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Prevalence and clinical correlations of somatostatin receptor-2 (SSTR2) expression in neuroblastoma.

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Abstract

Alternative radiolabeled targeted agents are being investigated for children with relapsed neuroblastoma who do not respond to I¹³¹-metaiodobenzylguanidine (MIBG) therapy. DOTATATE targets somatostatin receptors (SSTR), particularly SSTR2, which are expressed on neuroblastoma cells. We investigated SSTR2 expression in neuroblastoma tumors (36 high risk; 33 non-high risk patients) and correlated SSTR2 levels with clinical features, norepinephrine transporter (NET) expression and MIBG avidity. SSTR2 and NET immunohistochemistry scores (0-3) were calculated on biopsies using digital image analysis based on staining intensity and distribution. Clinical data were correlated with SSTR2 expression. Median SSTR2 score for 69 patients was 1.31 (0.26-2.55). Non-high risk neuroblastoma was associated with higher SSTR2 score (p=0.032). SSTR2 expression did not correlate with age, INSS stage, *MYCN* amplification and histology. Higher SSTR2 scores were observed in MIBG-avid versus non-avid NB. SSTR2 score were not significantly associated with NET score (r=-0.062, p=0.62). Twenty-six patients who relapsed or progressed had a median SSTR2 score of 1.33 (0.26-2.55). Patients with neuroblastoma including relapsed or progressive disease showed SSTR2 expression at diagnosis, suggesting they could be candidates for radiolabeled DOTA-conjugated peptide imaging or therapy.

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Keywords

Neuroblastoma; somatostatin receptor-2; SSTR2; norepinephrine transporter

Introduction

The outcome for high risk (HR) neuroblastoma (NB) patients who relapse remains poor, with 5-year overall survival (OS) of only 4-8%.^{1,2} Personalized therapeutic approaches for patients with relapsed NB consist of cytotoxic chemotherapy regimens for example, irinotecan and temozolomide; immunotherapy agents such as dinutuximab; radiolabeled targeted therapy including I¹³¹-metaiodobenzylguanidine (I¹³¹-MIBG); and molecularly targeted agents for example, crizotinib for ALK-mutated tumors³⁻⁷. Current therapies for relapsed neuroblastoma have response rates of approximately 15-50% with better responses seen in localized non-high-risk patients and those with late relapse.^{1-3,8-11} However, in the patients that are salvaged from first relapse, 50-90% will have a further relapse within 5 years and time for disease control shortens with each subsequent relapse.^{1,11,12} I¹³¹-MIBG has shown efficacy and symptomatic clinical response in one-third of patients with relapsed NB, and alternative radioactive targeted agents are being investigated for patients that do not respond to MIBG therapy.⁹

Transmembrane G-protein-coupled receptors for somatostatin, a regulatory hypothalamic peptide, are detected in many tissues and are highly expressed in neuroendocrine tumors and other cancers.¹³⁻¹⁷ All five somatostatin receptor subtypes have been detected in neuroblastoma tumors, particularly somatostatin receptor-2 (SSTR2), using different in vitro and in vivo techniques including immunohistochemistry, autoradiography (with radioactive somatostatin analogues), Western blot and reverse transcriptase polymerase chain reaction (RT-PCR).^{15,18-21} There may be intratumoral and intertumoral variations in SSTR expression due to the heterogeneity within neuroblastoma.²⁰ Increased somatostatin and the presence of high affinity SSTRs are associated with more differentiated *MYCN*-non-amplified neuroblastic tumors and more favorable disease.^{14,19,22,23}

Somatostatin analogues have been used in diagnosis and treatment of neuroendocrine tumors for many years.^{23,24} Radiolabeled DOTA-conjugate peptides bind to specific SSTRs and have demonstrated superior sensitivity and specificity for clinical detection of neuroendocrine tumors when combined with positron emission tomography (PET)/computerized tomography (CT), compared to traditional octreotide scintigraphy.^{25,26} Early phase studies with radiolabeled-DOTA-conjugate peptides such as ¹⁷⁷Lu-DOTATATE and ⁹⁰Y-DOTATOC have demonstrated efficacy and safety in adults with SSTR-positive tumors.²⁷

DOTATATE ((DOTA⁰-Tyr³) octreotate) has a high specificity for SSTR2, and when combined with radionuclide Gallium-68 (⁶⁸Ga) and Lutetium-177 (¹⁷⁷Lu) is a potential candidate for targeted diagnosis and therapy for neuroblastoma.²⁸⁻³⁰ We recently demonstrated higher uptake of ⁶⁸Ga-DOTATATE in high-expressing SSTR2 NB xenografts compared to low-expressing SSTR2 NB xenografts, using micro-PET CT and autoradiography.³¹ In addition, histological co-localization of SSTR2 and ⁶⁸Ga-DOTATATE

was demonstrated in high-expressing SSTR2 NB xenografts using immunohistochemistry and autoradiography, and tumor growth inhibition was achieved using ^{177}Lu -DOTA-TATE in high-expressing SSTR2 xenograft models. Preliminary studies using DOTA-conjugates for imaging and treatment in children with relapsed and refractory neuroblastoma have demonstrated safety and some clinical benefit; however, the NB patient subgroup most likely to respond to treatment is unknown.^{29,30,32} In addition, MIBG non-avid NB patients, who would require alternative therapy to ^{131}I -MIBG and who are potential candidates for ^{177}Lu -DOTATATE, have low expression of norepinephrine transporter (NET)³³, but it is not known whether their tumors express SSTR2 or if SSTR2 and NET expression levels are correlated.

This study aimed to describe the prevalence of SSTR2 expression in neuroblastoma tumors and correlate SSTR2 expression with clinical features, clinical outcome, norepinephrine transporter expression and MIBG avidity.

Materials and Methods

Sixty-nine patients with neuroblastoma, 36 high-risk (HR) and 33 non-high risk (NHR), from Children's Oncology Group (COG) high-risk study A3973³⁴(N=16) & The Hospital for Sick Children (HSC) (N=53) were included in the study. Tumor samples from diagnosis in the COG study (N=16) were selected from the same cohort of patients (N=27) used in a prior publication from Dubois et al, describing NET protein expression and MIBG avidity.³³ For these 16 patients, results of the clinical features and centrally reviewed MIBG scans were obtained from the COG study database. For the remaining patients, clinical features and outcomes were obtained from HSC electronic health records, including local MIBG reports.

15 NB patients from the HSC cohort had post-treatment samples analysed in addition to their diagnostic pre-treatment biopsy. Post-treatment samples were taken at varying time intervals during therapy and not specifically at relapse.

Immunohistochemistry

SSTR2 and NET immunohistochemistry staining was performed for all patient samples (N=69). NET expression was done using the same methodology and antibody as the COG study.³³ Formalin-fixed, paraffin-embedded neuroblastoma specimens were sectioned at 5 μm and slides were subjected to heat-induced epitope retrieval with 0.01M Citrate Buffer pH 6 and blocked with 3% methanolic hydrogen peroxide followed by 10% normal horse serum. Slides were then incubated with rabbit anti-SSTR2 antibody (Epitomics #3582-1) (1:100 dilution) or mouse anti-NET antibody (MAb Technologies #NET17-1) (1:1000 dilution) for 60 minutes. Antibody localization was detected using the ImmPRESS Anti-Rabbit or Anti-Mouse Ig Peroxidase Reagent Kit (Vector Laboratories, #MP-7401 or #MP-7452) and visualized with 3,3'-diaminobenzidine. Sections were then counterstained with hematoxylin. Cerebellum and adrenal medulla was used as the positive controls for SSTR2 and NET, respectively. Tonsil was used as the negative for both SSTR2 and NET.

SSTR2 and NET immunostaining were digitally analyzed in a quantitative fashion using Aperio Image Scope v9.1.19.1567 and IHC Membrane Algorithm version 8.001. The staining intensity thresholds were set as follows: strong (3+) 210, moderate (2+) 180, weak (1+) 160. The entire tumour area on each section was scanned by the computer and the total number of cells for each staining intensity calculated. For each antibody, a modified H-score was calculated for each tumor with the following formula: $[(1 \times (\% \text{ cells } 1+) + 2 \times (\% \text{ cells } 2+) + 3 \times (\% \text{ cells } 3+)]/100$. The scores lay on a continuous scale from 0 to 3.

Statistical analysis

Wilcoxon rank-sum tests were performed to assess the correlation of SSTR2 expression with age at diagnosis (<18 months vs. 18 months), INSS stage (non-stage 4 vs. stage 4)^{35,36}, *MYCN* status (non-amplified vs. amplified), International Neuroblastoma Pathology Classification (INPC) histology (favorable vs. unfavorable)³⁷, MIBG avidity (non-avid vs. avid) and risk group (low/intermediate vs. high). (Table 1) The Children's Oncology Group (COG) neuroblastoma risk stratification definitions were used to describe low-, intermediate- and high-risk as per P9641, A3961 and A3973 studies.^{34,38,39}

A Spearman correlation coefficient was calculated between SSTR2 and NET scores. (Figure 2) Wilcoxon ranked sum test was used to compare SSTR2 expression pre and post treatment, as well as NET.

Results

Fifty-two percent of patients had high-risk NB (36/69) with 52% of tumors showing unfavorable histology according to INPC (N=36/68) and 20.3% were *MYCN*-amplified (N=14). (Table 1) The median score for SSTR2 expression for the whole cohort was 1.31 (range: 0.26-2.55). (Figure 1) Risk group was the only clinical feature significantly associated with SSTR2 score (p=0.032), with lower scores for the 36 high-risk (HR) patients (median=1.16) compared to the 33 non-high risk (NHR) low or intermediate risk patients (median=1.54). (Table 1)

There were 53 MIBG-avid (76.8%) and 16 MIBG-non-avid (23.1%) patients. There was a trend for higher SSTR2 scores in MIBG-avid (median=1.417) versus non-avid NB patients (median=0.954) but this did not meet statistical significance (p=0.077). (Table 1) SSTR2 scores were not statistically significantly associated with age, INSS stage, *MYCN* amplification or histology. (Table 1)

The median NET score of the cohort was 1.20 (range: 0.10-2.52). SSTR2 score was not significantly associated with NET score ($r=-0.062$, $p=0.62$) (Figure 2). Of the 69 patients in the cohort, there were 20 who relapsed and 6 who progressed. (Table 2) At initial diagnosis, 18/26 of relapsed/refractory patients were high-risk, 5/26 were *MYCN*-amplified, and 23/26 were MIBG-avid. The median SSTR2 score of the patients who relapsed was 1.32 (range: 0.26-2.55) and median NET score was 1.17 (range: 0.40-1.80). There were 2 patients with events unrelated to NB that were not included: one developed a secondary malignancy and the other died without disease relapse or progression.

For the 15 NB patients (9 HR and 6 NHR) from the SickKids cohort who had pre- and post-treatment samples, median SSTR2 score for the pre-treatment group was 1.34 (0.28-2.45) and post-treatment was 1.44 (0.66-2.55). There was no significant difference between pre- and post-treatment SSTR2 scores ($p=0.61$). Median NET score for pre-treatment group was 1.21 (0.45-1.80) and post-treatment was 2.00 (0.62-2.51). Post-treatment NET scores were significantly higher than pre-treatment ($p=0.01$).

Discussion

This study used immunohistochemistry (IHC) on tumor samples from a large cohort of mixed-risk neuroblastoma patients at initial diagnosis to determine the prevalence of SSTR2 expression. This is the first study in NB to determine a quantitative SSTR2 IHC score by digital analysis of specimens. We showed a range of SSTR2 expression seen in a variety of neuroblastoma tumors at time of diagnosis. (Figure 1) Although previous smaller studies in NB have demonstrated SSTR2 expression in 77-90% of neuroblastoma tumors, they vary in the definition of SSTR2 positivity, analytical techniques (autoradiography, RT-PCR and IHC) used and stage(s) of disease.^{15,20,21} In addition, some of the assays only detect total SSTR2 expression of a sample (Western immunoblot and RT-PCR), while IHC can determine both level and percentage of positively stained cells.

The SSTR2 score in this study was calculated on a continuous scale of 0-3 to incorporate the percentage of cells staining at different intensities across the tumor sample. Although there is no standard IHC scoring system for SSTR2, semi-quantitative scoring systems such as Allred score, H-score and immunoreactive score (IRS), have been used to assess protein expression of a range of tumor biomarkers, including epidermal growth factor receptor and estrogen receptors.⁴⁰⁻⁴² In keeping with the DuBois et al NB study evaluating NET expression and MIBG, we used the methodology for immunohistochemical staining for NET and had a similar approach to NET and SSTR2 scoring by taking into account staining intensity and percentage of positive cells.³³ However, in our study IHC scores were determined by image analysis and quantified the number of tumor cells positive for each intensity (0-3), rather than a product of manual determination of percentage positive cells and average staining intensity (0-300).³³

We found SSTR2 protein expression levels to be significantly higher in low/intermediate risk in comparison to high-risk neuroblastoma tumors. This is consistent with other studies in which SSTRs have been shown to be up-regulated in more differentiated disease, with increased somatostatin concentrations correlating with NB cell differentiation.^{14,22,23} SSTRs have also been shown to be down-regulated following disease progression. Although SSTR2 expression has been reported to be associated with favorable histology and prognosis, using our quantitative immunostaining method we did not find SSTR2 to be significantly associated with favorable histology, lack of *MYCN* amplification, low stage of disease or age < 18 months.^{14,15} Moertel et al used autoradiography with radiolabeled somatostatin analogues to determine SSTR2 expression in 30 NB specimens and found that expression was associated with lower risk, non-*MYCN*-amplified disease and favorable outcome.¹⁵ In addition, Raggi et al used RT-PCR to detect SSTR2 mRNA in human cell lines and tumors from 54 patients, and determined SSTR2 expression to be positively

correlated with OS and EFS, negatively related to tumor stage and *MYCN* amplification.¹⁹ One of the limitations of our study is that as we only had archival specimens available, we did not compare SSTR2 expression using techniques used in other case series other than IHC.

There is no standardized way to clinically interpret the degree of SSTR2 expression of NB tumors in terms of uptake on DOTA-conjugate peptide scans, or to identify which relapsed and progressive patients with NB would benefit from DOTA-conjugate peptide therapy. In our preclinical studies, we found that xenografts expressing high levels of SSTR2 as determined by Western blot and RT-PCR showed a higher uptake of ⁶⁸Ga-DOTA-TATE PET/CT.³¹ We also demonstrated co-localization of SSTR2 and DOTATATE uptake using autoradiography-IHC. Early studies in relapsed and refractory NB have used ⁶⁸Ga-DOTATATE PET/CT to screen for patients for DOTA-conjugate peptide therapy, using a threshold of degree of uptake greater than the liver.^{29,30} In addition, Kong et al demonstrated positive staining for SSTR2 by IHC in 5/5 eligible patients with tissue samples.²⁹ They noted some patients had weak SSTR2 expression on biopsy with high ⁶⁸Ga-DOTATATE uptake, possibly due to tumor heterogeneity and sampling.²⁹ It will be important to correlate quantitative SSTR2 immunostaining on tissue samples in a prospective study with ⁶⁸Ga-DOTATATE uptake in larger cohorts of patients to determine optimal cut-offs or thresholds for eligibility for treatments.

The clinical significance of the finding in this study that SSTR2 was more highly expressed in non-high risk NB patients suggests that radiolabeled DOTA-conjugate peptide imaging could be helpful in non-high risk group to detect extent of disease at diagnosis. In clinical studies, sensitivity of ⁶⁸Ga-DOTA-TOC PET/CT on a per-lesion basis is higher than ¹³¹I-MIBG (94.4% vs 76.9%) including detecting marrow disease, and is also a quicker scan that can be done within an hour of injection rather than next day for MIBG.^{43,44} However, further studies would be needed to understand if imaging is detecting more differentiated and less clinically significant disease, and the risk-benefit ratio of DOTA-conjugate peptide scanning in non-high-risk patients.

Focusing on the cohort of 18 high-risk and 8 non-high-risk patients with relapsed and progressive NB, who would be the potential candidates for targeted ¹⁷⁷Lu-DOTATATE therapy, we found similar SSTR2 expression compared to the total cohort (median 1.33 (0.26-2.55) vs 1.31 (0.26-2.55)). (Table 2) For a small group of patients (N=15) who had pre and post-treatment samples analysed, there was no significant difference in SSTR2 expression. Of note, it was beyond the scope of the study to examine tumor biopsy samples at the time of relapse for SSTR2 expression. Studies have demonstrated that SSTR2 can be detected at relapse, and also that the SSTR status may change or SSTR not be expressed at time of progression or after chemotherapy.^{20,21}

Within the cohort we were also interested in whether DOTA-conjugate imaging and treatment could be used as an alternative for MIBG-non-avid patients. However, we did not find a statistically significant difference between SSTR2 scores for MIBG avid versus non-avid NB, and patients who were MIBG-avid had higher median SSTR2 scores than non-avid patients. As expected, most of the patients in the MIBG-non-avid group were non-high-risk

patients, so one could have expected the SSTR2 expression to be higher in this group. In addition, all the patients in the relapsed/refractory group who had high SSTR2 tumor expression were MIBG-avid. NET expression is correlated to MIBG avidity but despite our preclinical work finding low NET expression in high SSTR2 NB cell lines, using IHC and western blot detection methods we did not find any correlation between NET expression and SSTR2 expression in our cohort. (Figure 2)³³ Based on these findings of SSTR2 expression in MIBG-avid tumors, it would still be feasible to use DOTA-conjugate imaging and treatment in patients who have not responded to ¹²³I-MIBG.

From this study, we can conclude that SSTR2 expression is present in many NB tumors at time of diagnosis, particularly non-high-risk patients, but expression is variable and does not correlate with prognosis or other clinical and diagnostic features. Further studies will be required to evaluate the prevalence of SSTR2 in relapsed patients with NB and to determine the optimal threshold of immunostaining that correlates with clinical DOTA uptake. The role of DOTA-conjugate peptide imaging in the non-high-risk population still needs to be determined, but this technique can also be helpful in detecting metastatic disease. Moreover, DOTA-conjugate peptide therapy has so far demonstrated minimal toxicity, some efficacy and increased convenience.^{29,30,45} and has potential as a personalized approach to relapsed or refractory NB with demonstrated SSTR2 expression, particularly in heavily pretreated patients who have not responded to other therapies. A necessary step, however, will involve establishing standardized IHC thresholds for SSTR2 positivity in conjunction with a standard scoring system for DOTA-conjugate peptide imaging, in order to select eligible patients for DOTA-conjugate peptide therapy.

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Abbreviation table

COG	Children's Oncology Group
CT	Computerized tomography
DOTATATE	(DOTA ⁰ -Tyr ³)-octreotate
DOTATOC	(DOTA ⁰ -Phe ¹ -Tyr ³)octreotide
EFS	Event-free survival

⁶⁸Ga	Gallium-68
HR	High-risk
IHC	Immunohistochemistry
INPC	International Neuroblastoma Pathology Classification
IRS	Immunoreactive score
INSS	International Neuroblastoma Staging System
¹⁷⁷Lu	Lutetium-177
MIBG	Metaiodobenzylguanidine
NB	Neuroblastoma
NET	Norepinephrine Transporter
NHR	Non-high risk
OS	Overall survival
PET	Positron emission tomography (PET)
RT-PCR	Reverse Transcriptase – Polymerase Chain Reaction
SSTR2	Somatostatin receptor 2

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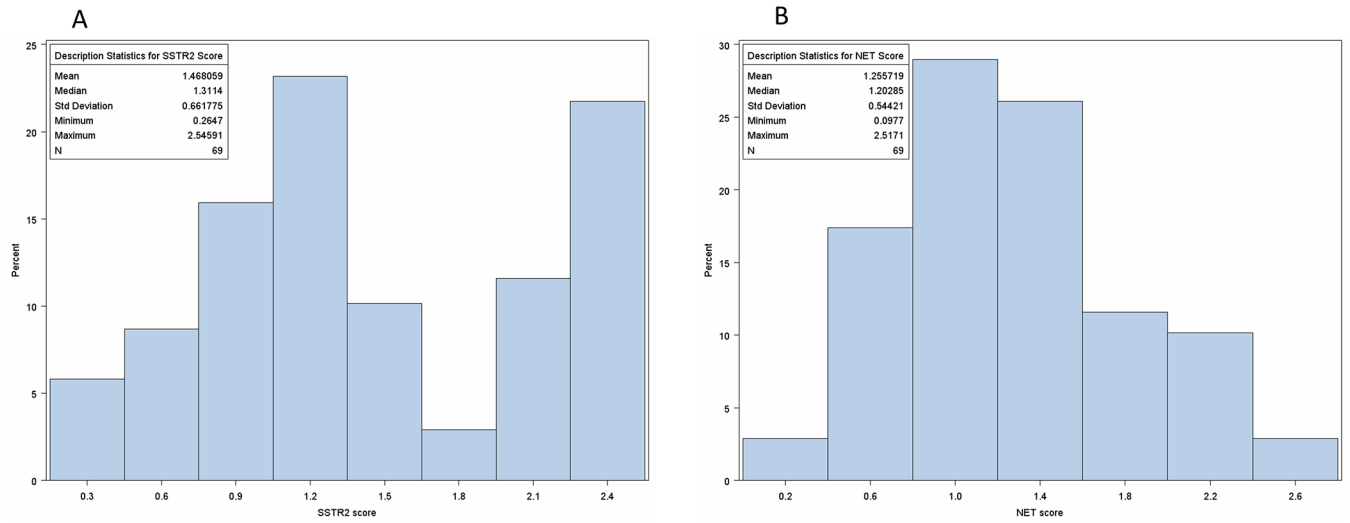


Figure 1: Distribution of Somatostatin Receptor-2 (SSTR2) (**1A**) and Norepinephrine Transporter (NET) Receptor (**1B**) immunohistochemistry scores from 69 high-risk (N=36) and non-high risk tumors (N=33) neuroblastoma tumors from initial diagnostic biopsy. SSTR-2 expression median score of 1.31 (range: 0.26-2.55) and NET expression median score of 1.20 (range: 0.10-2.52).

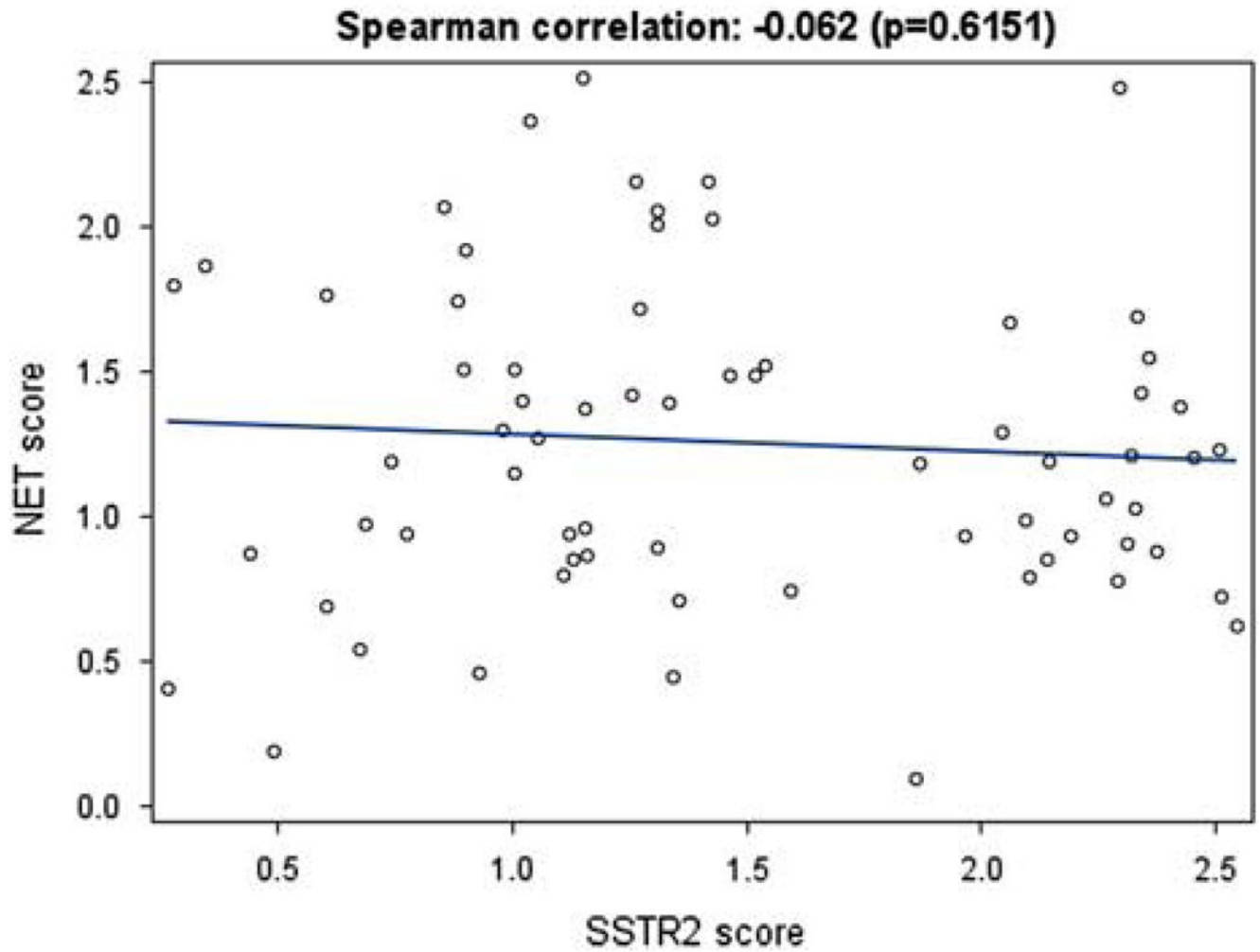


Figure 2: Scatter plot demonstrating the correlation of Somatostatin Receptor-2 (SSTR2) and Norepinephrine Transporter (NET) immunohistochemistry scores using digital analysis from neuroblastoma tumors (N=69). No statistical correlation was demonstrated between SSTR2 scores (X-axis) and NET scores (Y-axis) ($r = -0.062$, $p = 0.62$).

Table 1:

Comparison of clinical features and Somatostatin Receptor-2 (SSTR2) immunohistochemistry scores of mixed-risk neuroblastoma cohort (N=69).

Descriptive statistics for SSTR2 scores included are median, mean and standard deviation. P-value refers to Wilcoxon rank-sum tests for difference between clinical features. COG risk group (low/intermediate vs high) was the only variable with statistically significant SSTR2 scores;

Clinical Feature		N	%	Median	Mean	SD	P-value
Age	< 18 months	37	53.6%	1.4174	1.4958	0.6795	0.6346
	18 months	32	46.4%	1.2054	1.4359	0.65	
INSS Stage	Non-stage 4	32	46.4%	1.3339	1.4807	0.6718	0.8804
	Stage 4	37	53.6%	1.3083	1.4571	0.662	
MYCN status	Not amplified	55	79.7%	1.4174	1.5517	0.6857	0.0992
	Amplified	14	20.3%	1.2325	1.1394	0.4377	
INPC Histology	Favorable	32	46.3%	1.4469	1.5752	0.6576	0.1669
	Unfavorable	36	52.2%	1.2054	1.3429	0.6388	
MIBG Avidity	Non-avid	16	23.2%	0.9544	1.2424	0.6473	0.0767
	Avid	53	76.8%	1.4174	1.5362	0.6568	
COG Risk Group	Low/Intermediate	33	47.8%	1.5409	1.6519	0.6611	0.0320 *
	High	36	52.2%	1.1563	1.2996	0.6247	

* refers to statistically significant p-value.

INSS: International Neuroblastoma Staging System, INPC: International Neuroblastoma Pathology Classification, MIBG: Meta-iodobenzylguanidine, SD: standard deviation, COG: Children's Oncology Group.

Table 2.

Clinical and biological characteristics of the 26 patients who relapsed or progressed with neuroblastoma.

Clinical Feature		N	%			
Event type	Relapse	20	76.9			
	Progression	6	23.1			
Risk group	Low/intermediate	8	30.8			
	High	18	69.2			
MYCN status	Non-amplified	21	80.8			
	Amplified	5	19.2			
MIBG avidity	Non-avid	3	11.5			
	Avid	23	88.5			
	Median	Range	Mean	SD	N	
SSTR2	1.33	0.26-2.55	1.47	0.73	26	
NET	1.17	0.40-1.80	1.13	0.40	26	

MIBG: Meta-iodobenzylguanidine, NET: norepinephrine transporter, SSTR2: Somatostatin Receptor-2.