

## ***N-myc* oncogene amplification in neuroblastoma: prognostic implications and the development of targeted therapies.**

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Amplification of *N-myc* oncogene in neuroblastoma: prognostic implications and development of targeted therapies

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### **Summary**

Neuroblastoma is the most common extracranial solid neoplasm in childhood, characterized by high clinical heterogeneity. This study aimed to investigate the correlation between *MYCN* oncogene amplification, disease staging, and prognosis, as well as to review emerging therapies. An integrative literature review was conducted in the PubMed, SciELO, and Google Scholar databases, selecting 10 articles published between 2018 and 2026. The results indicate that *MYCN* overexpression is the main predictor of poor prognosis and resistance, directing the patient to a high-risk protocol. Strategies such as targeted therapies (via PI3K/Akt/mTOR) and immunotherapies (anti-GD2 and CAR-T) emerge as ways to overcome this aggressiveness.

It is concluded that precision oncology is indispensable to the management of the disease, highlighting the strategic role of the biomedical scientist in molecular diagnosis and translational research.

**Descriptors:** Neuroblastoma. N-Myc Proto-Oncogene Protein. Prognosis. Molecular Targeted Therapy. Immunotherapy.

## **1 INTRODUCTION**

Neuroblastoma (NB) is the most prevalent extracranial solid neoplasm in the pediatric population, being the most common malignant tumor diagnosed in infants and children under five years of age. (Cochran et al., 2020). Characterized by remarkable biological heterogeneity, the NB presents a variable clinical spectrum, ranging from spontaneous regression in low-risk cases to aggressive metastatic progression that is refractory to treatment, in high-risk cases. This variation in biological risk assessment requires robust stratification systems, such as the International Risk Group of Neuroblastoma (INRGSS), which uses clinical and molecular factors to determine prognosis and the intensity of the therapeutic protocol (Van Nimwegen et al., 2022).

The basis of this aggressiveness lies in specific genetic alterations, with the amplification of oncogene *MYCN* is the most determinant biological marker of unfavorable prognosis (Zafar et al., 2021). Located on chromosome 2p24.3, *MYCN* acts as a potent transcription factor that, when overexpressed, it drives pathways of unrestricted cell proliferation, angiogenesis, and resistance to chemotherapy. Present in about 20% of cases, this genetic amplification allows the tumor cells avoid programmed cell death (apoptosis) mechanisms and activate signaling pathways crucial for tumor survival, such as the PI3K/Akt/mTOR pathway (Sharma et al., 2021).



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In this context, early identification of *MYCN* gene status plays a key role.

in clinical management and in the allocation of patients to intensive treatment protocols. Unlike

Unlike other tumors, *MYCN* amplification is such a robust predictor of poor prognosis that classifies the patient immediately into the high-risk category, regardless of the stage.

anatomical or age-related (Van Nimwegen et al., 2022). Accurate detection of this change through

Molecular biology techniques are essential, since *MYCN*-amplified patients exhibit

significantly lower rates of overall survival and event-free survival (Park et al.,

2018).

In addition to conventional stratification, the development of new therapeutic approaches has

The goal was to overcome the resistance induced by this genetic alteration. Among the emerging strategies,

Of particular note is immunotherapy with anti-GD2 monoclonal antibodies, which has become the standard.

maintenance therapy in high-risk cases, and CAR-T cell-based therapies, which represent

The frontier of cancer research by targeting specific neuroblastoma antigens (Cafferkey)

et al., 2024). The combined use of multimodal treatments and molecularly targeted therapies has been...

shown to be the key to improving survival and reducing morbidity and mortality associated with the subtypes.

more aggressive forms of the disease (Medscape, 2025).

Given this scenario, the present study aims to investigate the correlation between the

*MYCN* gene amplification, the staging and prognosis of neuroblastoma in children, as well

how to conduct a review of emerging therapeutic approaches that seek to neutralize the effects

Biological agents of this oncogene and overcome tumor aggressiveness.

## 2 METHODOLOGY

This study consisted of an integrative literature review. The bibliographic survey

It was carried out between August 2025 and June 2026, through a structured search in the databases of

SciELO, PubMed, and Google Scholar data, without language restrictions, considering articles.

published in the last five years. The following descriptors were used, obtained from the

Health Sciences Descriptors (DeCS) and Medical Subject Headings (MeSH): "Neuroblastoma",

"*MYCN* Amplification", "*N-Myc* Proto-Oncogene Protein", "Prognosis", "Targeted Therapy" and

"Immunotherapy", combined using the connectors "AND" and "OR". For the article search, it was

The following strategy was adopted, structured in the following databases: "*Neuroblastoma*" AND "*MYCN*

Amplification" OR "*N-Myc* Proto-Oncogene Protein" AND "Prognosis" AND "Targeted Therapy"

OR "Immunotherapy".

Studies related to neuroblastoma in pediatric patients were considered for inclusion;

studies that addressed the correlation between *MYCN* gene amplification and clinical prognosis; and studies that described emerging therapies, such as anti-GD2 antibodies and CAR-T cells, focused on high-risk treatment.

Studies conducted solely on animal models were not considered; studies that addressed other types of solid tumors unrelated to the neural crest; duplicate studies in the databases of data; and works that did not present the complete text available for analysis.

### 3 RESULTS

Based on a structured search conducted in the PubMed Central (PMC), SciELO, and Google databases.

Academically, 1,860 studies were identified. Of these, 340 articles were excluded because they were...

duplicates across platforms, leaving 1,520 studies for analysis. After a careful reading of

Of the titles, 1,150 articles were excluded because they did not have a direct relationship with molecular biology.

from *MYCN* or because they deal with pediatric neoplasms in general. In the abstract reading stage,

330 studies were excluded for not meeting the inclusion criteria — such as lack of focus.

in new therapeutic strategies or unavailability of the full text—totaling 40 articles.

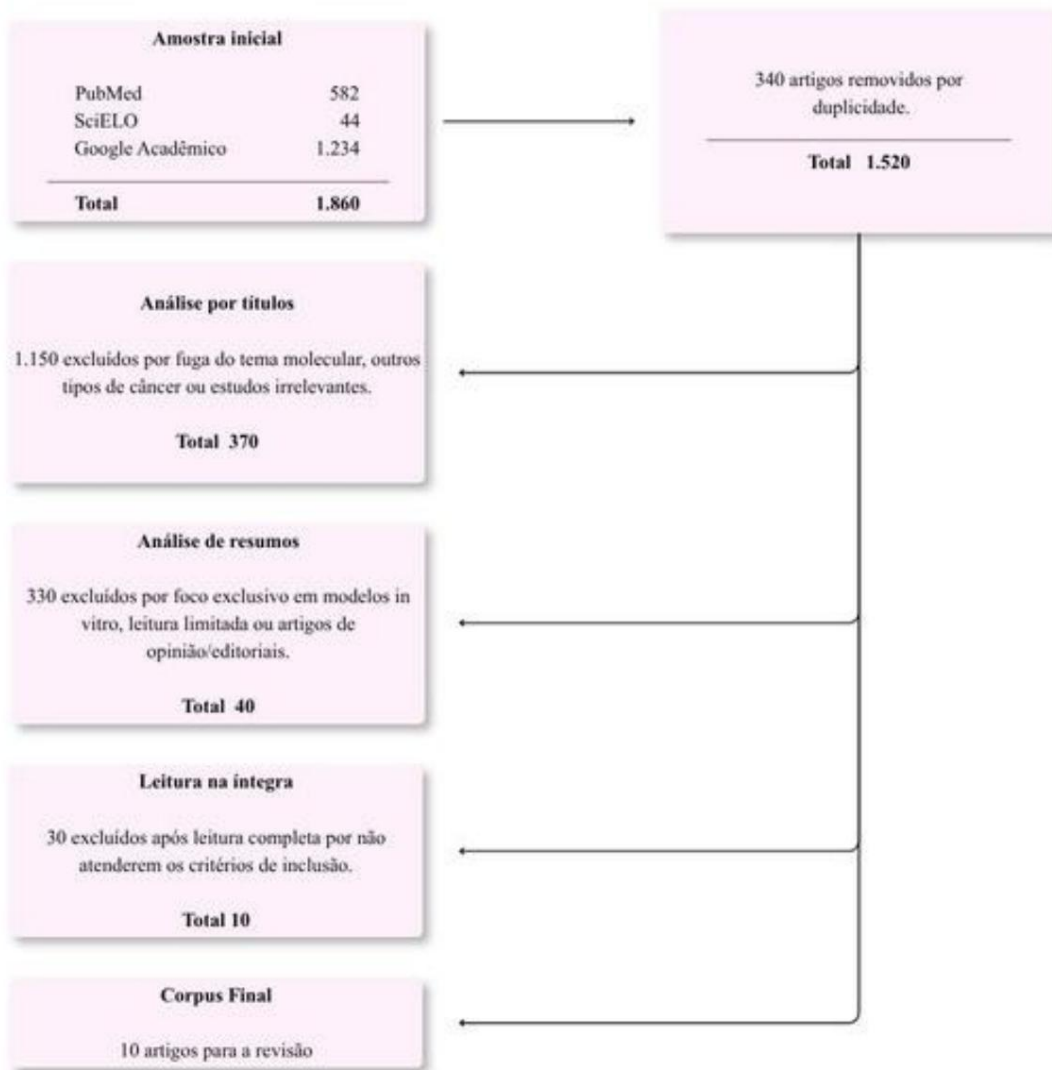
for full reading. After a complete and systematic analysis of the content, 10 were selected.

articles that best answered the research question, correlating *MYCN* amplification

with risk stratification and current therapeutic management, comprising the final corpus of this analysis.

(Figure 1).

Figure 1. Flowchart for article selection, 2026.



Source: Author's own work, 2026.

The systematic analysis of the 10 selected studies allowed for a critical overview of the role...  
 The central role of molecular biology in the clinical management of neuroblastoma. The data show that...  
 The aggressiveness of the disease is not a random factor, but rather the direct result of axis dysregulation.  
 specific molecular parameters, in which MYCN amplification acts as the main catalyst for  
 resistance.  
 Unlike traditional oncological approaches, which are primarily based on staging.  
 Anatomically, the corpus of this work reveals a transition towards precision oncology. The articles  
 They discuss three fundamental pillars: molecular characterization as a determinant of risk, the  
 interdependence between signaling routes (especially the PI3K/Akt/mTOR route) and strategies

border crossings, which seek to circumvent the historical difficulty of reaching MYCN directly through of innovative immunotherapies. A synthesis of this evidence, detailing the goals of each study and its... The respective findings are presented in Table 1.

Table 1. Characteristics of the selected scientific articles, according to title, authors, year of publication, focus of the investigation and results, 2026.

Title	Authors	Year	Focus of the investigation	Results
The role of the <i>MYCN</i> oncogene and its therapeutic implications in neuroblastoma	Sharma et al.	2021	Analyze the molecular biology of <i>MYCN</i> overexpression and strategies for neutralize its oncogenic function.	It disrupts cell survival genes; indirect inhibitors emerge as the main pathway to neutralize it.  the tumor.
Neuroblastoma: from molecular pathogenesis to therapy	Cochran et al.	2020	Review the molecular pathogenesis and evolution of neuroblastoma treatment protocols.	Risk stratification based on <i>MYCN</i> status is the central pillar for defining intensive multimodal protocols.  and
Targeting <i>MYCN</i> and neuroblastoma	Kappel et al.	2023	Investigating signaling pathways regulated by <i>MYCN</i> potential pharmacological targets.	<i>MYCN</i> acts in the control of metabolic pathways; the blocking of cooperating proteins (BET inhibitors) is effective in reducing tumor size.
CAR T-cell therapy for neuroblastoma: al. current status and future directions	Cafferkey et al.	2024	Describe the development and challenges of CAR-T cells in the treatment of high-risk neuroblastomas.	Targeted cellular immunotherapy It presents itself as a promising alternative for the control of persistent refractory diseases.  and
Prognostic	Van	2022	Assess the impact of amplification Patients with	

relevance of <i>MYCN</i> amplification in neuroblastoma	Nimwegen et al.		<i>MYCN</i> 's work on overall survival and amplification of event-free survival.	Genetic mutations exhibit drastically reduced survival rates. smaller, which consolidates <sup>the</sup> <i>MYCN</i> as a predictor of poor prognosis.
Molecular mechanisms of MYCN-driven neuroblastoma oncogenesis in	Zafar et al. 2021	Analyze the	cellular mechanisms by which <i>MYCN</i> overexpression promotes oncogenesis.	The <i>MYCN</i> protein It accelerates protein synthesis <sup>and</sup> and blocks <sup>the</sup> apoptosis, ensuring a proliferative advantage over <sup>and</sup> tumor aggressiveness.
Anti-GD2 Immunotherapy for Neuroblastoma: Current State and Future Directions	Park et al. 2018	Assessing	the clinical benefit of maintenance anti-GD2 immunotherapy on long-term survival with anti-GD2 monoclonal antibodies in	high-risk patients. GD2 increased significantly. event-free survival in high-risk groups.
Targeting the PI3K/Akt/mTOR pathway in neuroblastoma	Sharma et al.	2023 To	elucidate the role of the PI3K/Akt/mTOR pathway in <i>MYCN</i> -mediated aggression.	<sup>THE</sup> <sup>via</sup> PI3K/Akt/mTOR is fundamental <sup>the</sup> for tumor survival <sup>under</sup> Stress, making it a critical target for co-therapies.
New insights into the biology and treatment of high-risk neuroblastoma	Medscape (Org)	2025 Review	updates to the integration of treatment protocols and new molecular diagnostic tools.	allows <sup>one</sup> intervention <sup>and</sup> Early intervention tailored for high-progression phenotypes.

The Landscape of Kappel et al. 2022 Mapping new drugs in			targeted clinical trials in neuroblastoma resistant neuroblastomas.	The future of clinical management lies in targeted therapies that destabilize the MYCN protein and strengthen the host's immune response.

#### 4. DISCUSSION

A combined analysis of the literature reveals that neuroblastoma is no longer understood solely by... its clinical anatomy and began to be interpreted through its molecular signature. The central role The point of convergence among the consulted authors regarding the *MYCN* oncogene is that it is described by Zafar. et al. (2021) as a protein synthesis accelerator that blocks apoptosis, conferring to the cells tumors possess an unparalleled proliferative advantage. This biological aggressiveness translates directly into in epidemiological and clinical data; while Cochran et al. (2020) reinforce that NB is the tumor The most common extracranial solid in children is the presence of MYCN amplification, which determines... the clinical outcome.

The prognostic relevance of this genetic alteration is emphasized by Van Nimwegen et al. (2022), who Studies show that patients with *MYCN amplification* have higher overall survival rates. drastically lower. This view is corroborated by Cochran et al. (2020), who position the status MYCN as the central pillar for high-risk stratification, justifying the immediate adoption of more intensive and multimodal therapeutic protocols, regardless of the patient's age or of the tumor staging.

However, the historical challenge lies in the difficulty of directly targeting the MYCN protein, due to due to its "non-drugable" molecular structure. In light of this, Sharma et al. (2021) discuss the need of indirect strategies to neutralize its function. One of the most promising avenues for this The intervention is via the PI3K/Akt/mTOR pathway. According to Sharma et al. (2023), this signaling pathway is fundamental for the tumor to survive under stress conditions, making it a critical target for Co-therapies that aim to sensitize MYCN-amplified cells to conventional treatment. Complementing this view, Kappel et al. (2023) highlight that blocking cooperating proteins, Like BET inhibitors, it offers an effective alternative to reduce tumor burden by to destabilize the metabolic control exerted by the oncogene.

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The transition to precision oncology is also evident in the advancement of immunotherapies. Park et al. (2018) demonstrate that the introduction of anti-GD2 monoclonal antibodies significantly increased event-free survival in high-risk groups, establishing a new gold standard for the phase for treatment maintenance. Going further, Cafferkey et al. (2024) point to CAR-T cells as the current frontier, offering hope for cases of refractory or persistent disease, in which therapies Conventional treatments fail to eradicate the aggressive tumor clone.

Finally, the scenario outlined by Medscape (2025) and Kappel et al. (2022) suggests that the future of Clinical management of neuroblastoma depends on the integration of early molecular biopsies and... developing drugs that not only attack the tumor, but also strengthen the immune response.

The host's immune response. In short, current literature is unanimous: overcoming mediated aggression The *MYCN* approach requires a methodology that combines a deep understanding of cell biology with... Targeted therapies that destabilize the tumor's survival networks.

## FINAL CONSIDERATIONS

This study fulfilled its objective by correlating the amplification of the *MYCN* oncogene with the unfavorable clinical outcome in neuroblastoma, demonstrating that this biological marker is the central pillar for the immediate stratification of patients into the high-risk category. The review of Literature has shown that, although the molecular structure of *MYCN* makes direct blocking unfeasible, the development of targeted therapies directed at cooperating pathways (such as PI3K/Akt/mTOR and the BET proteins) and the advancement of immunotherapeutic frontiers (through anti-GD2 antibodies and cells) T-CAR) represent established strategies for overcoming tumor aggressiveness and resistance.

Thus, the theoretical hypotheses that the multimodal approach based on genetic signature

The improvements in event-free survival were confirmed by the scientific evidence analyzed.

The relevance of this research for biomedical professionals lies in the need to provide a foundation for...

His expertise lies in precision oncology and highly complex molecular diagnostics. Being the

Neuroblastoma is a rapidly progressive neoplasm, and the technical responsibility for its diagnosis falls to a biomedical professional.

through the accurate conduct and interpretation of molecular assays that detect *MYCN* status .

The accuracy of your laboratory report is the critical factor that determines the allocation of the pediatric patient.

in intensive therapeutic protocols. Furthermore, this study reinforces the role of the biomedical professional in

Translational research, since the elucidation of new signaling routes and monitoring by

The methods of molecular biopsies depend directly on the development of benchtop scientific research.

conducted by this professional.

As a suggestion for future research, clinical studies evaluating the following are recommended.



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long-term efficacy and toxicity profiles of co-therapies combining pathway inhibitors

Cellular treatment with CAR-T cells in patients with disease recurrence is also recommended.

To investigate the impact of incorporating early molecular screening methodologies into routine practice.

public laboratory, aiming to optimize the interval between initial diagnosis and the start of treatment.

targeted.

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