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# The Level of Procalcitonin in Severe COVID-19 Patients: A Systematic Review and Meta-Analysis

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Alireza Shahriary, Thomas P. Johnston,  
and Amirhossein Sahebkar

## Abstract

There is data from individual clinical trials suggesting that procalcitonin (PCT) may be a prognostic factor in the severity of COVID-19 disease. Therefore, this systematic review and meta-analysis was performed to investigate PCT levels in severe COVID-19 patients. We searched Embase, ProQuest, MEDLINE/

PubMed, Scopus, and ISI/Web of Science for studies that reported the level of PCT of patient with severe COVID-19. We included all studies regardless of design that reported the level of PCT in patients with severe COVID-19. We excluded articles not regarding COVID-19 or not reporting PCT level, studies not in severe patients, review articles,

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25 editorials or letters, expert opinions, com- 67  
 26 ments, and animal studies. Nine studies were 68  
 27 included in the analysis. The odds of having 69  
 28 more severe COVID-19 disease was higher in 70  
 29 subjects with elevated PCT ( $\geq 0.05$  ng/mL) 71  
 30 compared with those having low procalcitonin 72  
 31 ( $< 0.05$  ng/mL) [ $n = 6$ , OR(95% 73  
 32 CI) = 2.91(1.14, 7.42),  $p = 0.025$ ). After esti- 74  
 33 mating the mean and standard deviation val- 75  
 34 ues from the sample size, median, and 76  
 35 interquartile range, a pooled effect analysis 77  
 36 indicated higher serum PCT concentrations in 78  
 37 patients with severe versus less severe disease 79  
 38 [ $n = 6$ , SMD(95% CI) = 0.64(0.02, 1.26), 80  
 39  $p = 0.042$ ]. The results of this study showed 81  
 40 that PCT is increased in patients with severe 82  
 41 COVID-19 infection. 83

42 **Keywords**

43 Coronavirus · Procalcitonin · Meta-analysis · 86  
 44 SARS-CoV-2 · COVID-19 · Viral infection 87

and resource issues, and in the early stage of the 67  
 disease, the positive rate of the test was relatively 68  
 low [7]. Therefore, there is an urgent need to 69  
 identify reliable biomarkers that may help predict 70  
 the risk for illness severity so that appropriate 71  
 care can be allocated earlier [8]. 72

One of the most studied biomarkers in this 73  
 field is procalcitonin (PCT), which has both diag- 74  
 nostic and prognostic utility [9]. PCT has become 75  
 a promising biomarker for early detection of bac- 76  
 terial infections in modern clinical practice [10]. 77  
 However, the expression of this biomarker may 78  
 differ in severe COVID-19 infection with some 79  
 studies reporting that PCT is increased in patients 80  
 with severe COVID-19 [11, 12] and others report- 81  
 ing that PCT in these patients is normal [13, 14]. 82  
 These differences could reflect the fact that the 83  
 clinical characteristics differ in those patients 84  
 with mild versus severe COVID-19 infections [9]. 85

We have aimed to resolve this issue by carry- 86  
 ing out a systematic review of PCT levels in 87  
 COVID-19 cases. 88

89 **25.2 Methods**

90 **25.1 Introduction**

91 Patients with unexplained pneumonia were initially 92  
 93 reported in Wuhan, China, in December 94  
 95 2019. A novel coronavirus named severe acute 96  
 97 respiratory syndrome coronavirus 2 (SARS- 98  
 99 CoV-2) was detected in samples of the lower 100  
 respiratory tract of the infected patients, and the 101  
 disease was termed COVID-19 (coronavirus- 102  
 2019) [1]. This disease is rapidly spreading 103  
 across the world [2] and has turned into a pan- 104  
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This study was performed according to the 90  
 Preferred Reporting Items for Systematic 91  
 Reviews and Meta-Analyses (PRISMA) state- 92  
 ment [15]. We aimed to compare the levels of 93  
 PCT in patients with severe versus non-severe 94  
 COVID-19 infections, to determine if it can be 95  
 used as a biomarker to predict disease course. A 96  
 threshold level of 0.05 ng/mL PCT was taken to 97  
 discriminate between severe and non-severe 98  
 infection for each study. 99

100 **25.2.1 Search Strategy**

We searched, Embase, ProQuest, MEDLINE/ 101  
 PubMed, Scopus, and ISI/Web of Science for 102  
 studies that reported the level of PCT in patients 103  
 with COVID-19 infections. A date limit was set 104  
 from December 2019, and the search was per- 105  
 formed up to April 5, 2020. The reference list of 106  
 articles was reviewed using forward and back- 107

Currently the gold standard for the diagnosis 63  
 of COVID-19 is reverse transcription polymerase 64  
 chain reaction (RT-PCR) testing, but, in many 65  
 countries, testing has been limited due to time 66

ward citation tracking to identify other eligible documents. No language limits were applied. type, and levels of PCT. Disagreements on the extracted data were resolved by consensus.

### 25.2.2 Selection Criteria

We included all studies regardless of study design, targeting prospective or retrospective studies that met the following criteria:

1. Reported the level of PCT in COVID-19 patients displaying serious symptoms
  2. Reported the level of PCT in COVID-19 patients displaying non-serious symptoms
- We excluded:

1. Articles not regarding COVID-19 or not reporting PCT level
2. Studies not including both severe and non-severe COVID-19 patients

3. Review articles, editorials or letters, expert opinions, comments, and animal studies

At least two reviewers independently evaluated titles and abstracts and selected relevant studies for inclusion. If this could not be done reliably using the title and abstract of an article, the full text version was retrieved for detailed analysis. Any disagreement was resolved by a third independent reviewer. The reasons for exclusion of studies were recorded.

At first we searched databases that included Scopus (7), Embase (9), ProQuest (14), PubMed (2), and Web of Science (2) and identified 31 total studies. In the next step we removed the duplicate articles and retained 19 records to include Scopus (7), Embase (2), ProQuest (10), PubMed (1), and Web of Science (2). Subsequently, the records were screened by title and abstract and six further studies were excluded. The remaining 16 records were screened further using the full text, and this left nine articles that met all requirements for inclusion in the study.

### 25.2.3 Data Extraction

Data were extracted independently by at least two reviewers and included authors, year of publication, clinical setting, sample size, sample

### 25.2.4 Quality Assessment

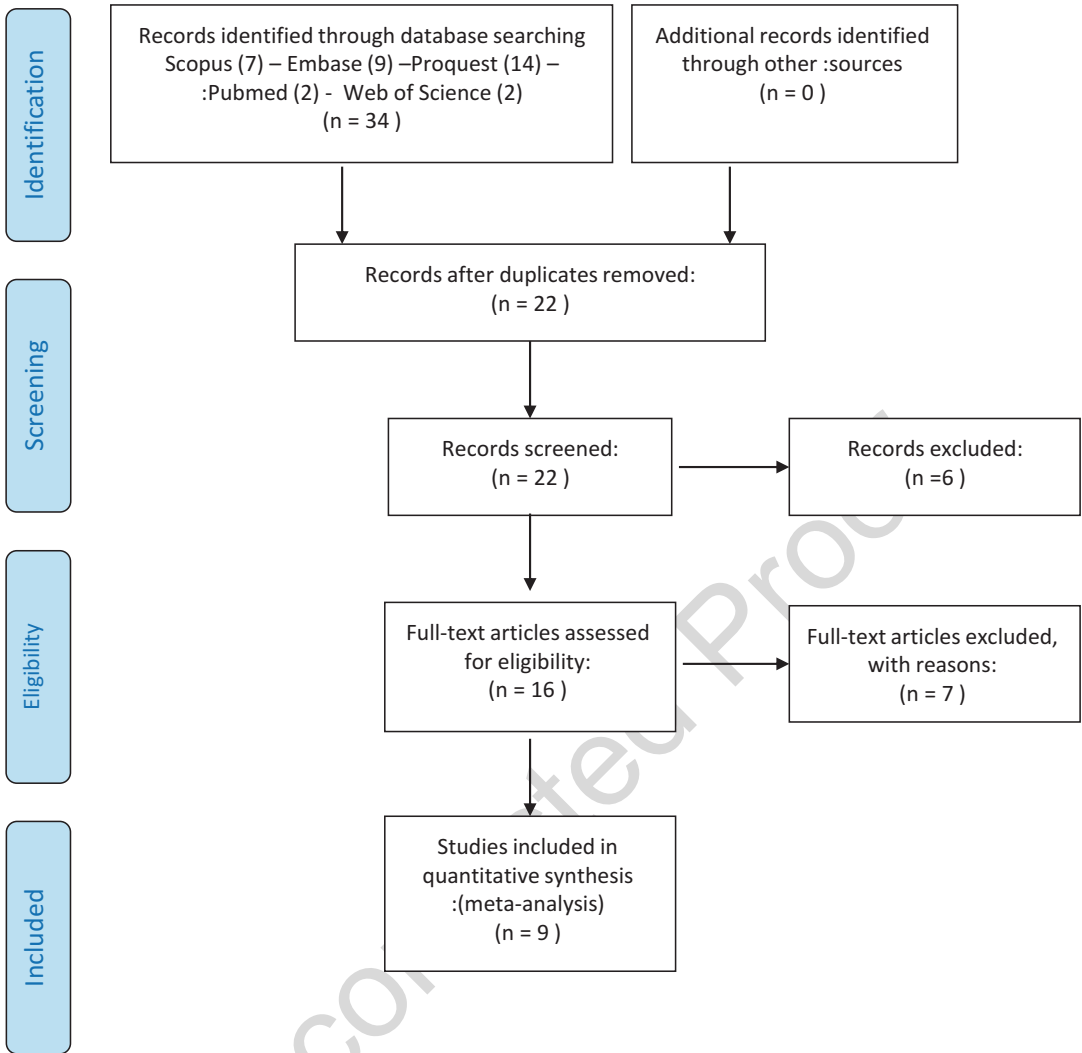
The methodological quality assessment of studies was performed independently by two authors using Quality Assessment of Diagnostic Accuracy Studies (QUADAS-2) criteria, as recommended by the Cochrane Collaboration [16]. Disagreements about inclusion criteria, data extraction, and quality assessment were resolved by consensus.

### 25.2.5 Data Synthesis and Statistical Analysis

We applied random-effects meta-analyses with inverse variance weighting to calculate pooled estimates and 95% confidence intervals (CIs). We estimated the sample mean and standard deviation (SD) from the sample size, median, and interquartile range via the Wan et al. approach [17] to compute a meta-analysis of outcomes and demonstrate effectiveness. In addition, we used odds ratios (ORs) and 95% CIs to compare the rates of disease severity. Heterogeneity ( $I^2$  statistics) was assessed and reported using Cochran's Q-test [18]. We also plotted Galbraith graphs to display heterogeneity. Egger's test and a visual inspection of funnel plots were carried out to evaluate publication bias between studies [19]. In general, a PCT concentration  $\geq 0.05$  ng/mL was considered as being high in our analysis. Values below 0.05 ng/mL were considered normal or low. Two groups of patients with either low or high disease were also examined. STATA version 14.0 (Stata Corp, College Station, TX) and R version 3.6.3 were used to conduct the analyses.

## 25.3 Results

A summary of study selection process and characteristics of the included studies are summarized in Fig. 25.1 and Table 25.1, respectively.



**Fig. 25.1** PRISMA flow diagram summarizing the study selection process

### 25.3.1 Results of PCT Analyses Using Odds Ratios

Results were obtained for PCT with an odds ratio in 6 out of the 9 studies which met the selection criteria. A randomized effect analysis showed significant risk according using a forest plot to visualize the results (Fig. 25.2). The odds of having more severe COVID-19 disease was higher in subjects with elevated PCT compared to those

with low PCT levels ( $n = 6$ ,  $OR(95\% \text{ CIs}) = 2.91 (1.14, 7.42), p = 0.025$ ) (Fig. 25.3).

The Q-test showed a significant variations in odds ratios attributable to heterogeneity of the procalcitonin data ( $I^2 = 73.5\%, p = 0.002$ ), which can also be seen in the Galbraith plot (Fig. 25.3). There was no publication bias in the PCT studies according to results of Egger’s test analyses and by using a funnel plot visualization ( $p = 0.434$ ) (Fig. 25.4).

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**Table 25.1** Final studies included in meta-analysis

No	Authors	Year	Procalcitonin		Sample size	Setting	Samples	Study design	
			Patients with worse condition	All Patients					
t1.1	1	Qiu et al. [20]	2020	N = 19 0.32 (0.19)	N = 36 0.24 (0.17)	36	Three hospitals in Zhejiang province, China	Children with coronavirus disease 2019	Retrospective
t1.2	2	Zhang et al. [12]	2020	N = 56 0.1 (0.06-0.3)	N = 138 0.07 (0.04-0.1)	138	No. 7 hospital of Wuhan	Patients diagnosed as COVID-19	Prospective
t1.3	3	Wang et al. [21]	2020	N = 36 27 (75.0)	N = 138 49 (35.5)	138	Zhongnan Hospital of Wuhan University	Patients with confirmed NCIP	Retrospective
t1.4	4	Guan et al. [22]	2020	N = 173 16/117 (13.7)	N = 1099 35/633 (5.5)	1099	552 hospitals in 30 provinces, autonomous regions, and municipalities in mainland China	Patients with laboratory-confirmed COVID-19	Retrospective
t1.5	5	Huang et al. [13]	2020	N = 13 0.1 (0.1-0.4)	N = 41 0.1 (0.1-0.1)	41	Wuhan, China	Patients with laboratory-confirmed 2019 nCoV infection	Prospectively
t1.6	6	Liu et al. [23]	2020	N = 69 2/69 (2.90)	N = 80 2/80 (2.50)	80	Department of Infectious Diseases, Union Hospital, Tongji Medical College, Huazhong University of Science and Technology	Patients with severe type COVID-19	Retrospective
t1.7	7	Shi et al. [16]	2020	N = 82 0.06(0.03-0.10)	N = 416 0.07(0.04-0.15)	416	Renmin Hospital of Wuhan University, located in Wuhan, Hubei province, China, was assigned responsibility for the treatment of patients with severe COVID-19 by the Wuhan government	Severe patients admitted to Renmin Hospital of Wuhan University with laboratory-confirmed COVID-19	Retrospective
t1.8	8	Peng et al. [24]	2020	N = 16 0.20(0.15-0.48)	N = 96 0.11(0.06-0.20)	112	Wuhan Union Hospital West Hospital	Patients with cardiovascular diseases infected with new coronavirus pneumonia	Retrospective
t1.9	9	Zhou et al. [16]	2020	N = 11 0.21(0.11-0.42)	N = 9 0.05(0.03-0.06)	20	West District of Union Hospital of Tongji Medical College	In-hospital severe patients with COVID-19	Retrospectively

t1.1

t1.2

t1.3

t1.4

t1.5

t1.6

t1.7

t1.8

t1.9

t1.10

t1.11

t1.12

t1.13

t1.14

t1.15

t1.16

t1.17

t1.18

t1.19

t1.20

t1.21

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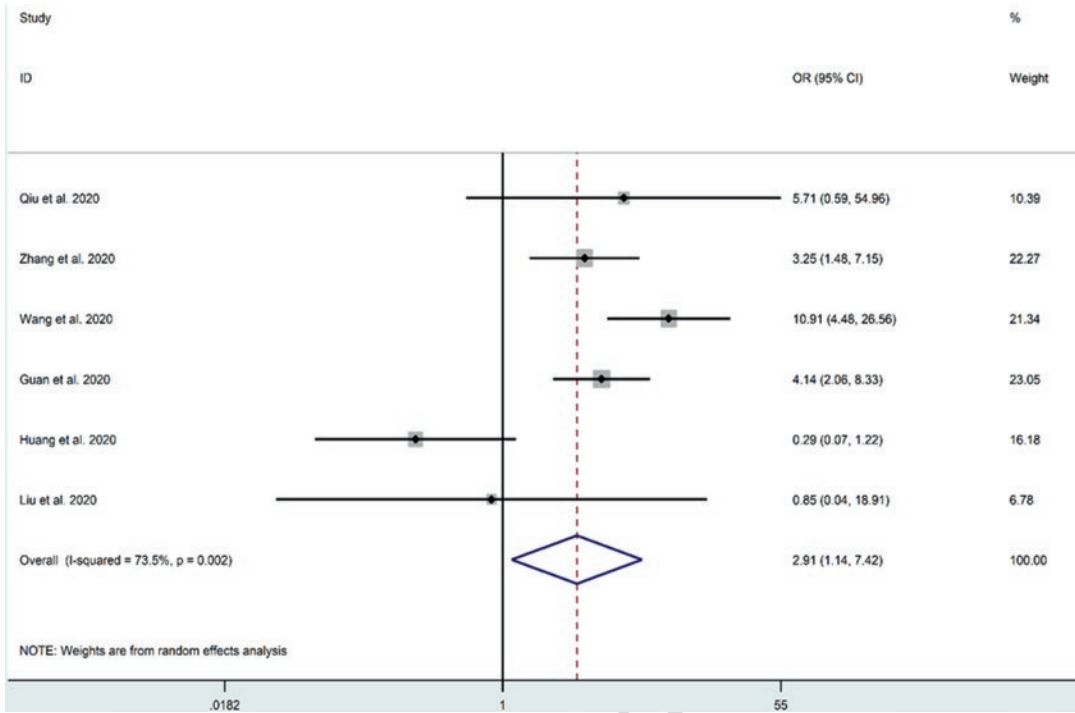
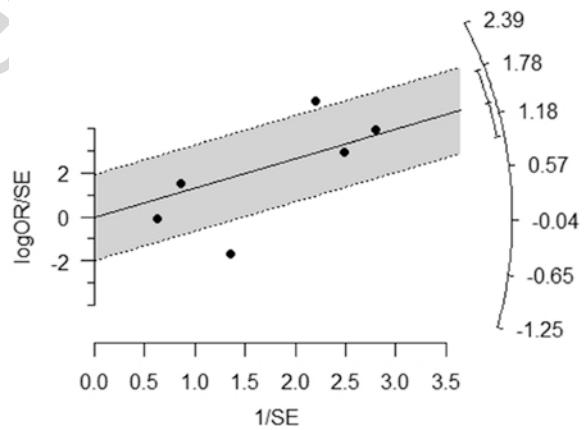


Fig. 25.2 Forest plot results for odds ratio of procalcitonin laboratory data

Fig. 25.3 Galbraith (radial) plot for odds ratio of trials used by procalcitonin laboratory data

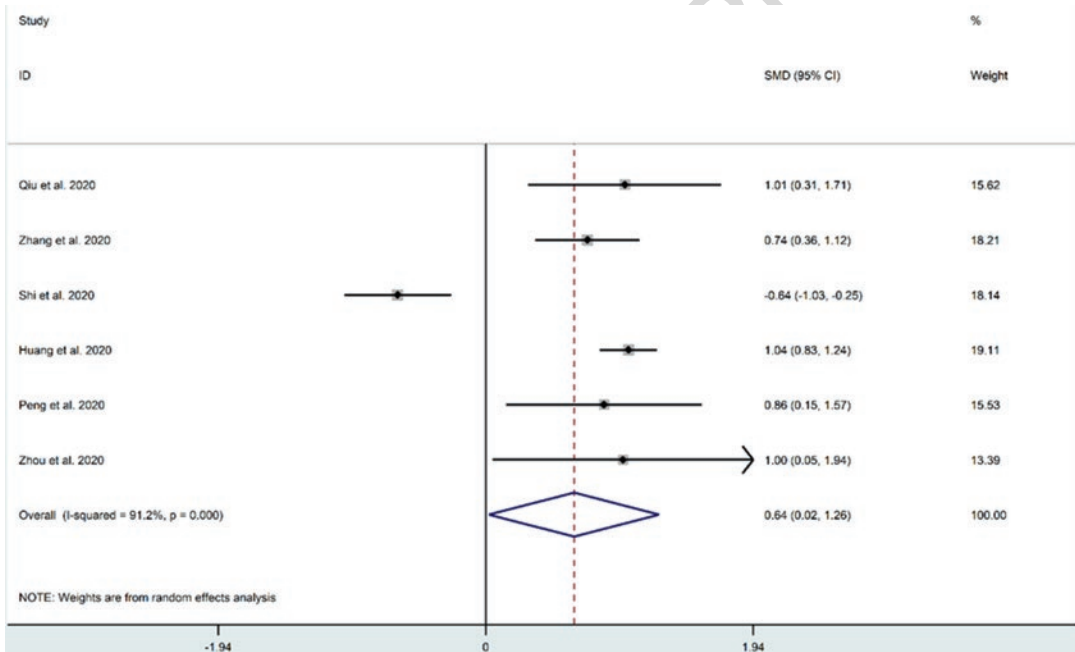
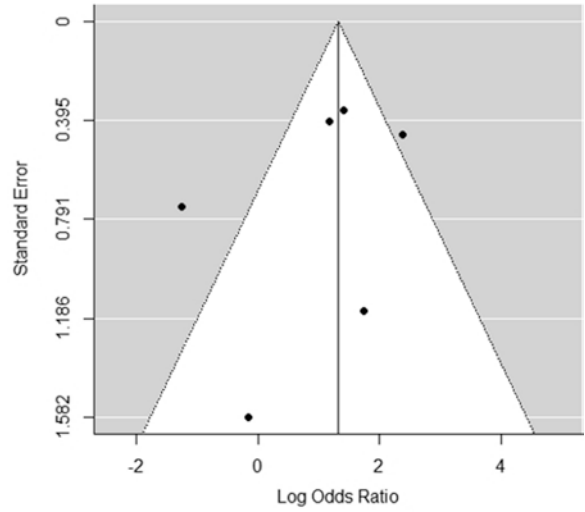


207 **25.3.2 Results of PCT Analyses Using**  
208 **Standardized Mean Difference**

209 All PCT results included in the six studies were  
210 analyzed using the standardized mean difference  
211 (SMD). For most studies (except the study by  
212 Qiu et al. [20]), the median and interquartile  
213 range of PCT were reported instead of the mean

and standard deviation. After estimating the val- 214  
ues of the mean and standard deviation from the 215  
sample size, median, and interquartile range, a 216  
pooling effect analysis indicated a significantly 217  
higher PCT level ( $n = 6$ ,  $SMD(95\%$  218  
 $CI) = 0.64(0.02, 1.26)$ ,  $p = 0.042$ ) in subjects with 219  
severe versus less severe COVID-19 disease 220  
(Fig. 25.5). 221

**Fig. 25.4** Funnel plot for verification of publication bias in the meta-analysis of odds ratio of procalcitonin laboratory data



**Fig. 25.5** Forest plot results for standardized mean difference of procalcitonin laboratory data

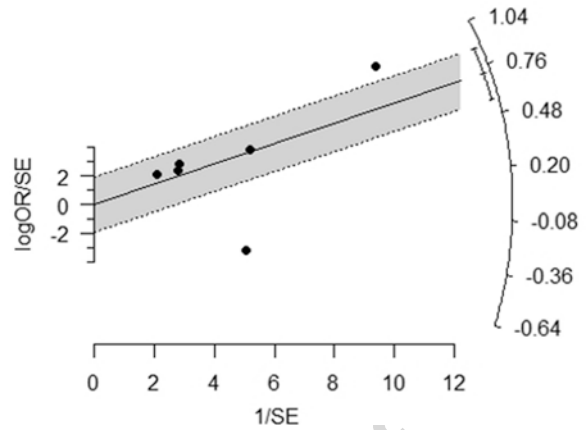
222 Next, we applied a random effects analysis  
 223 since there was heterogeneity between the studies  
 224 for PCT based on the Q-test and radial plot  
 225 results ( $I^2 = 91.2\%$ ,  $p < 0.001$ ) (Fig. 25.6).

226 Additionally, according to Egger’s test and visu-  
 227 alization with a funnel plot, there was no publica-  
 228 tion bias in the results of the meta-analysis  
 229 ( $p = 0.711$ ) (Fig. 25.7).

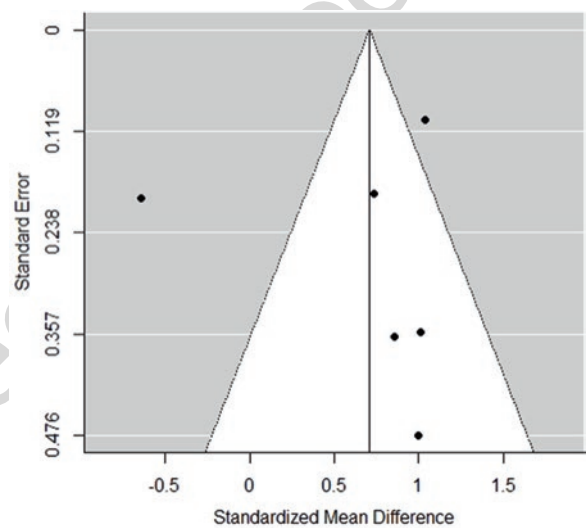
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**Fig. 25.6** Galbraith (radial) plot for standardized mean difference of trials using procalcitonin laboratory data



**Fig. 25.7** Funnel plot for verification of publication bias in the meta-analysis of standardized mean difference of procalcitonin laboratory data



**25.4 Discussion**

231 This study demonstrated that high levels of PCT  
 232 are associated with disease severity in patients  
 233 infected with COVID-19. PCT is known to be  
 234 elevated in bacterial infections and is currently  
 235 used for diagnosis and decision-making regard-  
 236 ing antibiotic treatment duration in respiratory  
 237 infections [25]. Its synthesis is upregulated in  
 238 bacterial infections and downregulated in viral  
 239 infections [26]. PCT is produced by the thyroid C  
 240 cells in healthy people. In the presence of bacte-  
 241 rial infections, PCT production is activated in all  
 242 parenchymal tissues and its level increases rap-  
 243 idly. PCT production by these tissues is stimu-

lated both directly by bacterial endotoxins and  
 lipopolysaccharides and indirectly by inflamma-  
 tory mediators that include tumor necrosis factor-  
 alpha (TNF-a), interleukin (IL)-1, and IL-6.  
 However, mediators of viral infection such as  
 interferon-gamma (IFN- $\gamma$ ) decrease the PCT  
 level, which makes it a more specific marker for  
 bacterial infections [26]. Nevertheless, the find-  
 ing of this study showed that PCT is increased in  
 patients with severe COVID-19. This suggests  
 that in some severe COVID-19 cases, there is a  
 bacterial co-infection that increases their PCT  
 levels. This hypothesis is supported by the work  
 of Zhou et al. in which it was reported that the  
 most of severe COVID-19 patients have viral

infection and secondary bacterial infection [16]. Bacterial co-infection is a poor prognostic feature in these patients [27] and may contribute to the death of these patients [28]. Thus, a PCT level determination, in addition to helping with identification of severe patients, may guide physicians in determinations of bacterial co-infection. This would allow them to initiate early antibiotic therapy that may prevent further deterioration of health.

The results of this study are contrary to the results of the study by Lippi and Plebani, which reported that the PCT value would remain within the reference range in severe coronavirus-infected patients [29]. However, the meta-analysis in the present study was of wider scope, including multiple studies that would have accounted for more patient variables. Nevertheless, further studies are required to address this issue.

## 25.5 Conclusion

The results of this study showed that PCT in patients with severe COVID-19 disease is increased, which suggests that it may play an important role in predicting severity and outcome of infection. Therefore, a PCT level determination may guide physicians in cases of suspected bacterial co-infection to initiate early antibiotic therapies that may prevent further deterioration of health and death.

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**Conflict of Interests** None.

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