Neutrophil CD64 as a diagnostic marker for neonatal sepsis: Meta-analysis

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Abstract

Background. Neutrophil CD64 (nCD64) is a promising marker for diagnosing bacterial infections. Several studies have investigated the performance of nCD64 for diagnosing neonatal sepsis and the results are variable. Interest in nCD64 for detecting serious bacterial infections is increasing rapidly.

Objectives. The aim of the present study was to carry out a meta-analysis to systematically evaluate the diagnostic accuracy of nCD64 in neonatal sepsis. As far as the authors know, no previous studies have undertaken this.

Material and methods. A review of studies from PubMed, Embase and the Cochrane Library, from inception through June 2015, found 7 studies (involving 2213 neonates) fulfilling the inclusion criteria. These 7 studies were subjected to a bivariate meta-analysis of sensitivity and specificity and a summary receiver operating characteristic (SROC) curve; $I^2$ was used to test heterogeneity, and the source of heterogeneity was investigated by influence analysis and meta-regression.

Results. The pooled sensitivity and specificity were 80% (95% CI, 69—88%) and 83% (95% CI, 71—90%), respectively. The area under the SROC curve (AUC) was 0.88 (95% CI, 0.85—0.91). The studies had substantial heterogeneity ($I^2 = 87.1$%).

Conclusions. The results showed that nCD64 is a reliable biomarker for diagnosing neonatal sepsis (AUC = 0.88).

Key words: meta-analysis, neonatal infection, CD64, neonatal sepsis
Neonatal sepsis is one of the most common causes of morbidity and mortality for neonates all over the world, particularly in developing countries. The incidence of neonatal sepsis is approximately 3–40 per 1000 live births, and the mortality rate ranges from 9% to 20%. It is difficult to identify neonatal sepsis early because of a lack of specific clinical manifestations. The signs are hard to distinguish from non-infectious disorders such as maladaptation, respiratory distress syndrome and aspiration syndromes. Blood culture is regarded as the reference standard for the identification of serious bacterial infection, but it is time-consuming (2–4 days) and has high false negative/positive rates. This means that broad-spectrum antibiotics are applied to all suspected neonates in case of potential serious outcomes. As a result, drug-resistant strains appear and neonatal healthcare costs escalate.

Several biochemical markers have been studied for the early diagnosis of neonatal sepsis, especially C-reactive protein (CRP) and procalcitonin (PCT). However, the specificity and the value of these markers are not sufficiently reliable. Therefore, a persistent search for better biomarkers of neonatal sepsis is still very necessary. CD64, a high affinity receptor that binds monomeric IgG, is normally expressed by monocytes and weakly on resting neutrophils. The expression of neutrophil CD64 (nCD64) is considered to be a very early phase of the host’s immune response to bacterial infection, increasing about one hour after invasion. It is stimulated by inflammatory cytokines, then increases in a graded manner. nCD64 expression remains stable for more than 24 h. The development of flow cytometric technology (FCM) has made it possible to measure nCD64 quickly and precisely with minimal blood volumes.

Interest in nCD64 for detecting serious bacterial infections is increasing rapidly. The performance of nCD64 in diagnosing neonatal sepsis has been investigated in several studies and the results are variable. Taking all the above into consideration, the aim of the present study was to carry out a meta-analysis to systematically evaluate the accuracy of nCD64 in diagnosing neonatal sepsis.

A study was considered eligible for inclusion in the present review if it provided data on nCD64 for neonates with or without sepsis. Moreover, nCD64 measurement had to be performed when suspected sepsis presented before antimicrobial therapy. In the septic group, patients had either culture-proven or clinically diagnosed sepsis; in the non-septic group, neonates had benign clinical disorders. Only studies written in English were included.

Furthermore, the studies had to provide sufficient information to construct a 2 x 2 contingency table with false and true positives and negatives provided. All studies that involved healthy neonates and patients older than 28 days were excluded. Animal experiments, reviews, correspondences, case reports, expert opinions and editorials were excluded.

Neonatal sepsis diagnosed in the first 72 h of life was considered early onset sepsis (EOS); after 72 h it was considered late onset sepsis (LOS). The Quality Assessment of Diagnostic Accuracy Studies (QUADAS) tool was used to assess the methodological quality of the studies included. If agreement could not be reached, differences were resolved by a 3rd investigator (ZM).

Material and methods

Search strategy and selection criteria

Two investigators systematically searched the PubMed, Embase and the Cochrane Library databases for studies that assessed the accuracy of nCD64 in the diagnosis of neonatal sepsis. The PubMed and the Cochrane Library combined search term used was (CD64) AND (neonatal sepsis OR neonatal infectious OR sepsis), and the Embase combined search term was (CD64) AND (sepsis). The databases were searched from their inception through June 2015.

A bivariate mixed-effects regression model was performed to synthesize the pooled sensitivity, specificity, positive/negative likelihood ratios (P/N LRs) and the diagnostic odds ratio (DOR). This model did not transform pairs of sensitivity and specificity of individual studies into a single indicator of diagnostic accuracy, but ensured the two-dimensional nature of the data, taking into account any correlations between pairs of studies. A summary receiver-operating characteristic (SROC) curve was also constructed, plotting sensitivity vs specificity, and...
the area under the curve (AUC) was calculated. Statistical heterogeneity among the studies was evaluated by I^2 statistics. Values of 25, 50 and 75% for the I^2 test were considered low, moderate and high statistical heterogeneity, respectively. The publication bias of the included studies was assessed by Deek’s funnel plot asymmetry test. The Spearman correlation coefficient between the logits of sensitivity and specificity was used to evaluate the presence of a threshold effect in the accuracy of nCD64. Fagan’s nomogram was used to calculate post-test probability (PTP). All the above analyses were performed using the Midas Module in Stata software, v. 12 (Stata Corporation, College Station, USA) and Metadisc 1.4 (XI Cochrane Colloquium, Barcelona, Spain). A p-value < 0.05 was considered statistically significant.

Results

Study selection process

The database search retrieved 308 studies. After reviewing the titles and abstracts, 266 articles were excluded, consisting of 98 duplicates, 15 case reports, 93 commentaries, 9 meta-analyses, 13 reviews, 12 meeting abstracts and poster presentations and 26 that did not investigate the diagnostic accuracy of neutrophil CD64 as a marker for sepsis. A further 35 were excluded after a full text review, leaving 7 studies for inclusion.19–25 The 35 articles included 22 in which the reference group or control group did not correspond to the definitions of the present meta-analysis, 2 that involved adult/pediatric or mixed populations and 13 for which 2 × 2 contingency tables could not be made (Fig. 1).

Characteristics of the studies included

Seven studies were included in the review. The 2213 neonates in these studies came from different parts of the world. Among these 2213 patients, 869 (39%) had sepsis (culture-proven or clinical) and 1344 were non-septic but with other critical conditions. The study population sizes ranged from 32 to 1156. All the studies were carried out in newborn intensive care units (NICUs) and nCD64 expression was measured using flow cytometry analysis. The types of study design were either prospective case-control or cohort studies. The quality of the 7 studies was generally high, satisfying the majority of the QUADAS criteria (Table 1).

Diagnostic accuracy of nCD64

Significant heterogeneity between studies was demonstrated (I^2 = 87.1%) for DOR. The pooled sensitivity of nCD64 for the diagnosis of neonatal sepsis was 80% (95%CI, 69–88%), and the specificity was 83% (95%CI,
71–90%) (Fig. 2). The pooled DOR was 19 (95%CI, 6–57), whereas the pooled P/N LRs were 4.6 (95%CI, 2.5–8.6) and 0.24 (95%CI, 0.14–0.41), respectively. The area under the SROC curve for CD64 was 0.88 (95%CI, 0.85–0.91) (Fig. 3). Fagan’s nomogram for likelihood ratios indicated that using nCD64 expression to diagnose neonatal sepsis increased the post-probability to 54% when the results were positive and reduced the post-probability to 6% when the results were negative (Fig. 4). The effect of the diagnostic threshold was not significant (p-value = 0.71 > 0.05). Deek’s funnel plot asymmetry test revealed the existence of publication bias with asymmetry in the data (p-value = 0.03 < 0.05) (Fig. 5).

Discussion

Neonatal sepsis is one of the most common causes of neonatal deaths. Diagnosing neonatal sepsis is a serious challenge, because there is no single test that can be used for its early confirmation or exclusion.14,26 Recently, many researchers have focused on nCD64 as a marker of neonatal sepsis.8,16,17 In the light of this, the current meta-analysis was undertaken to estimate the efficiency of nCD64 for diagnosing neonatal sepsis.

As noted earlier, PCT is a very promising diagnostic marker of neonatal sepsis.27 The sensitivity of PCT is 81% and the specificity is 79%. In the present study, the sensitivity and specificity of nCD64 were 80 and 83% respectively, which is similar to PCT. CRP is also an excellent marker and has been applied in clinical practice.28 The sensitivity of CRP ranges from 30 to 97%, and the specificity ranges from 75 to 100%.29 In the present meta-analysis, the sensitivity of nCD64 ranges from 57 to 89%, and the specificity ranges from 62 to 100%, indicating that nCD64 is a reli-
able marker in the diagnosis of neonatal sepsis. Positive and negative likelihood ratios (P/N LRs) and post-test probability (PTP) are also relevant for clinicians. They both show whether a patient with a positive or negative test actually has sepsis or not. A PLR of 4.6 indicates that a neonatal with sepsis is 4.6 times more likely to have a positive test result than a neonatal without. The PTP for a positive test result is 54% with a given pretest probability of 20%. Likewise, a NLR of 0.24 reduces the PTP to 6% for a negative result. The area under the SROC curve is 0.88. However, significant statistical heterogeneity exists in the analysis ($I^2 = 87.1\%$). Still, the interpretation of the above findings should not be ignored.

Several methods were tried to find the source of the high heterogeneity, including the threshold effect, publication bias, influence analysis and meta-regression. The different cutoff values for nCD64 did not account for the statistical heterogeneity through the analysis of threshold effect ($p = 0.71 > 0.05$). Deek’s funnel plot asymmetry test showed the existence of publication bias ($p$-value = 0.03 < 0.05), which is a source of heterogeneity. No valuable information was found through the sensitivity analysis and meta-regression. The meta-regression analysis included study design (prospective cohort or control-case study) and the gestational age of the neonates (preterm or not). The lack of a uniform definition of neonatal sepsis may potentially contribute to the high heterogeneity, especially for the clinically septic but culture-negative newborns, although the concept of clinical sepsis is widely used. This means that the selected studies use different criteria for the definition of sepsis. Thus, the spectrum of disorders and disease varies among the studies included. Considering that age may be a potential source of heterogeneity, studies in which over 15% of the neonates were older than 28 days were excluded. Studies involving healthy neonates as controls in whom nCD64 will not be applied in routine clinical testing were also excluded, so they cannot be representative of the population studied in the current meta-analysis.

This meta-analysis has several limitations. First, substantial heterogeneity was detected among the studies included, but none of the study characteristics accounts
for the majority of this heterogeneity. The studies differ in many ways, especially the definition criteria for neonatal sepsis and the postnatal age of the enrolled neonates. This important limitation will continue to exist in the research in this field until a uniform definition of neonatal sepsis is formulated. Second, a wide range of cut-off values in the reported nCD64 tests caused a wide variation in sensitivity and specificity. Even when the same method of measuring nCD64 expression was used, cut-off values were still different. Third, publication bias was detected. Studies with satisfactory results are more likely to be published, which can lead to overestimates of diagnostic accuracy. To overcome this problem, the authors searched again for further studies, but could not find additional relevant articles. Finally, only 1 study evaluated the performance of neutrophil CD64 in diagnosing EOS and only 2 evaluated it for LOS. The rest of the studies scarcely reported the percentage of EOS and LOS. So the accuracy of nCD64 for the diagnosis of EOS and LOS cannot be assessed.

In conclusion, nCD64 is a helpful marker for diagnosing neonatal sepsis. A study by Dhlamini et al. also shows that nCD64 has a high negative predictive value for excluding neonatal sepsis. But the results of nCD64 tests cannot be used alone to diagnose neonatal sepsis, as neonatal sepsis is a pathophysiological process rather than a specific syndrome and is too complex to be described by a single test. Further studies to determine the optimal cut-off values and to formulate a uniform definition of neonatal sepsis are urgently required.

References

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