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### Pediatric RSV Diagnostic Testing Performance

#### Citation for published version:

Onwuchekwa, C, Atwell, J, Mora Moreo, L, Menon, S, Machado, B, Siapka, M, Agarwal, N, Rubbrecht, M, Aponte-Torres, Z, Rozenbaum, M, Curcio, D, Nair, H, V Kalina, W, Vroling, H, Gessner, B & Begier, E 2023, 'Pediatric RSV Diagnostic Testing Performance: A Systematic Review and Meta-analysis', *The Journal of* Infectious Diseases. https://doi.org/10.1093/infdis/jiad185

#### **Digital Object Identifier (DOI):**

10.1093/infdis/jiad185

#### Link: Link to publication record in Edinburgh Research Explorer

**Document Version:** Peer reviewed version

**Published In:** The Journal of Infectious Diseases

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1	Pediatric RSV Diagnostic Testing Performance: A Systematic Review and Meta-analysis
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12	
13	Running title: RSV diagnostic testing in children
14	Keywords: Respiratory Syncytial Virus Infections; epidemiology; diagnosis; sensitivity and
15	specificity, children
16	Abstract word count: 200
17	Main text word count: 3486
18	Article summary: We investigated whether adding other specimen types to
19	nasopharyngeal/nasal aspirate or swab RT-PCR increased RSV detection in children, as
20	found in adults. RT-PCR was the most sensitive diagnostic test and adding additional
21	specimen types modestly increased RSV detection.

#### 22 ABSTRACT

Background: Adding additional specimen types (e.g., serology or sputum) to nasopharyngeal
swab (NPS) RT-PCR increases respiratory syncytial virus (RSV) detection among adults. We
assessed if a similar increase occurs in children and quantified under-ascertainment associated
with diagnostic testing.

Methods: We searched databases for studies involving RSV detection in persons <18 years</li>
using ≥2 specimen types or tests. We assessed study quality using a validated checklist. We
pooled detection rates by specimen and diagnostic tests and quantified performance.

Results: We included 157 studies. Added testing of additional specimens to NP aspirate (NPA), NPS and/or nasal swab (NS) RT-PCR resulted in statistically non-significant increases in RSV detection. Adding paired serology testing increased RSV detection by 10%, NS by 8%, oropharyngeal swabs by 5%, and NPS by 1%. Compared to RT-PCR, direct fluorescence antibody tests, viral culture, and rapid antigen tests were 87%, 76%, and 74% sensitive, respectively (pooled specificities all ≥98%). Pooled sensitivity of multiplex versus singleplex RT-PCR was 96%.

37 Conclusions: RT-PCR was the most sensitive pediatric RSV diagnostic test. Adding multiple
38 specimens did not substantially increase RSV detection, but even small proportional increases
39 could result in meaningful changes in burden estimates. The synergistic effect of adding
40 multiple specimens should be evaluated.

41 Keywords: Respiratory Syncytial Virus Infections; epidemiology; diagnosis; sensitivity and
42 specificity, children

#### 44 Introduction

Respiratory syncytial virus (RSV) is the most common cause of acute lower respiratory tract infection (LTRI) in children [1]. Globally, an estimated 33 million episodes of RSV LRTI, 3.6 million hospitalizations, and 101,400 deaths occur yearly [2]. RSV is associated with 28% of all childhood acute lower respiratory infection (ALRI) episodes and 13-22% of ALRI mortality in young children [3]. Virtually all children experience RSV within the first two years of life; nearly 40% of RSV-associated hospitalizations in children <5 years of age occur within the first 6 months [2, 4].

52 No vaccines or RSV-specific treatments are currently available, and care is largely supportive. Because viral identification has limited impact on clinical decision-making, RSV diagnostic 53 testing is not universally performed, and RSV is, therefore, likely underdiagnosed. Diagnostic 54 55 testing has become more common in recent years and often involves upper respiratory tract specimens, including nasopharyngeal aspirates (NPA), swabs (NPS), nasal swabs (NS), washes 56 (NW), or oropharyngeal swabs (OPS). Testing methodologies include viral culture, rapid 57 antigen detection tests (RADT), immunofluorescence (IF) tests (including direct fluorescent 58 antibody (DFA)), and reverse transcriptase polymerase chain reaction (RT-PCR). Previously, 59 60 viral culture was considered the gold standard test for RSV, especially in children with high 61 viral shedding; however, it requires several days and special specimen handling, which limits 62 use in clinical settings [5, 6]. DFA tests and RADTs had replaced viral culture for routine clinical testing; more recently, molecular methods have become the gold standard due to their 63 higher sensitivity [6, 7]. Additionally, serum testing can identify recent infections by a  $\geq$ 4-fold 64 increase in RSV-specific antibody titers between paired acute and convalescent specimens or 65 pre- and post-season specimens. However, up to one-third of children do not seroconvert after 66 infection [6, 8]. The accuracy of each diagnostic test is influenced by specimen type, specimen 67 quality, age of the patient and RSV prevalence [6, 9, 10]. 68

69 Because disease incidence is a crucial component of decision-making for preventive interventions, it is critical to estimate RSV incidence accurately. In theory, some testing 70 methods and specimen sources may underestimate incidence in children, and even small 71 72 proportional underestimations could translate to high numbers of missed cases. Our recent review suggested that adding testing of an additional specimen type to nasal/NP swab RT-PCR 73 can increase adult RSV detection by 28-45% [11], depending on the specimen type. However, 74 75 similar data are needed to assess potential underestimation in other age groups. Therefore, we conducted a systematic review and meta-analysis to quantify RSV under-ascertainment due to 76 77 different specimens and testing modalities among children. We also summarize the evidence on the diagnostic accuracy of available testing modalities and evaluate factors affecting test 78 performance. 79

#### 80 Methods

81 Protocol and registration

This meta-analysis was part of a larger review on RSV under-ascertainment due to diagnostic approaches in adults and children. The protocol adhered to the Preferred Reporting Items for Systematic Review and Meta-Analyses for Diagnostic Test Accuracy Studies (PRISMA-DTA) and was registered on the PROSPERO database (registration No. CRD42022313209) [12].

86 Study eligibility criteria

Eligibility criteria were developed using the Population-Intervention-Comparator-Outcomes-Time Frame-Setting (PICOTS) framework (Supplementary Table 1). We identified primary studies reporting on RSV detection in children and young adults (age <18 years), using  $\geq 2$ types of specimens or diagnostic tests (index and reference) in the same population. Eligible studies reported the number of RSV cases detected by each testing method or specimen type or presented 2-by-2 tables of test performance.

#### 93 Information sources and search strategy

We searched Embase, MEDLINE (via PubMed), and Web of Science for publications from 01 94 January 2000 to 27 December 2021. We also searched non-indexed sources and RSV research 95 96 networks, such as Respiratory Syncytial virus Consortium in Europe (RESCEU project, https://resc-eu.org/) and Respiratory Syncytial Virus Network (ReSViNET, 97 https://www.resvinet.org/). The bibliographies of selected systematic reviews and meta-98 analyses were also screened for relevant articles (Supplementary Table 2). Detailed search 99 strategy and search terms are presented in Supplementary Table 3. 100

101 Study selection

Retrieved articles were screened based on the eligibility criteria (Supplementary Table 1).
Titles and abstracts were first assessed to identify potentially relevant articles. We subsequently
conducted independent duplicate full-text screening, with disagreements resolved by a third
reviewer. We used DistillerSR to manage the study selection process [13].

106 Data extraction

107 We created a structured extraction form on DistillerSR, previously used for a similar review in

adults [11]. All extracted data were independently cross-checked to minimize errors.

109 Definitions for data extraction

The extracted data items are summarized in Supplementary Table 4. We defined children as persons <18 years of age, with the following subgroups: infants (< 12 months), young children (1- < 5 years), and older children (5-18 years). Consistent with the PRISMA-DTA guidance, the index test was defined as the test under evaluation, while the reference test was defined as the comparator test [14]. A true positive (TP) result was positive by both the index and reference test, and a false positive (FP) result was positive by the index test but negative by the reference test. A false negative (FN) result was negative by the index test but positive by the reference test, and a true negative (TN) result was negative by both index and reference tests.

118 Risk of bias and applicability

The risk of bias (RoB) assessment of included studies was performed by one reviewer and verified by a second reviewer using the Quality Assessment of Diagnostic Accuracy Studies (QUADAS-2) tool (Supplementary Table 5) [15]. Disagreements were resolved by consensus, and when required, a third reviewer resolved the conflict. We developed an extension to the main QUADAS-2 tool to facilitate bias assessment when comparing specimens using a single test method.

125 Diagnostic accuracy measures and synthesis of results

The analytic approach for this review and meta-analysis is detailed elsewhere [11]. Briefly, we describe the included studies and summarize key characteristics of the study population and testing approach (including specimens used and diagnostic method).

We compared the RSV detection rate (DR) under combined testing (using multiple specimens or diagnostic tests) and test-specific (reference) conditions, presenting this as detection rate ratios (DRR). For this comparison, RT-PCR using NPA, NPS, or NS was used as the reference testing method in order of sample preference, and all positives were considered true positives.

The data from 2-by-2 tables reporting TP, FP, FN, and TN were used to calculate individual studies' test sensitivity and specificity (with 95% confidence intervals). The sensitivity and specificity point estimates were pooled in meta-analyses for RADTs, DFA and viral culture, using RT-PCR as a common reference. All meta-analyses were performed under random-effects assumptions [5, 16] and were restricted to analyses with ≥3 studies. Each index-reference comparison was considered a separate experiment, presented separately, and

139 contributed to the meta-analysis estimates. All analyses were performed in R version 4.2.0,140 using the meta4diag and metafor packages [17].

141 Subgroup and sensitivity analyses

In the meta-analysis of test performance, we stratified findings by the type of RT-PCR used as the reference test, either singleplex or multiplex. We also stratified test performance by the following pediatric age categories for which estimates were available: infants, all under 5 years (without further disaggregation), and all under 18 years (without further disaggregation). Estimates limited to older children were not available.

We also assessed the impact on the pooled results of study quality, based on the QUADAS-2
assessment, by conducting separate meta-analyses, including only studies with a low risk of
bias.

#### 150 **Results**

151 Study selection process

Our search identified 8066 unique references, of which 157 studies were included in the review (Figure 1 and Supplementary Table 6). The most common reasons for exclusion at full-text level were studies conducted in adults or not reporting estimates for those aged <18 years separately, studies conducted before 2000, and those not reporting on RSV detection by multiple methods.

157 Study characteristics

Supplementary Table 7 provides detailed descriptions of included studies. Over 60% of studies
were conducted in the Americas and Europe, and 93% used a cross-sectional design (Table 1).
Most studies included hospitalized children (67%), and the most common clinical presentation
was LRTI, including bronchiolitis, pneumonia and non-specified acute respiratory infections
(67.5%).

NPAs were collected in 85 studies (54.1%), while 53 (33.8%) used NPS. Multiplex and singleplex RT-PCR were evaluated in 86 (54.8%) and 50 (31.8%) studies, respectively, while RADTs were evaluated in 56 studies (35.7%). Among included studies, 55 (35.0%) reported on children 0-5 years of age, 62 (39.5%) included children 0-18 years, and 6 (3.8%) were conducted on infants 0-1 year. Thirty-four studies (21.7%) did not clearly specify the age range of children included in the study.

169 Risk of bias and applicability

Supplementary Table 8 provides a detailed assessment of the risk of bias and applicability at 170 domain level. Overall, 20 studies (12.3%) had a low risk of bias assessment in all domains, 171 while 37 (22.7%) and 99 studies (60.7%) had a high or unclear risk of bias in at least one 172 domain, respectively. Figure 2 provides a graphical summary of the risk of bias assessment at 173 174 domain level for all eligible studies. In the patient selection domain, nearly half of the studies had an unclear risk of bias assessment, and 18 studies had a high risk of bias. Common reasons 175 for unclear or high risk of bias in this domain were unclear patient population and study designs 176 that could be interpreted as case-control designs. Risk of bias concerns for the index and 177 reference tests were often due to failure to report whether test operators were blinded or if 178 179 previously characterized stored specimens were used. Risk of bias in the index test domain was 180 not performed in 10% of studies where sample types were compared using the same test 181 (considered reference, e.g., RT-PCR of two different specimen types). Risk of bias in the domain of flow and timing was assessed as unclear in 50 studies (32%), often because it was 182 not explicitly reported that both the index and reference tests were conducted in the same period 183 of illness and whether all included patients had the same reference test (e.g., studies applying 184 185 a composite reference). Less than 10% of studies had high/unclear applicability concerns in any of the 3 relevant domains. 186

#### 187 Results of individual studies

Supplementary Table 9 provides the number of RSV infections detected by each diagnostic test and specimen type combination obtained in each study. It also provides overall RSV detection by combination testing and test-specific detection rates. Supplementary Table 10 displays the diagnostic performance, in terms of sensitivity and specificity, of included studies stratified by test type.

193 Synthesis of results

194 Increase detection of RSV by combined specimen testing

Testing of multiple respiratory specimens by any method (i.e., RADT, DFA, RT-PCR or paired 195 196 serology) resulted in a small overall increase in RSV detection (all non-statistically significant trends) for the specimen types examined (Table 2 and Supplementary Figure 1). Using RT-197 PCR on NPA/NPS or paired specimens, additional testing of OPS to NPS/NPA increased RSV 198 199 detection by 5% (DRR 1.05, 95% CI 0.96-1.14). Adding testing of NS to NPA/NPS increased detection by 8% (DRR 1.08, 95% CI 0.94-1.25), while adding NPS to NPA increased RSV 200 201 detection by 1% (DRR 1.01, 95% CI 0.94-1.08) (Figure 3). Excluding 4 studies with high or 202 unclear risk of bias did not change these results substantially (Supplementary Figure 2) [18-21]. Adding paired serology testing to RT-PCR on NPS resulted in a 10% increase in RSV 203 detection in one study (DRR 1.10, 95% CI 0.95-1.27) [22]. Stratified data allowed calculation 204 of detection ratios by age group for added paired serology: 0 - 6 months: DRR 1.02 (95% CI 205 0.76-1.36), 7 months-2 years: DRR 1.12 (95% CI 0.94-1.33), 3-5 years: DRR 1.04 (95% CI 206 0.73 – 1.48), 6-17 years: DRR 1.29 (95% CI 0.65 – 2.54). Adding sputum RT-PCR increased 207 RSV detection by 11.7% compared to OPS RT-PCR alone in one study (124/440 versus 208 111/440) [23]. Adding RT-PCR of BAL increased detection by 26.7% compared to OPS-PCR 209 alone in one study (19/76 versus 15/76) [24]. 210

211 Performance of RADT versus RT-PCR

We included 21 studies performing 28 comparisons of RADT as index tests compared to RT-212 PCR with a pooled sensitivity of 75% (95% CI 69-80%) with a range of 26-100% (Figure 4). 213 214 Stratified by population age, the pooled sensitivity was 63% in studies limited to children 0-5 years (95% CI 49-76%) and 78% in studies including children up to 18 years (95% CI 74-215 82%). When stratified by the type of RT-PCR test, the pooled sensitivity of RADT against 216 singleplex RT-PCR was 70% (95% CI 59-79%) and 76% against multiplex RT-PCR (95% CI 217 71-81%). Overall, the specificity of RADT was 97% (95% CI 95-98%) with little change when 218 219 stratified by age groups or RT-PCR type.

220 Performance of DFA versus RT-PCR

The pooled sensitivity of DFA against RT-PCR was 87% (95% CI 78-93%) and ranged between 72-96%, with similar results obtained with singleplex and multiplex RT-PCR references (Figure 5). Stratified by age groups, pooled sensitivity was 82% in studies limited to children 0-5 years (95% CI 76-87%) and 88% in studies reporting on all children 0-18 years (95% CI 73-95%). The pooled specificity of DFA was 99% (95% CI 97-100%), with similar results obtained when stratified by age groups and RT-PCR types.

227 Performance of viral culture versus RT-PCR

Viral culture had a pooled sensitivity of 76% compared to RT-PCR (95% CI 71-81%) with a

- range of 75-100% and a pooled specificity of 100% (95% CI 99-100%). It was not possible to
- assess the performance of viral culture by age or type of RT-PCR reference owing to the small
- number of studies included in the analysis (Figure 5).

232 Performance of multiplex versus singleplex RT-PCR and agreement between multiple233 platforms

Most of the commonly used multiplex platforms performed well against singleplex RT-PCR. The pooled sensitivity of multiplex RT-PCR was 96% (95% CI 92-98%), while pooled specificity was 100% (95% CI 98-100%) (Supplementary Figure 3). The performance of multiplex against singleplex RT-PCR showed little variation by subject age range.

#### 238 Discussion

To our knowledge, this is the first systematic review and meta-analysis to report on both 239 specimen and diagnostic test performance in children compared to common reference methods. 240 Overall, we found that additional testing of other specimen types to NPS/NPA RT-PCR did not 241 substantially increase RSV detection in pairwise comparisons (1-10% average increase 242 depending on the additional specimen type). However, given the higher disease burden among 243 children, this could meaningfully increase incidence estimates. Variation within individual 244 studies was also modest, with a maximum reported increase in detection of 24% associated 245 with additional RT-PCR testing of nasal swabs. While RT-PCR was the most sensitive 246 diagnostic method, DFA was comparatively more sensitive than RADT and viral culture in 247 detecting RSV infections. All 3 diagnostic tests had specificities above 97%. Similar to prior 248 findings in adults [11], using multiplex PCR rather than singleplex resulted in a small 249 250 decrement in sensitivity (4%).

Our findings suggest that in children presenting with LRTI, routine testing by RT-PCR using a single NPA or NPS specimen is unlikely to miss RSV cases or only a relatively small proportion. And, while we found that additional testing of BAL or sputum may increase detection more substantially, the small number of studies we found means we cannot draw conclusions on the value of adding lower respiratory specimens. This contrasts with results obtained from our review in adults, where adding an additional specimen to NPS increased

257 RSV detection by 28–45%, depending on specimen type [11]. This difference between children and adults may be due to several reasons. Firstly, our analysis of RSV detection in adults used 258 259 NPS as a reference specimen type because this was most commonly used in adults and is less 260 sensitive than NPA for RSV detection. Secondly, most studies in adults collected additional lower respiratory tract specimens, such as sputum, which may produce higher viral copies for 261 detection [19, 25, 26]. Thirdly, on average, children exhibit higher levels and more extended 262 263 periods of RSV shedding than adults [27-30], have a smaller upper airway space, and may present earlier during illness. These factors may increase the likelihood of finding a positive 264 265 result from a single upper airway specimen.

We identified one study that investigated the added RSV detection using paired serology, resulting in a 10% increase in RSV detection in children with community-acquired pneumonia [22]. As an indicator of recent infection, a 4-fold increase in acute-convalescence serology titers has been shown to increase RSV detection in adults [31]. However, the interpretation of serological titers may be complicated by a lower likelihood of seroconversion in children [8, 32]. This may be partly attributed to IgG maternal antibodies in infants, producing high baseline antibody levels and effectively masking the immune response during infection [33].

273 Compared to specimen type and number, we observed a greater variation by diagnostic method. Compared to RT-PCR, DFA was most sensitive (87%), followed by viral culture (76%) and 274 RADT (75%), with the latter value consistent with other pediatric studies and higher than 275 reported in adults [5]. Our results document a higher sensitivity of these testing modalities in 276 children compared to our results in adults [11], which may be linked to higher viral shedding 277 in children, as prior infection in adults results in a greater immune response and lower shedding. 278 Also, NPA is used more frequently for RSV diagnosis in children, which may increase the 279 sensitivity of RADTs, owing to the higher viral yield of NPA compared to NPS [19]. 280 Regardless, adjusting pediatric RSV incidence estimates upwards by 13-26% may be 281

reasonable depending on the testing method used when studies use RADT, DFA or viral cultureinstead of RT-PCR.

284 Strengths and limitations

One of the key strengths of our review was the added search of relevant grey literature sources and RSV networks (Supplementary table 2). This addition resulted in a large number of relevant studies representing global data. We also included a range of clinical settings, clinical presentations, and multiple specimen and testing combinations, giving the results of this review wider utility.

A limitation of our review is that 60% of studies had an overall high/unclear risk of bias. Most 290 of the included studies also did not include sufficient demographic data or information about 291 292 specimen storage and handling; hence the effect of these factors could not be evaluated within 293 this meta-analysis. Appropriate reporting of DTA studies following the STARD (Standards for Reporting of Diagnostic Accuracy) guidance will mitigate this limitation [34]. Furthermore, 294 295 prominent reasons for the attribution of a high risk of bias included the use of a case control design, which may inadvertently increase participants who are the "sickest of the sick" or the 296 "wellest of the well" [35], thereby inflating diagnostic accuracy [36]. In addition, a bias could 297 have been introduced when a non-automated index test was interpreted with prior knowledge 298 of the reference test [36]. 299

300 Our comparison of diagnostic test performance used RT-PCR as a common reference. A 301 limitation of this approach is that the true accuracy of RT-PCR remains a relatively unknown 302 quantity, and implicit assumptions of its accuracy are made. However, our analysis shows that 303 RT-PCR consistently detects >95% of pediatric RSV infections identified by all testing 304 modalities (Supplementary Table 9), with DFA, RADT and viral culture detecting the other

5% missed by RT-PCR. These findings support our approach of using RT-PCR as a "gold
standard" reference for test accuracy evaluations.

307 Our review also did not stratify the analysis by retrospective and prospective studies because 308 few studies were adequately similar to analyze test performance. We also did not capture the 309 duration of specimen storage, as this may also influence diagnostic test performance.

#### 310 Conclusions

Compared to adults, where we observed a substantial statistically significant increase in RSV 311 detection, the magnitude of the added benefit in a pediatric population was more modest, 312 especially regarding additional specimen types. Given the modest levels of increased 313 detection in published pairwise comparisons, assessing the combined effects of adding multiple 314 315 specimen types would likely be the most efficient way to study this issue further. In adults, 316 each added specimen type identifies some unique additional cases creating a synergistic effect in the cumulative increase in RSV detection [37]. Further, including saliva as an additional 317 specimen type in such studies would be warranted. While we did not find RSV studies 318 examining saliva, it has recently been shown to be a high-yield specimen type in adults [37] 319 and for SARS-CoV-2 detection [38]. Such studies could confirm the appropriate level of 320 321 adjustment for epidemiologic data when a single NPA or NPS specimen type is used in disease incidence studies. 322

Under-ascertainment was documented for studies that used diagnostic methods other than RT-PCR. This finding might be incorporated into RSV incidence estimates, future estimates of vaccine effectiveness, and population-level impact once RSV vaccines become available. RT-PCR remains the most sensitive test for RSV, having the advantages of rapid turn-around time and lower running costs. Although RADTs may offer an acceptable alternative in some settings. Further studies are needed to understand the combined effect of specimen and diagnostic test choice, specimen handling and other external factors on RSV detection. Even
small proportional increases in RSV detection, when applied to large numbers of RSV
infections, could result in meaningful changes in burden estimates.

332

#### 333 Footnotes

#### 334 Conflict of Interest,

P95 was contracted by Pfizer to conduct this study. Neha Agarwal, Zuleika Aponte-Torres, 335 336 Belen Machado, Sonia Menon, Laura Mora Moreo, Chukwuemeka Onwuchekwa, Michelle Rubbrecht, Mariana Siapka, Hilde Vroling are employees of P95, which received funding from 337 Pfizer in connection with the development of this manuscript. Harish Nair received grants from 338 Innovative Medicine Initiative, National Institute of Health and Care Research, Bill and 339 Melinda Gates Foundation, WHO and Pfizer, and consultancy or honoraria and speaker fees 340 from Sanofi, Merck, Novavax, ReViral and GSK (all paid to institution). Jessica E. Atwell, 341 Elizabeth Beiger, Daniel Curcio, Bradford D. Gessner, Warren Kalina, and Mark Rozenbaum 342 are employees of Pfizer and may hold stock or stock options. 343

#### 344 Funding

345 This work was supported by Pfizer Inc., USA.

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470 471 472 473 474 475 476 477 478	Figure LegendsFigure 1. PRISMA flow chart showing the study selection processFigure 2. Summary risk of bias assessment of included studies based on the QUADAS-2 toolFigure 3. Increase in RSV detection rate due to the addition of another specimen testing to reference RT-PCR of Nasopharyngeal aspirate/swab or Nasal swab(Abbreviations: ARI= acute respiratory infection, URTI= upper respiratory tract infection, LRTI= lower respiratory tract infection, NR= not reported, CI= confidence interval)Figure 4a. Sensitivity of Rapid antigen detection tests (RADT) test using reverse transcription polymerase chain reaction (RT-PCR) as reference test

- 480 Figure 4b. Specificity of Rapid antigen detection tests (RADT) test using reverse
- 481 transcription polymerase chain reaction (RT-PCR) as reference test
- 482 (Abbreviations: TN= true negative, FP= false positive, TN + FP= all reference negative)
- 483 Figure 5a. Sensitivity of Direct fluorescent antibody test (DFA) using reverse transcription
- 484 polymerase chain reaction (RT-PCR) as reference test
- 485 (Abbreviations: TP= true positive, FN= false negative, TP + FN= all reference positives)
- 486 Figure 5b. Specificity of Direct fluorescent antibody test (DFA) using reverse transcription
- 487 polymerase chain reaction (RT-PCR) as reference test
- 488 (Abbreviations: TN= true negative, FP= false positive, TN + FP= all reference negative)
- 489 Figure 5c. Sensitivity of Viral culture using reverse transcription polymerase chain reaction
  490 (RT-PCR) as reference test
- 491 (Abbreviations: TP= true positive, FN= false negative, TP + FN= all reference positives)
- 492 Figure 5d. Specificity of Viral culture using reverse transcription polymerase chain reaction
  493 (RT-PCR) as reference test
- 494 (Abbreviations: TN= true negative, FP= false positive, TN + FP= all reference negative)