Systematic review and meta-analysis of the clinical survival significance of Sialyl-Tn expression in histological tissues from cancer patients

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Background: The expression of Sialyl-Tn (STn) has been associated with different mechanisms of cancer. Several studies using histological tissue samples from different types of cancer showed differences in the STn expression patterns, but few studies have explored the association between STn expression and decreased overall survival. Therefore, a comprehensive analysis of the clinical significance of STn in cancer patients was performed.

Methods: A systematic review and meta-analysis were done of scientific articles published and available from different sources, including MEDLINE/PubMed, Web of Science, Cochrane Library and LILACS. Odds ratio (OR) was represented by the number of deaths during follow-up in positive-STn versus negative-STn patients. OR from each study was pooled and the results were presented in a forest plot.

Results: Eleven articles were included in the systematic meta-analysis. The estimated OR of 3.26 (95% CI: 2.12–5.02) was obtained. Furthermore, subgroups studies of gastric cancer showed an OR of 3.74 (95% CI: 2.47–5.66) and other cancer types reported an OR of 2.92 (95% CI: 1.16–7.32).

Conclusions: This study showed that survival time was significantly longer in patients with negative-STn tumors than in those with positive results.

Keywords: Sialyl-Tn (STn); cancer; survival; meta-analysis


do: 10.21037/tcr.2019.11.53

View this article at: http://dx.doi.org/10.21037/tcr.2019.11.53

Introduction

It has been shown that abnormal glycosylation plays an important role during the carcinogenesis and can lead to the overexpression of mucin-associated carbohydrates as Sialyl-Tn (STn) antigen (1). The biosynthesis of STn involves different pathways, first the Tn antigen is produced through the addition of N-acetylgalactosamine in serine or threonine residues of a mucin protein. Once synthesized, Tn antigen suffers modifications that bounds to different structures such as STn antigen (Saα2,6GalNAcα-O-Ser/Thr) by addition of sialic acid, antigen T by addition of galactose (Galβ1,3GalNAcα-O-Ser/Thr) and Core 3 by addition of N-acetylglucosamine (2).

Over the last 40 years, research efforts were focused mainly on uncovering the role of STn antigen during cancer development. Through the 80s and 90s, authors aimed to describe the STn expression in different types of cancers and its clinical use (3). The results showed that expression of STn can be associated with different mechanisms of cancer, but mainly involved in the tumor
However, the activation of pathways can depend on the cancer type or subtype (5). These results brought the idea of using immunotherapy, based on anti-STn vaccine, as an attractive approach for the treatment of cancer patients. Therefore, in the early 2000s, some clinical trials (6) were developed in order to explore the potential of the anti-cancer vaccine targeting STn but, despite the great effort involved in the application of the vaccine, the clinical trial failed in phase III (3). Still, taking in consideration the positive and negative results obtained in the past, and the new information that has emerged in the last decade (7), it can be concluded that STn is an attractive target to be revised and contemplated in the design of novel immunotherapies.

Several studies using histological tissue samples from different types of cancer showed differences in the STn expression patterns (Figure 1). Reported data from histological tissues noted consistent high frequencies in colorectal (8), colon (9), gastric (10) and pancreatic cancer (11). Other studies registered some variability in the expression of STn, as for example in breast (12), ovarian (13), bladder (14), liver (15) and cervical cancer (16). In contrast, in normal tissues, STn expression is found to be rare or low compared to the matching cancer cells (17). This heterogeneity can be due to several factors including antibodies used, staining protocol, number of samples evaluated, among others. Many monoclonal antibodies have been developed for STn detection, however B72.3 and TKH2 are the most commonly used anti-STn antibodies according to the literature (18). Overall, it can be concluded that STn expression is higher in the majority of cancers than in healthy tissues. In fact, STn has been postulated as a good tumor marker of carcinogenesis and potentially useful for diagnosis.

In the literature above, only a few studies have explored the association between STn expression and decreased overall survival, but contradictory results can be found among cancer types and subtypes. Therefore, more studies are needed in order to obtain reliable and accurate information regarding the prognostic value of STn in cancer. The very few data available assessing the prognostic value of STn antigen in carcinomas drove us to determine the association between the expression of this mucin-associated carbohydrate antigen and patients survival. For this reason a comprehensive analysis of the clinical significance of STn in cancer patients was performed using a systematic literature review and meta-analysis.

**Methods**

**Study design and search strategy/data source**

A systematic review was performed independently by three investigators (ALGS, RL, AAN) using published articles on the clinical significance of the expression of STn in histological tissue samples from cancer patients. The studies were identified through searching in English terms using the online sources of MEDLINE/PubMed, Web of Science, Cochrane Library and LILACS. The grey literature was searched using Google Scholar. The following keywords were taken from the Medical Subject Headings library of PubMed and used in combination: “Sialyl-Tn antigen” AND “Neoplasms” AND “Mortality”. The search strategy for Medline was developed first and then adapted for the remaining sources. The search was done for titles and/or abstracts and was carried out between August 2018 and February of 2019.

**Selection of studies**

Three investigators selected the studies for the systematic review and meta-analysis if they reported on the expression of STn using histological tissue samples from carcinoma samples. A total of 108 relevant studies were identified and exported to EndNote X9 (Philadelphia, USA). After removing duplicates, 79 studies were considered as candidates for this review. Then, the following exclusion
criteria were applied: (I) STn expression from serum; (II) studies evaluating immunotherapy; (III) studies from cell lines or animal models; (IV) reviews and letters to the editors; (V) manuscripts without abstract; (VI) no manuscript available. The inclusion criteria for our study was: (I) STn expression from formalin-fixed, paraffin-embedded specimens of carcinomas samples; (II) using TKH2 or B72.3 monoclonal antibodies; (III) including survival and mortality data; (IV) prospective cohort or retrospective study design; (V) studies with data of at least five-years follow-up (Figure 2).

After preliminary screening of the titles of the studies, all abstracts were evaluated for eligibility (some studies were assessed in full length), based on the established criteria. There was no restriction in the year of publication, recruitment period or sample size.

Data extraction

The present study was done based on the criteria of Preferred Reporting Items for Systematic Reviews and Meta-Analysis (PRISMA) (Figure 2). Data from all the selected articles were extracted independently by two investigators (ALGS and RL) into a predefined database, including first author information, year of publication, sample type and size, antibody used, percentage of STn positive and negative (STn+ and STn–) expression, survival data and follow-up of patients (Table 1). GetData Graph Digitizer 2.26 was used to extract survival data from studies that only reported the analysis by Kaplan-Meier graphs; mortality was defined as a death of the patient or recurrence of the disease in period of time of 5 years. In case of discrepancy among investigators, the final consensus was done by another investigator (AAN). The quality of the methodology used in each study was evaluated using The Newcastle-Ottawa Scale (NOS) (19) (Table S1).

Statistical analysis

Odds ratio (OR) was represented by the number of deaths during follow-up in STn+ versus STn– patients. Five-year follow-up data were extracted; otherwise, the maximum reported follow-up was considered. All statistical analyses were undertaken using the software RStudio version 1.0.153. OR for mortality, comparing STn+ with STn–, was pooled using a random-effects model and afterwards presented in a forest plot with corresponding 95% confidence interval (CI). Statistical heterogeneity was assessed using the I² index. Finally, publication bias was assessed by visual inspection of funnel plots and with the Egger bias test (20).
<table>
<thead>
<tr>
<th>#</th>
<th>Author/study</th>
<th>Population</th>
<th>Antibody (primary)</th>
<th>Sialyl-Tn, n (%)</th>
<th>Death, n (%)</th>
<th>Survival, n (%)</th>
<th>Follow-up (months)</th>
</tr>
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<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>STn+</td>
<td>STn–</td>
<td>Death+</td>
<td>Death–</td>
<td></td>
</tr>
<tr>
<td>1</td>
<td>Ma et al. (1993). Expression of sialyl-Tn antigen is correlated with survival-time of patients with gastric carcinomas</td>
<td>85 gastric carcinomas: 19 stage-I, 17 stage-II, 33 stage-III and 16 stage-IV</td>
<td>TKH2</td>
<td>53 (62.35%)</td>
<td>32 (37.64%)</td>
<td>28 (32.94%)</td>
<td>5 (5.88%)</td>
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<td>(32.94%)</td>
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<tr>
<td>2</td>
<td>Werther et al. (1994). Mucin-associated sialosyl-Tn antigen expression in gastric cancer correlates with an adverse outcome</td>
<td>31 gastric carcinomas: 11 well, 8 moderately, 8 poorly differentiated and 4 mucinous</td>
<td>TKH2</td>
<td>21 (67.74%)</td>
<td>10 (32.25%)</td>
<td>18 (58.06%)</td>
<td>8 (25.80%)</td>
</tr>
<tr>
<td>3</td>
<td>Yamada et al. (1995). Sialosyl Tn antigen expression is associated with the prognosis of patients with advanced gastric cancer</td>
<td>53 gastric carcinomas</td>
<td>TKH2</td>
<td>27 (50.94%)</td>
<td>26 (49.06%)</td>
<td>18 (33.96%)</td>
<td>10 (18.86%)</td>
</tr>
<tr>
<td>4</td>
<td>Miles et al. (1995). Expression of sialyl-Tn in gastric cancer: correlation with known prognostic factors.</td>
<td>147 gastric carcinomas: 38 well, 49 moderately, and 71 poorly differentiated (survival analysis: 139 cases)</td>
<td>B723</td>
<td>95 (68.34%)</td>
<td>44 (31.65%)</td>
<td>90 (64.74%)</td>
<td>35 (25.17%)</td>
</tr>
<tr>
<td>5</td>
<td>Victorzon et al. (1996). Sialyl Tn antigen is an independent predictor of outcome in patients with gastric cancer.</td>
<td>242 gastric carcinomas; 74 stage-I, 59 stage-II, 73 stage-III and 36 stage-IV disease. (survival analysis: 237 cases)</td>
<td>B72.3</td>
<td>148 (62.44%)</td>
<td>89 (37.55%)</td>
<td>111 (46.83%)</td>
<td>41 (17.29%)</td>
</tr>
<tr>
<td>6</td>
<td>Itzkowitz et al. (1990). Sialosyl-Tn. A Novel Mucin Antigen Associated with Prognosis in Colorectal Cancer Patients</td>
<td>128 colorectal carcinomas</td>
<td>TKH2</td>
<td>112 (87.50%)</td>
<td>16 (12.50%)</td>
<td>30 (23.42%)</td>
<td>0 (0%)</td>
</tr>
<tr>
<td>7</td>
<td>Franchi et al. (1996). Prognostic implications of Sialosyl-Tn antigen expression in sinonasal intestinal-type adenocarcinoma</td>
<td>30 intestinal-type adenocarcinoma of the nasal cavities and paranasal sinuses</td>
<td>TKH2</td>
<td>21 (70%)</td>
<td>9 (30%)</td>
<td>16 (53.33%)</td>
<td>2 (6.66%)</td>
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<td>(53.33%)</td>
<td>(6.66%)</td>
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<td>8</td>
<td>Terasawa et al. (1998). Expression of Tn and Sialyl-Tn Antigens in the Neoplasia Transformation of Uterine Cervical Epithelial Cells</td>
<td>83 cervical invasive carcinomas</td>
<td>TKH2</td>
<td>42 (50.60%)</td>
<td>41 (49.39%)</td>
<td>9 (10.84%)</td>
<td>7 (8.43%)</td>
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<td>(49.39%)</td>
<td>(49.39%)</td>
<td>(10.84%)</td>
<td>(8.43%)</td>
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<tr>
<td>9</td>
<td>Ghazizadeh et al. (1997). Mucin Carbohydrate Antigens (T, Tn, and Sialyl-Tn) in Human Ovarian Carcinomas: Relationship With Histopathology and Prognosis</td>
<td>38 ovarian carcinomas</td>
<td>B72.3</td>
<td>32 (84.21%)</td>
<td>6 (15.78%)</td>
<td>26 (68.42%)</td>
<td>2 (5.26%)</td>
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<td>(15.78%)</td>
<td>(15.78%)</td>
<td>(68.42%)</td>
<td>(5.26%)</td>
</tr>
<tr>
<td>10</td>
<td>Takao et al. (1999). Mucin Core Protein Expression in Extrahepatic Bile Duct Carcinoma Is Associated with Metastases to the Liver and Poor Prognosis</td>
<td>67 bile duct carcinomas</td>
<td>TKH2</td>
<td>54 (80.59%)</td>
<td>13 (19.40%)</td>
<td>39 (58.20%)</td>
<td>11 (16.41%)</td>
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<td>(19.40%)</td>
<td>(19.40%)</td>
<td>(58.20%)</td>
<td>(16.41%)</td>
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<tr>
<td>11</td>
<td>Costa et al. (2015). Abnormal Protein Glycosylation and Activated PISK/Akt/mTOR Pathway: Role in Bladder Cancer Prognosis and Targeted Therapeutics</td>
<td>96 bladder carcinomas, 49 invasive</td>
<td>TKH2</td>
<td>57 (59.37%)</td>
<td>39 (40.62%)</td>
<td>30 (31.25%)</td>
<td>10 (10.41%)</td>
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</table>

STn, Sialyl-Tn; STn+, positive staining for Sialyl-Tn; STn–, negative staining for Sialyl-Tn; Death+, death among positive staining for Sialyl-Tn; Death–, death among negative staining for Sialyl-Tn, (%) Percentage based on total population.
Results

Identified studies

The conducted literature search displayed a total of 108 articles. Once duplicates were eliminated, 79 relevant records were identified and their titles and abstracts reviewed. After applying the inclusion and exclusion criteria, 11 articles were included for this systematic review and meta-analysis (Figure 2 and Table 1).

Characteristics of studies

Eleven articles reporting on the association between positive-STn staining in histological tissue samples and cancer patient’s overall survival were included. The year of publication of the studies ranged from 1990 to 2015, therefore, the recruitment period of the participants was different across the selected studies. Five of eleven articles were carried out in Japan (13,15,16,21,22) and two in United Stated of America (8,23). Additionally, single studies were performed in Finland (5), Portugal (14), United Kingdom (24) and Italy (25).

All articles assessed the expression of STn by immunohistochemical staining using monoclonal antibodies (TKH2 or B72.3). Carcinoma tumors analyzed ranged from stage I to IV, and included samples from primary tumors and metastatic lesions. Depending of the cancer type, the results were organized in the following categories: five studies of gastric cancer (5,21-24); one study of colorectal cancer (8); one of bladder cancer (14); one of ovarian cancer (13); one of sinonasal cancer (25); one of cervical cancer (16), and finally one of liver cancer (15). A total of 662 patients samples were positive and 325 patients were negative for STn.

The mortality in cancer patients was determined by the percentage of deaths in the STn+ population and in the STn− one. Among gastric cancer patients, mortality or recurrence was found in 77.03% (265/344) patients with STn+ and 46.77% (94/201) that were STn−. In colorectal cancer, 26.79% (30/112) with STn-positive staining and no deaths were found in patients with STn-negative (0/16). Mortality of 52.63% (30/57) among STn+ expression group was reported in bladder cancer patients and 25.64% (10/39) with STn−; in ovarian cancer 26 deaths were found among patients with STn-positive and 2 more with STn-negative (81.25% and 33.33%, respectively). In sinonasal cancer, 76.19% (16/21) were positive for STn and 22.22% (2/9) were negative for STn; and for cervical cancer, it was reported 9/42 deaths with positive and 7/41 with negative staining for STn (21.42% vs. 17.07%). In liver cancer, both positive (39/54) and negative (11/13) STn expressing patients, showed high mortality rates (72.22% vs. 84.61%) (Table 1 and Figure 3).

OR of survival

An OR of 3.26 was obtained, when all studies among different types of cancers were pooled, meaning that those patients positive for STn have three times higher probability of death comparing with those that are negative.
(OR: 3.26; 95% CI: 2.12–5.02) (Figure 3). No publication bias was found (Egger’s test =0.12; P=0.89), as also shown by the funnel plot (Figure 4). Following the general analysis, cancer subgroups were analyzed. When studies reporting gastric cancer were grouped, the results exposed an OR of 3.74 (95% CI: 2.47–5.66) (Figure 5). The remaining articles were grouped together as other cancer types, showing an OR of 2.92 (95% CI: 1.16–7.32) (Figure 6).

**Discussion**

In the present study, a total of eleven articles were analyzed including a total of 987 histological tissues from patients with different cancer types, showing STn+ expression in
of STn with an increased risk of mortality (5,8,13,14,21,22). Based on our results, positive-STn is related with a strong clinical impact in cancer (OR =3.26), it shows that survival time was significantly longer in patients with negative-STn tumors than in those with positive results (Figure 3). The heterogeneity of this study was low (I² =27%), suggesting a correlation of positive-STn expression with a higher incidence of aggressiveness and decreased survival of patients with different cancer types, across the selected studies.

Conclusions
To our knowledge this is the first systematic review and meta-analysis that shows evidence of the association between the expression of STn in survival patients from different types of cancers. Results from this study confirm that there is scientific evidence of the positive relation between STn expression and poor prognosis in different types of cancer from histological tissue samples. Currently, research groups are working on uncovering the potential of the STn antigen combined with markers of proliferation (Ki-67) (28), estrogen receptor (ER) (29), and their relationship with circulating tumor cells (CTC) (30) in early cancer diagnosis. In addition, with this new evidence and information, STn has again drawn the attention to be used as a potential candidate for immunotherapy (7), that might lead into the development of novel and effective treatment for cancer patients.

Acknowledgments
We are very grateful to M.Sc. Julio Vega for helping with the statistical analysis.

Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was approved by Research Committee and the Ethics Committee from the Regional High Speciality Hospital of the Yucatan Peninsula (No. CONBIOETICA-31-CEI-002-20170731).
References


Table S1 Newcastle Ottawa scale for assessment of quality of included studies

<table>
<thead>
<tr>
<th>Studies</th>
<th>Selection</th>
<th>Comparability</th>
<th>Outcome</th>
<th>Conclusion</th>
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<tbody>
<tr>
<td>Ma et al. (1993)</td>
<td>☆☆</td>
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<td>☆☆☆☆</td>
<td>Fair</td>
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<td>Werther et al. (1994)</td>
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<td>Yamada et al. (1995)</td>
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<td>Yamada et al. (1995)</td>
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<td>Victorzon et al. (1996)</td>
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<td>Itzkowitz et al. (1990)</td>
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<td>Takao et al. (1999)</td>
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<td>Costa et al. (2015)</td>
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<td>Ghazizadeh et al. (1997)</td>
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<td>Franchi et al. (1996)</td>
<td>☆</td>
<td>☆☆</td>
<td>☆☆☆☆</td>
<td>Good</td>
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☆ represents if individual criterion within the subsection was fulfilled. A study can be awarded a maximum of four stars for selection; two stars for comparability and three stars for outcome. The total score of the studies were rated as good, fair, or poor quality. Thresholds for converting the Newcastle-Ottawa scales to AHRQ standards (good, fair, and poor) (19): Good quality: 3 or 4 stars in selection domain AND 1 or 2 stars in comparability domain AND 2 or 3 stars in outcome/exposure domain. Fair quality: 2 stars in selection domain AND 1 or 2 stars in comparability domain AND 2 or 3 stars in outcome/exposure domain. Poor quality: 0 or 1 star in selection domain OR 0 stars in comparability domain OR 0 or 1 stars in outcome/exposure domain.