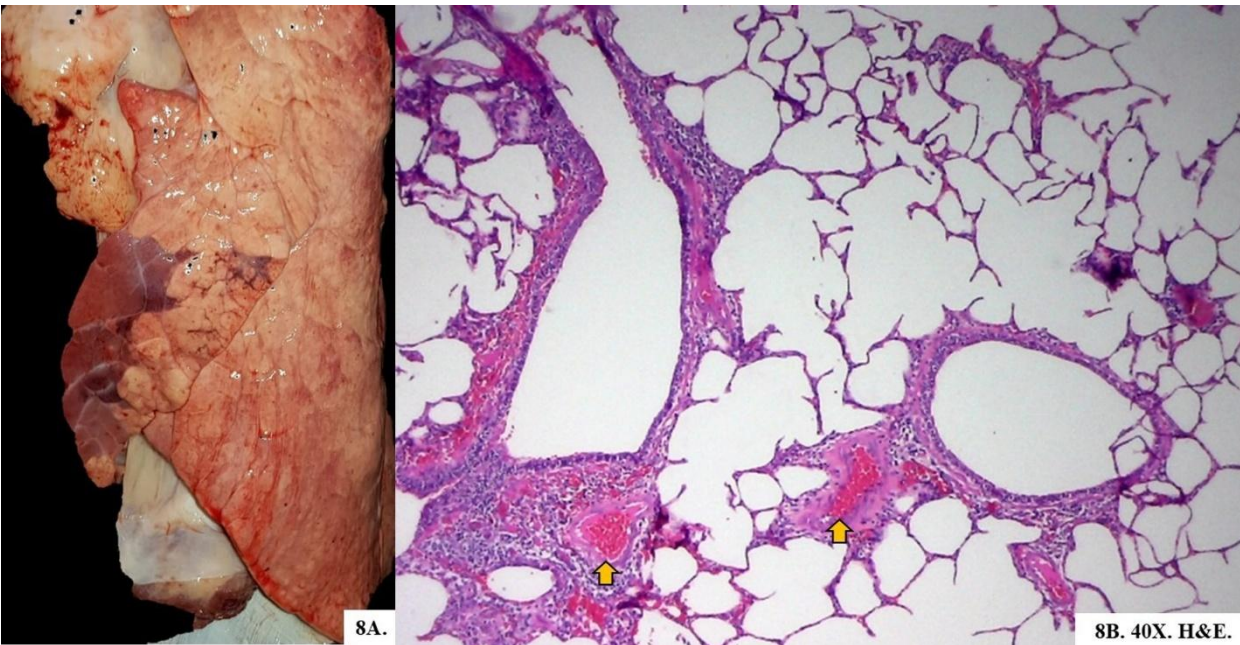
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515 **Figure 7.** Macroscopic and microscopic (histopathologic) images from a SPES2 sample collected
 516 during the study. 7A. Gross image of the lung showing lesions corresponding to a SPES2 score. 7B.
 517 Lung tissue section at 40× magnification (H&E stain). The arrow indicates moderate activation of
 518 bronchus-associated lymphoid tissue (BALT).

519



520

521 **Figure 8.** Macroscopic and microscopic (histopathologic) images from an MADEC2 sample
 522 collected during the study. 8A. Gross image of the lung showing a MADEC2 score. 8B. Lung tissue
 523 section at 40× magnification (H&E stain). The arrows indicate mild pulmonary congestion.