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Article

Effect of Neoadjuvant Chemotherapy on Tumor Infiltrating Lymphocytes in Resectable Gastric Cancer: Analysis from a Western Academic Center

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Simple Summary: In this investigation, we analyzed the number, type and location of immune cells within surgically resected gastric cancer specimens treated with or without preoperative chemotherapy. We hypothesized that chemotherapy can stimulate the host immune system evidenced by an increased number of anti-tumor infiltrating lymphocytes in the tumor microenvironment. We found significantly elevated levels of immune cells within chemotherapy-treated tumors compared to chemotherapy-naïve specimens. We also revealed important associations between survival and immune lymphocytes in the tumor related stromal tissue. Together, we add evidence supporting the immunostimulatory role of chemotherapy and underscore the potential utility of immunotherapy in resectable gastric cancer.

Abstract: Tumor infiltrating lymphocytes (TILs) are an emerging biomarker predictive of response to immunotherapy across a spectrum of solid organ malignancies. The characterization of TILs in gastric cancer (GC) treated with contemporary, multiagent neoadjuvant chemotherapy (NAC) is understudied. In this retrospective investigation, we analyzed the degree of infiltration, phenotype, and spatial distribution of TILs via immunohistochemistry within resected GC specimens treated with or without NAC at a Western center. We hypothesized that NAC executes immunostimulatory roles evidenced by an increased number of anti-tumor TILs in the tumor microenvironment. We found significantly elevated levels of conventional and memory CD8+ T cells, and total TILs (CD4+, CD8+, T_{reg}, B cells) within chemotherapy-treated tumors compared to chemotherapy-naïve specimens. We also revealed important associations between survival and pathologic response with enhanced TIL infiltration. Taken together, our findings advocate for an immunostimulatory role of chemotherapy and underscore the potential synergistic effect of combining chemotherapy with immunotherapy in resectable gastric cancer.

Keywords: gastric cancer; tumor infiltrating lymphocytes; neoadjuvant chemotherapy; immunotherapy

Introduction

Gastric cancer (GC) is among the most common and aggressive gastrointestinal (GI) cancers worldwide.^{1,2} While accounting for only 1.5% of new cancer diagnoses in the United States, nearly half of patients present with advanced disease.³ Although perioperative chemotherapy regimens

have evolved and garnered modest improvements in OS when administered in the neoadjuvant setting, 5-year survival in advanced GC remains less than 40%.⁴ Thus, the need for improved anti-tumor therapies for gastric cancer is paramount.

Immunotherapy, specifically immune checkpoint blockade (ICB), has revolutionized the care of several solid organ malignancies such as cutaneous melanoma, non-small cell lung cancer (NSCLC), and renal cell carcinoma.⁵⁻⁸ Recent randomized control trial (RCT) data has established adjuvant ICB therapy in resected stage II/III esophageal cancer as standard of care in light of significantly prolonged disease-free survival with immunotherapy.⁹ While consensus guidelines currently recommend ICB immunotherapy in unresectable or metastatic GC that harbor established biomarkers predictive of response to immunotherapy, its utility for potentially resectable GC tumors warrants further investigation.¹⁰

It is well known that the degree and phenotype of tumor infiltrating lymphocytes (TILs) is a prognostic marker for response to ICB.¹¹⁻¹⁴ In triple negative breast cancer and NSCLC, higher cytotoxic T cells (CD8+ T cells) demonstrate higher rates of overall response to ICBs along with improved progression-free and overall survival (PFS, OS, respectively) compared to those with lower CD8+ T cells.^{15,16} It is also known that conventional chemotherapeutic agents such as anthracyclines and platinum-based agents, the latter of which are frequently used to treat GC, can favorably alter the tumor microenvironment (TME) by inducing immunogenicity and synergizing the anti-tumor effect of host immunostimulatory agents.^{17,18}

The effect of contemporary multiagent chemotherapy on the degree of infiltration, phenotypes, and spatial distribution of TILs in potentially resectable GC is not well defined. The current body of work lacks analysis of memory immune cell subtypes and consideration of spatial (intratumoral versus stromal) TIL distributions.^{17,19,20} Furthermore, most studies that do report on GC are from Asia, which is known to have distinct disease biology treated with different chemotherapeutic regimens than in the West.²¹ Considering these differences, we sought to characterize the density and infiltrative patterns of conventional and memory TIL subtypes of GC treated with or without chemotherapy at a Western academic referral center. We hypothesized that chemotherapy favorably alters the TME of GC leading to increased levels of anti-tumor TILs.

Methods

Patient Cohort

After obtaining institutional review board consent, all adult patients with biopsy proven diagnosis of gastric adenocarcinoma who ultimately underwent a resection with curative intent from 2012-2020, either endoscopic or surgical, at our institution and had available formalin fixed, paraffin embedded (FFPE) tissue samples for histologic analysis were included in this study. A retrospective review of a prospectively maintained, clinically oriented database of patients was conducted. Review of the patient electronic health record was performed for missing data. After patients were identified, additional FFPE slides were requested from areas of invasive tumor at least 2 mm in diameter. Slides were reviewed during creation, and the areas of invasive disease were determined by a board-certified gastrointestinal pathologist (author JK).

Definitions and Immunologic Profile Characterization

TIL populations were characterized by multiplex immunohistochemistry (IHC) staining of associated cell surface (CD, cluster of differentiation) or intranuclear markers using the Vectra-7-tumor infiltrating lymphocyte kit (PerkinElmer, Waltham, MA). The included TILs and markers are as follows: B cells/CD220+, CD8 T cells/CD8+, CD4 T cells/CD4+, T regulatory (T_{reg}) cells/forkhead box P3 (FOXP3)+, CD8 memory T cells/CD8+/CD45RO+, CD4 memory T cells/CD4+/CD45RO+, memory Treg cells/CD4+/FOXP3+/CD45RO+, memory B cells/CD220+/CD45RO+, epithelial malignant cell/pan cytokeratin. TIL density was defined as the number of above stained immune cells per mm² designated within tumor or stroma regions of the tissue section. Total TILs were defined as the sum of CD4+, CD8+, T_{reg}, and B cells. Categorical assignment of high and low TIL density was

determined by the median value from the overall cohort. Clinical and pathologic staging of GC tumors were based on the latest National Comprehensive Cancer Network (NCCN) guidelines.¹⁰

To characterize the immunologic profile of GC tumors, we also analyzed EBV status, mismatch repair (MMR) protein expression, and tumor cell PD-L1. EBV status was determined by in situ hybridization (ISH) detection of EBV-encoded small RNA (EBER)-positive tumor cells (ARUP Laboratories, Salt Lake City, UT). Assessment of mismatch repair (MMR) protein expression was performed via immunohistochemistry (IHC) analysis of *MLH1*, *PSM2*, *MSH2*, and *MSH6* proteins (Leica; Wetzlar, Germany); deficiency (dMMR) was defined as loss of >95% of any one of the protein expressions in tumor cells. Programmed Death-Ligand 1 (PD-L1) expression was measured via the Combined Positivity Score (CPS), defined as the number of positive PD-L1 stained cells via IHC divided by the total number of tumor cells multiplied by 100; values greater than 1 were considered positive expression (Leica; Wetzlar, Germany).

Multiplex Immunohistochemistry

IHC was performed using an autostainer and then slides reviewed using image processing software following a previously employed protocol²²: Vectra 3.0 Automated Quantitative Pathology Imaging System (PerkinElmer) was used with the Bond RX autostainer (Leica). Slides were deparaffinized, heat treated in epitope retrieval solution 2 (ER2) antigen retrieval buffer for 20 minutes at 93 C (Leica), blocked in antibody (Ab) Diluent (PerkinElmer), incubated for 30 min with the primary antibody, 10 minutes with horseradish peroxidase-conjugated secondary polymer (anti-mouse/anti-rabbit, Perkin Elmer), and 10 minutes with horseradish peroxidase-reactive OPAL fluorescent reagents (Perkin Elmer). Slides were washed between staining steps with Bond Wash (Leica) and stripped between each round of staining with heat treatment in antigen retrieval buffer. After the final staining round, the slides were heat-treated in antigen retrieval buffer, stained with spectral 4',6-diamidino-2-phenylindole (PerkinElmer), and cover slipped with Prolong Diamond mounting media (ThermoFisher; Waltham, MA). Whole slide scans were collected using the 10× objective at a resolution of 1.0 μm. Then 10 regions of interest identified by a gastrointestinal subspecialty trained board-certified pathologist (author JK) were scanned for multispectral imaging with the 20× objective at a resolution of 0.5 μm. The multispectral images were analyzed with inForm software (PerkinElmer) to unmix adjacent fluorochromes; subtract autofluorescence; segment the tissue into tumor regions and stroma; segment the cells into nuclear, cytoplasmic, and membrane compartments; and to phenotype the cells according to cell marker expression.

Statistical Analysis

Parametric and nonparametric data are presented as means with standard deviations and medians with interquartile range, respectively. Categorical variables are expressed as absolute and relative frequencies (count and number). Categorical variables were compared using Chi-squared test; for continuous variables, parametric data was analyzed via Student's T test and non-parametric data with Mann-Whitney U test. Comparison of more than two groups of non-parametric data was performed via Wilcoxon sign-ranked test. Kaplan-Meier survival curves were generated to estimate time-to-event analyses for OS and RFS. All statistical analyses were performed in IBM SPSS version 28.0 (IBM, Chicago, IL, USA). Figures were constructed with SPSS or GraphPad Prism (version 10.0.0 for Windows, GraphPad Software, Boston, Massachusetts USA). Quantification of IHC staining of MMR, PD-L1, and TIL densities was completed with inForm Imaging Analysis Software (Akoya Biosciences, Marlborough, MA, USA). Statistical significance was considered $p \leq 0.05$.

Results

Patient Cohort

Demographic and clinicopathologic variables of the entire patient cohort and stratified by receipt of NAC are displayed in **Table 1**. Eighty patients were identified, 68 of which pathologic specimens suitable for histologic analysis. Most patients were male (59%, $n=40$), of Caucasian race

(67%, n=46), with a mean age of 63 years at the time of diagnosis (range 28-87 yrs, SD +/-15 yrs). All tumors were adenocarcinoma in origin. In the total cohort, most patients harbored clinical stage T3 tumors (52%, n=35) and node negative disease (N0 57%, n=39). Nearly 75% of patients received NAC (n=50), the most common regimen being combination folinic acid, fluorouracil, and oxaliplatin (FOLFOX, 38%, n=26). Neoadjuvant radiation was given to four percent of patients (n=3). Surgical resection consisted of total gastrectomy or subtotal gastrectomy in 93% of patients (n=63), the remaining 7% underwent endoscopic resection (n=5). Half of the cohort received adjuvant chemotherapy (52%, n=35).

Table 1. Demographic and clinicopathologic characteristics of the overall cohort and of patients with $\geq cT2N0-3$ disease stratified by receipt of neoadjuvant chemotherapy (NAC). *EMR*, endoscopic mucosal dissection; *ESD*, endoscopic submucosal dissection; *CRS*, chemotherapy response score; *US*, upfront surgery.

<i>Characteristic</i>	<i>Overall cohort</i> (n=68)	<i>Upfront Surgery</i> (n=18)	<i>NAC</i> (n=50)	<i>p value</i>	<i>$\geq cT2N0-3$ US</i> (n=11)	<i>$\geq cT2N0-3$ NAC</i> (n=46)	<i>p value</i>
Demographic characteristics							
Sex, n (%)							
Male	40 (58.8)	8 (44.4)	32	0.148	7 (63.6)	30 (65.2)	0.921
Female	28 (41.2)	10 (55.6)	(64.0)		4 (36.4)	16 (34.8)	
			18 (36.0)				
Age at diagnosis, mean (SD)	62.8 (53.3, 73.3)	64.6 (+/- 18.0)	65.5 (+/- 13.5)	0.531	69.8 (+/- 16.6)	63.0 (+/- 13.4)	0.156
Race, n (%)							
White	46 (67.6)	13 (72.2)	33	0.944	7 (63.6)	31 (67.4)	0.944
Black/African American	6 (8.8)	1 (5.6)	(66.0)		1 (9.1)	4 (8.7)	
Asian	8 (11.8)	2 (11.1)	5 (10.0)		1 (9.1)	6 (13.0)	
American Indian/Alaskan native	1 (1.5)	-	6 (12.0)		-	-	
Other	7 (10.3)	2 (11.1)	1 (2.0)		2 (18.2)	5 (10.9)	
			5 (10.0)				
Clinicopathologic characteristics							
Clinical T stage, n (%)							
T1a	3 (4.4)	2 (11.1)	1 (2.0)	<0.001	-	-	<0.001
T1b	7(10.3)	5 (27.8)	2 (4.0)		-	-	
T2	12 (17.6)	7 (38.9)	5 (10.0)		7 (63.6)	5 (10.9)	
T3	35 (51.5)	3 (16.7)	32		3 (75.0)	32 (69.6)	
T4	1 (1.5)	-	(64.0)		-	1 (2.2)	
T4a	7 (10.3)	-	1 (2.0)		-	7 (15.2)	
T4b	2 (2.9)	1 (5.6)	7 (14.0)		1 (9.1)	1 (2.2)	
	1 (1.5)	-	1 (2.0)		-	-	

Missing			1 (2.0)					
Clinical N stage, <i>n</i>								
(%)		39 (57.4)	16 (88.9)	23	0.002	9 (81.8)	20 (43.5)	0.026
N0		28 (41.2)	2 (11.1)	(46.9)		2 (18.2)	25 (54.3)	
N1-2		1 (1.5)	-	26		-	1 (2.2)	
Missing				(52.0)				
			1 (2.1)					
Overall clinical								
stage, <i>n</i> (%)		19 (27.9)	14 (77.8)	5 (10.2)	<0.001	7 (63.6)	3 (6.5)	<0.001
Stage I		21 (30.9)	2 (11.1)	19		2 (18.2)	18 (39.1)	
Stage II		25 (36.8)	1 (5.6)	(38.0)		1 (9.1)	24 (52.2)	
Stage III		67 (98.5)	1 (5.6)	24		1 (9.1)	1 (2.2)	
Stage Iva		1 (1.5)		(48.0)		-	-	
Missing				1 (2.0)				
			1 (2.0)					
Tumor location, <i>n</i>								
(%)		45 (66.2)	15 (75.0)	30	0.091	10 (90.9)	26 (56.5)	0.034
Distal		18 (26.5)	2 (10.0)	(60.0)		-	16 (34.8)	
Proximal		4 (5.9)	-	16		-	4 (8.7)	
Linitis plastica		1 (1.5)	1 (5.0)	(32.0)		1 (9.1)	-	
Undefined				4 (8.0)				
			-					
Histologic subtype, <i>n</i> (%)								
		24 (35.3)	9 (50.0)	15	0.277	7 (63.6)	15 (32.6)	0.275
Intestinal		39 (57.4)	8 (44.4)	(30.0)		4 (36.4)	28 (60.9)	
Diffuse/Signet-		3 (4.4)	-	31		-	2 (4.3)	
ring		2 (2.9)	1 (5.6)	(62.0)		-	1 (2.2)	
Mixed				3 (6.0)				
Neuroendocrine				1 (2.0)				
<i>Preoperative and intraoperative characteristics</i>								
Neoadjuvant								
regimen, <i>n</i> (%)		17 (25.0)	-	-	-	-	17 (37.0)	-
Other		25 (36.8)	-	-		-	23 (50.0)	
FOLFOX		6 (8.8)	-	-		-	6 (13.0)	
FLOT								
Rounds	of	3.89	(+/-	-	-	-	4.0	(3.0-
chemotherapy		1.7)					4.0)	-
Neoadjuvant								
radiation, <i>n</i> (%)		65 (95.6)	-	45	0.288	11 (100)	43 (93.5)	0.288
No		3 (4.4)	-	(93.8)			3 (6.5)	
Yes				3 (6.3)				

Type of resection, <i>n</i>							
(%)	40 (58.8)	9 (50.0)	31	0.019	7 (63.6)	28 (60.9)	0.482
Partial gastrectomy	23 (33.8)	5 (27.8)	(62.0)		3 (27.3)	17 (37.0)	
Total gastrectomy	5 (7.3)	4 (20.0)	18		1 (9.1)	1 (2.2)	
EMR/ESD*			(36.0)				
			1 (2.0)				
Pathologic tumor characteristics							
Pathologic overall							
stage, <i>n</i> (%)	21 (30.9)	11 (61.1)	10	0.006	4 (36.4)	9 (19.6)	0.270
Stage I	20 (29.4)	1 (5.6)	(20.0)		1 (9.1)	18 (39.1)	
Stage II	20 (29.4)	5 (27.8)	19		5 (45.5)	15 (32.6)	
Stage III	7 (10.3)	1 (5.6)	(38.0)		1 (9.1)	4 (8.7)	
Stage IV			15				
			(30.0)				
			6 (12.0)				
Clinical to							
pathologic stage							0.201
change, <i>n</i> (%)	35 (51.5)	13 (72.2)	22	0.058	6 (54.5)	21 (45.7)	
No change	17 (25.0)	1 (5.6)	(44.0)		1 (9.1)	16 (34.8)	
Downstage	15 (22.1)	4 (22.2)	16		4 (36.4)	9 (19.6)	
Upstage	1 (1.5)	-	(32.0)		-	-	
Missing			11				
			(22.0)				
			1 (2.0)				
Histologic subtype,							
<i>n</i> (%)	24 (35.3)	9 (45.0)	15	0.443	7 (63.6)	15 (32.6)	0.275
Intestinal	39 (57.4)	10 (50.0)	(31.3)		4 (36.4)	28 (60.9)	
Diffuse/signet-ring	3 (4.4)		29		-	2 (4.3)	
Mixed	2 (2.9)	1 (5.0)	(60.4)		-	1 (2.2)	
Neuroendocrine			3 (6.3)				
			1 (2.1)				
Histologic							
differentiation, <i>n</i> (%)	42 (61.8)	7 (38.9)	35	0.075	3 (27.3)	31 (67.4)	0.047
Poor	5 (7.4)	2 (11.1)	(70.0)		2 (18.2)	3 (6.5)	
Poor-moderate	17 (25.0)	6 (33.3)	3 (6.0)		5 (45.5)	11 (23.9)	
Moderate	1 (1.5)	1 (5.6)	11		1 (9.1)	-	
Mod to well	3 (4.4)	2 (11.1)	(22.0)		-	1 (2.2)	
Well			-				
			1 (2.0)				
Margin status, <i>n</i> (%)							
R0	58 (85.3)	18 (100)		0.040	11 (100)	37 (80.4)	0.040

R1	10 (14.7)	-	40	-	9 (19.6)
R2	-	-	(80.0)	-	-
			10		
			(20.0)		
			-		
Treatment effect, <i>n</i>					
(%)	4 (8.0)	-	4 (8.0)	-	4 (8.7)
Minimal residual	21 (42.0)		21		20 (43.5)
disease (CRS 3)	22 (44.0)		(42.0)		19 (41.3)
Moderate response	3 (6.0)		22		3 (6.5)
(CRS 2)			(44.0)		
Poor response			3 (6.0)		
(CRS 1)					
Unknown					

Demographic and Clinicopathologic Characteristics of Upfront Surgery & NAC Cohorts

Patients who received NAC were significantly more likely to have clinically larger tumors and node positive disease resulting in higher overall clinical stage (**Table 1**). Of the overall study cohort, 84% of patients (*n*=57) met the current NCCN recommendations to receive preoperative chemotherapy (\geq T2N0-3); of these patients, 19% did not receive NAT (*n*=11) most commonly due to patient preference (55%, *n*=6) in the setting of cT2N0 disease. In the \geq cT2N0-3 cohort, those who received NAC were more likely to have positive node disease, proximal tumor location, and poor histologic grade.

TIL and Molecular Profiles of Upfront Surgery & NAC Cohorts

The intratumoral and stromal TIL phenotypes/densities and molecular profiles of the study cohort are detailed in Table 2, Figure 1. In the overall cohort, patients who received NAC had significantly higher intratumoral conventional CD8+ T cells (14.3 vs. 5.1, *p*=0.024) and total TILs (summation of CD4+, CD8+, T_{reg}, B cells; 19.3 vs. 7.9; *p*=0.047). There were no significant differences in TIL densities in the tumor stroma between the two groups. The prevalence of EBV positive, dMMR, and PD-L1 positive status was not different between the upfront surgery and NAC groups.

Table 2. Molecular phenotype and tumor infiltrating lymphocyte densities in the overall cohort (left) and in patients with \geq cT2N0-3 disease (right) stratified by location (intratumoral and stromal) and receipt of NAC. *US*, upfront surgery.

Molecular phenotype & tumor infiltrating lymphocyte profiles							
	Overall cohort (<i>n</i> =68)	Upfront Surgery (<i>n</i> =18)	NAC (<i>n</i> =50)	<i>p</i> value	\geq cT2N0-3 US (<i>n</i> =11)	\geq cT2N0-3 NAC (<i>n</i> =46)	<i>p</i> value
EBV status, <i>n</i> (%)							
Negative	65 (95.6)	18 (100)	47 (94.0)	0.288	11 (100)	43 (93.5)	0.384
Positive	3 (4.4)	-	3 (6.0)		-	3 (6.5)	
MMR, <i>n</i> (%)							
Proficient	60 (88.2)	17 (94.4)	43 (86.0)	0.340	10 (90.9)	40 (87.0)	0.720
Deficient	8 (11.8)	1 (5.6)	7 (14.0)		1 (9.1)	6 (13.0)	
PD-L1 status, <i>n</i> (%)							
Negative	41 (60.3)	11 (61.1)	30 (60.0)	0.934	7 (63.6)	27 (58.7)	0.764
Positive	27 (39.7)	7 (38.9)	20 (40.0)		4 (36.4)	19 (41.3)	

Tumor infiltrating lymphocytes densities – intratumoral								
CD8+ T cells, cells/mm ²								
Conventional (CD8+)	8.6 (3.4, 37.1)	5.1 (2.1, 8.5)	14.25 (4.3, 43.7)	0.024	3.6 (2.0, 8.1)	43.7	0.019	
Memory (CD8+/CD45RO+)	1.8 (0.8, 9.0)	1.0 (0.6, 4.8)	2.3 (1.1, 10.8)	0.119	0.7 (0.4, 3.1)	2.0 (1.0, 11.6)	0.050	
CD4+ T cells, cells/mm ²								
Conventional (CD4+)	3.4 (0.8, 8.0)	1.7 (0.6, 5.7)	4.5 (0.8, 9.2)	0.182	1.5 (0.3, 3.2)	4.2 (0.9, 8.3)	0.089	
Memory (CD4+/CD45RO+)	1.6 (0.4, 5.0)	0.8 (0.3, 3.3)	2.1 (0.6, 5.6)	0.254	0.6 (0.2, 1.7)	1.6 (0.5, 5.4)	0.119	
T _{reg} cells, cells/mm ²								
Conventional (CD4+/FOXP3+)	0.4 (0.1, 1.7)	0.5 (0.04, 1.5)	0.4 (0.1, 1.9)	0.671	0.3 (0.1, 1.1)	0.4 (0.1, 1.9)	0.442	
Memory (CD4+/CD45RO+)	0.2 (0.04, 1.7)	0.2 (0.02, 1.2)	0.2 (0.04, 0.9)	0.950	0.1 (0.04, 0.4)	0.2 (0.03, 0.8)	0.754	
B cells, cells/mm ²								
Conventional (CD220+)	0.02 (0.003, 0.15)	0.01 (0.01, 0.09)	0.04 (0.0, 0.16)	0.550	0.01 (0.0, 0.01)	0.03 (0.0, 0.16)	0.088	
Memory (CD220+/CD45RO+)	0	0	0		0.0	0.0		
All TIL (CD8+, CD4+, B cell)	13.6 (5.5, 49.6)	7.9 (4.1, 15.4)	19.3 (5.6, 53.9)	0.047	6.7 (2.8, 9.6)	18.8 (5.4, 53.9)	0.041	
ALL memory TILs	0.3 (0.03, 2.3)	0.11 (0.01, 0.49)	0.28 (0.05, 3.48)	0.098	0.05 (0.01, 0.3)	0.2 (0.04, 3.1)	0.048	
CD8:Treg ratio	23.2 (6.6, 3.4)	7.8 (3.7, 53.3)	25.5 (12.4, 54.6)	0.123	7.5 (3.1, 29.5)	25.5 (7.0, 55.2)	0.079	
Tumor infiltrating lymphocytes densities – stromal								
CD8+ T cells, cells/mm ²								
Conventional (CD8+)	4.9 (1.6, 19.0)	6.4 (1.2, 21.4)	4.6 (1.8, 18.1)	0.597	2.3 (1.1, 8.4)	4.6 (1.9, 19.6)	0.203	
Memory (CD8+/CD45RO+)	3.3 (1.0, 11.6)	4.4 (0.9, 13.8)	3.1 (1.0, 8.8)	0.396	1.45 (0.6, 7.3)	3.1 (1.0, 10.2)	0.385	
CD4+ T cells, cells/mm ²								
Conventional (CD4+)	23.2 (6.3, 53.1)	23.6 (7.1, 73.7)	23.2 (6.0, 50.4)	0.906	8.9 (4.5, 21.2)	23.2 (7.0, 54.2)	0.143	
Memory (CD4+/CD45RO+)	9.7 (2.4, 29.0)	11.1 (3.0, 40.1)	8.4 (2.3, 21.0)	0.359	4.2 (1.8, 21.2)	8.4 (2.6, 22.7)	0.454	
T _{reg} cells, cells/mm ²								
Conventional (CD4+/FOXP3+)	1.3 (0.2, 3.4)	1.0 (0.5, 4.9)	1.4 (0.2, 2.8)	0.592	0.8 (0.2, 4.1)	1.4 (0.2, 3.3)	0.716	
Memory (CD4+/CD45RO+)	0.5 (0.1, 1.8)	0.6 (0.2, 1.7)	0.5 (0.1, 1.3)	0.254	0.5 (0.1, 1.8)	0.5 (0.1, 1.7)	0.952	
B cells, cells/mm ²								
Conventional (CD220+)	1.2 (0.4, 6.7)	1.3 (0.6, 10.0)	0.9 (0.3, 5.6)	0.294	0.9 (0.2, 1.3)	1.1 (0.3, 6.6)	0.379	
Memory (CD220+/CD45RO+)	0.1 (0.02, 1.2)	0.2 (0.02, 1.7)	0.1 (0.2, 0.6)	0.555	0.3 (0.02, 0.3)	0.1 (0.02, 0.9)	0.201	
All TIL (CD8+, CD4+, B cell)	36.1 (8.6, 74.9)	31.0 (8.7, 97.8)	37.4 (8.3, 71.1)	0.889	15.6 (5.3, 39.4)	37.4 (10.6, 73.3)	0.110	
CD8:Treg ratio	3.6 (2.3, 10.3)	3.4 (1.4, 16.2)	3.6 (2.4, 9.6)	0.479	2.3 (1.2, 4.8)	3.6 (2.3, 9.6)	0.152	

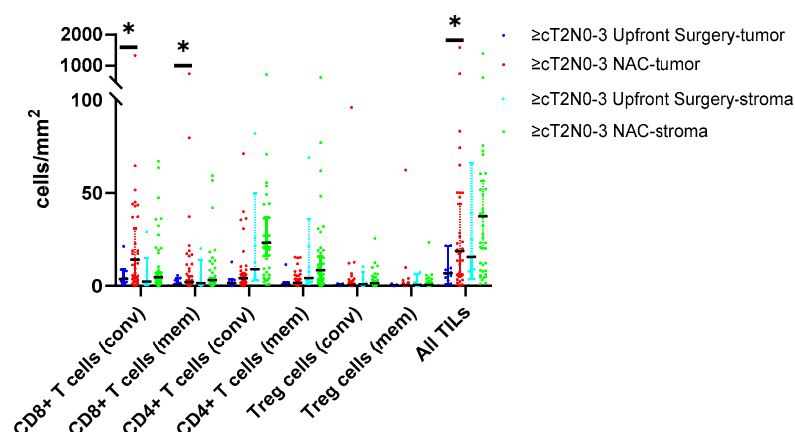
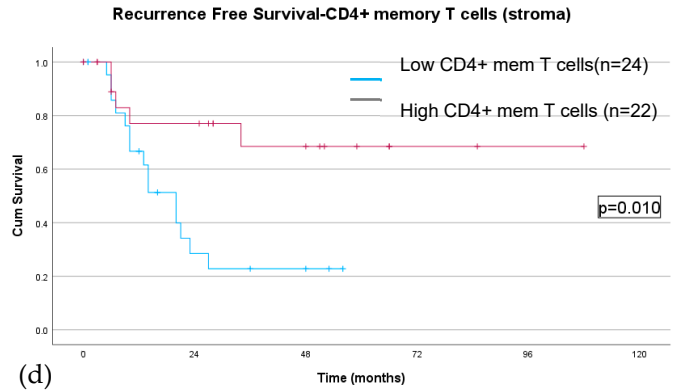
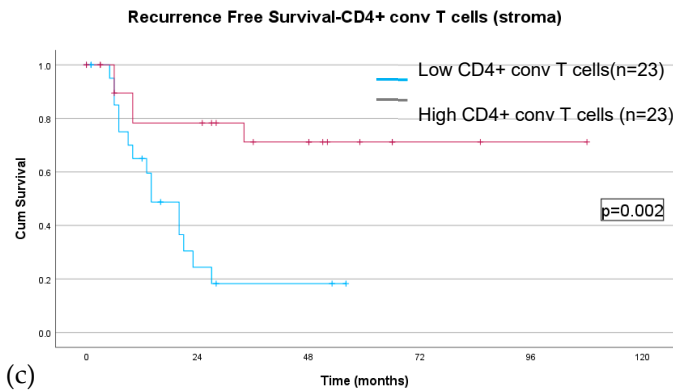
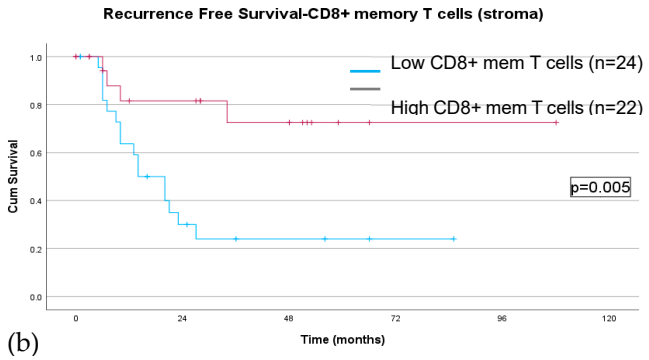
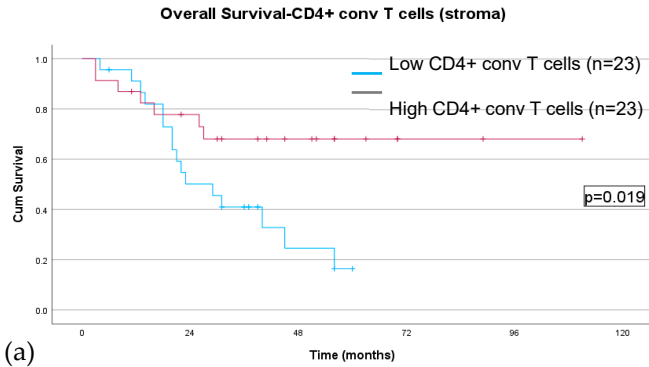


Figure 1. Scatter plot of tumor infiltrating lymphocyte (TIL) densities in \geq cT2N0-3 disease stratified by receipt of neoadjuvant chemotherapy (NAC) and tissue location (intratumoral vs. stromal). *Conv*, conventional; *mem*, memory; *Treg*, T regulatory. Black line represents median with error bars for 95% confidence interval. *Statistical significance $p < 0.05$.

In the subset of patients with \geq cT2N0-3 disease, conventional CD8+ T cells (14.2 vs. 3.6, and total conventional TILs (18.8 vs. 6.7, $p = 0.041$) continued to be significantly upregulated in the tumor tissue of those who underwent NAC. Additionally, in this select cohort, intratumoral memory CD8+ T cells (2.0 vs. 0.7, $p = 0.050$) and total memory TILs (0.2 vs. 0.05, $p = 0.048$) were increased in tumors treated with NAC. Again, no differences in TIL densities in the stromal component nor molecular phenotypes (EBV, MMR, PD-L1 positivity) was appreciated between the two cohorts. Although the CD8+ T cell to Treg ratio was substantially increased in tumor tissue of patients who received NAC, the difference only trended toward statistical significance (25.5 vs. 7.5, $p = 0.079$).

TIL density & Oncologic Outcomes

The median follow-up time in the overall cohort was 43 months (range 30-65 mos) with death occurring in nearly half the overall cohort (47.1%, $n = 32$) and distant recurrence in over a third of patients (36.8%, $n = 25$). Peritoneal dissemination was the most common form of metastasis (11/25, $n = 11$). In both the overall and \geq cT2N0-3 cohorts, various high (defined as upper half from median value) TIL populations in the stromal but not intratumorally were associated with significantly longer OS and RFS. Figure 2 and Supplemental Figures S1 and S2 display the statistically significant Kaplan-Meier curves stratified by TIL phenotype with associated log-rank analyses estimating median survival for OS and RFS.



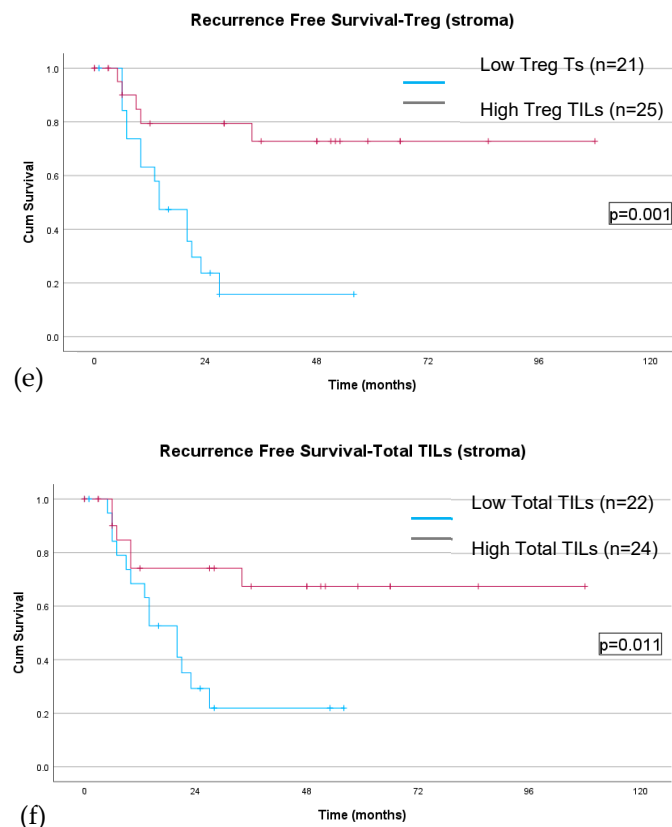


Figure 2. a-f. Kaplan-Meier survival curves with log-rank tests demonstrating longer (a) overall survival and (b-f) recurrence free survival with high vs. low **stromal** TILs in patients with \geq T2N0-3 disease treated with NAC. Median survival follows in parentheses. (a) low (29.0 mos) versus high (NR, not reached) CD4+ T cells; $p=0.019$ (b) low (14.0 mos) versus high (NR) CD8+ memory T cells $p=0.005$ (c) low (14.0 mos) versus high (NR) CD4+ T cells; $p=0.002$ (d) low (20.0 mos) versus high (NR) CD4+ memory T cells; $p=0.010$ (e) low (14.0 mos) versus high (NR) T regulatory cells; $p=0.001$ (f) low (20.0 mos) versus high (NR) total TILs; $p=0.011$.

TIL density & Pathologic Response

Most patients who underwent NAC demonstrated a poor pathologic response to preoperative treatment (chemotherapy response score 1) (Table 1). There were no differences in response by chemotherapy regimen. No significant associations were observed between high/low TIL categories and pathologic response based on the median cutoff values; however, we found that the top quartile of densities of intratumoral CD8+ T cells (OR 4.976; CI 1.166-21.242; $p=0.030$) and total TILs (OR 6.667; CI 1.269-35.035; $p=0.025$) were associated with higher rates of near complete and moderate response (chemotherapy response score 3 and 2, respectively) compared to poor response. Similarly, stromal CD8+ conventional T cells (OR 11.812; IC 1.3254-103.038; $p=0.025$), CD8+ memory T cells (OR 14.0; CI 1.615-121.369; $p=0.017$), total TILs (OR 5.625; CI 1.062-29.799; $p=0.042$), and total memory TILs (OR 14.0; CI 1.615-121.369; $p=0.017$) were more likely to be associated with improved pathologic response.

Discussion

In the present investigation, we compared TIL phenotypes and infiltrative patterns in resected GC specimens from patients who did and did not undergo NAC. We hypothesized that among our cohort of patients treated at a Western academic center, NAC-treated tumors would demonstrate higher TIL densities in the TME compared to non-NAC counterparts. We found that in both the overall cohort and among those recommended to receive NAC (\geq T2N0-3), tumors from NAC

recipients demonstrated significantly increased intratumoral, but not stromal, TILs compared to patients foregoing NAC. Furthermore, we observed improved OS, RFS, and pathologic response in patients with high compared to low TIL infiltration who received NAC.

Based on the results of recent RCTs, the application of immunotherapy in GC has been limited to unresectable or metastatic disease harboring specific immunotherapy-responsive molecular phenotypes e.g. PD-L1 positive, MSI-H, TMB-H.^{7,22,23} The results of such trials have raised the potential that ICB therapy could be beneficial for resectable GC. The only published report from a phase III RCT utilizing combined chemotherapy plus ICB versus chemotherapy plus placebo for locally advanced GC/gastroesophageal junction (GEJ) tumors did not show a statistical difference in event-free survival at a median follow-up of nearly fifty months but did demonstrate a significant improvement in pathologic complete response with combination chemotherapy and ICB.²⁴ Recently, the phase III CheckMate-577 trial in resected esophageal/GEJ tumors reported significantly longer disease-free survival in patients treated with adjuvant nivolumab compared to placebo.⁹ Notably, these improved outcomes occurred independently of PD-L1 status, a finding that highlights alternative prognostic biomarkers predictive of response to immunotherapy. One such biomarker may be the degree of anti-tumor TIL infiltration within the TME.^{13,25} Higher intratumoral and stromal TIL infiltrate, particularly cytotoxic CD8+ T cells, have been associated with longer survival and higher rates of pathologic response after ICB therapy compared to those with lower TIL infiltrate in advanced solid organ tumors.^{11,13,16,26} Therefore, identifying mechanisms to increase tumor-targeting TIL populations into the TME may facilitate immunotherapy in resectable GC.

The use of NAC has become standard of care for localized GC.^{27,28} Mounting evidence suggests that while conventional chemotherapeutic agents play various immunosuppressive roles, they may also induce substantial immunogenicity and immunostimulation against malignancy by producing tumor-derived neoantigens, improving cytotoxic T cell recognition of tumor cells, and upregulating damage associated molecular patterns (DAMPs) and cell surface molecules recruiting effector cells to the TME.^{17,29} However, the data demonstrating the impact of contemporary, multiagent chemotherapy on the degree and phenotypes of TILs in GC is lacking. Thus, we aimed to analyze the TIL composition in GC tumors treated with and without NAC.

We found that postoperative GC specimens treated with NAC demonstrated significantly increased densities of intratumoral TILs compared to those that did not undergo NAC. In the overall cohort which included patients with overall clinical stage I-III disease, CD8+ conventional T cells and total TILs were substantially elevated in NAC-exposed tumors. For those whom NAC is recommended per NCCN guidelines (clinical stage \geq T2N0-3), the upregulation of TILs was even more widespread as both conventional and memory subtypes of CD8+ T cells and total TILs were increased within the tumor tissue. Notably, we did not appreciate differences in stromal TIL densities between the two groups although both conventional CD4+ and CD8+ T cells were at least two-fold greater in NAC cohort. Our observations that anti-tumor TILs are increased after NAC is consistent with present literature in a range of epithelial carcinomas including breast, non-small cell lung cancer, colorectal and ovarian.³⁰⁻³⁴ Our findings also corroborate that of Yu *et al.*, who reported increased CD4+ and CD8+ T cell populations in Asian patients after receiving a combination of preoperative 5-FU, platinum-based agent, with or without taxane and gastrectomy.¹⁹ Unlike Xing *et al.* and Hu *et al.*, we did not appreciate a significant difference in intratumoral or stromal T_{reg} cells, which may be secondary to known differences in Western versus Asian gastric cancer biology and differences in NAC regimens.^{20,35}

Notably, to our knowledge, we are the first to report the relationship between increased memory T cell infiltration and receipt of NAC in GC. Memory subtypes are known to play important roles in executing durable anti-tumor response.³⁶ Furthermore, recent preclinical data suggests that neoantigen stimulation of CD4+ T cells can facilitate generation of specialized memory CD4+ T cells that be utilized in adoptive T cell immunotherapy to prime effector CD8+ T cells in mitigating metastasis.³⁷ Lastly, both clinical and preclinical studies have shown that response to ICB is positively related to the proportion of memory T cells, suggesting the importance of memory phenotypes to

mediating host immune response.^{38,39} Taken together, we show that memory T cell subtypes are higher in NAC-treated tumors, which may portend improved tumor control with IT.

In addition to enhanced TIL infiltration in NAC-exposed specimens, we identified associations between high TIL phenotypes and improved OS and RFS in patients who received NAC. Interestingly, despite observing statistically significant higher densities of intratumoral TILs between NAC and upfront surgery cohorts, survival associations were only related to high stromal rather than intratumoral TILs. These findings support existing literature citing similar associations with higher stromal TILs and improved RFS in breast and ovarian carcinomas.^{11,19,31,40} Further, stromal TILs, particularly CD8+ T cells, are proposed to be a stronger prognostic biomarker of the response to ICBs and survival than intratumoral TILs as reported by a meta-analysis including 2559 patients with a variety of solid organ tumors treated with immune checkpoint inhibitors.¹¹ Potential explanations for this finding may be that the intratumoral TILs, while increased, may be over-exposed to tumor rendering them to an inactive, “exhausted” phenotype.⁴¹ Additionally, active cytotoxic cells at the tumor periphery or invasive margin may be more proximal to antagonizing the aggressive metabolic and immune re-programming occurring at the tumor borders, thus critical to controlling tumor growth and dissemination.⁴² To this point, higher stromal TILs in the primary tumor site have been shown to correlate with decreased metastatic burden, which is consistent with our associations between improved RFS with increased stromal TIL populations.^{36,43} Notably, we also observed improved pathologic response to preoperative chemotherapy in a select subset of patients with the highest quartile of intratumoral and stromal infiltrating immune cells, supporting previous work demonstrating similar results in other NAC-treated carcinomas.^{44–46} Nevertheless, given that distant metastases are the primary mode of failure for gastric cancer, there are evidently a multitude of mechanisms driving tumor immune evasion and progression that may be independent of the TILs that are associated with the primary tumor.⁴⁷

While this study adds novel perspective to the immune landscape of resectable GC after NAC in Western patients, our results should be considered in the context of its limitations. As a retrospective, single center endeavor, it is constrained by inherent selection bias, small sample size, and heterogeneity in data collection/reporting. Additionally, we recognize our TIL and immunologic profile characterization is far from exhaustive, yet we aimed to bridge gaps according to prior literature. In our spatial TIL analysis, while we added novelty in differentiating intratumoral and stromal TILs, we did not assess TILs specifically confined to the tumor invasive margin, a metric that has risen to certain prognostic value. Due to the retrospective, clinically-oriented nature of this study, we are not able to fully explain the relationships between intratumoral and stromal TILs with long-term oncologic outcomes. Lastly, while a strength of this study is the in-depth nature of our analysis of TILs in the TME of Western GC, our results may not be fully translatable to GC at-large considering that GC arising in Asia is known to be biologically distinct. Given these limitations, future work should be dedicated to prospective, protocol-based analysis further detailing specific TILs such as granzyme B CD8+ T cells, effector and central memory T cells, natural killer cells, and those of the “exhausted” phenotype.

Conclusions

The immune TME of GC is highly heterogenous. Identifying mechanisms to facilitate novel therapeutics, i.e. immunotherapy, in effort to improve outcomes in GC is paramount. In this investigation, we observed that resected GC treated with NAC boast higher intratumoral TILs, namely conventional CD8+ and total TILs, compared to tumors undergoing upfront surgery across all clinical stages of localized disease. Importantly, we also established that memory subtypes are upregulated in a subset of higher stage patients who meet consensus criteria for NAC. Further, we highlight the prognostic value of stromal rather than intratumoral TILs for GC undergoing NAC. Together, our novel findings affirm the need for further investigation into the complex interplay between the TME, TILs, chemo- and immunotherapy.

Supplementary Materials: The following supporting information can be downloaded at the website of this paper posted on Preprints.org.

Author Contributions: Conceptualization, Elliott Yee, Jeffrey Kaplan, Sachin Wani, Sunnie Kim, Martin McCarter and Camille Stewart; Data curation, Elliott Yee, Danielle Gilbert, Sachin Wani and Martin McCarter; Formal analysis, Elliott Yee, Danielle Gilbert and Camille Stewart; Investigation, Sunnie Kim; Methodology, Elliott Yee, Jeffrey Kaplan, Sachin Wani, Sunnie Kim, Martin McCarter and Camille Stewart; Resources, Camille Stewart; Supervision, Camille Stewart; Validation, Jeffrey Kaplan; Writing – original draft, Elliott Yee; Writing – review & editing, Danielle Gilbert, Jeffrey Kaplan, Sunnie Kim, Martin McCarter and Camille Stewart.

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Ethical statement/consent: The University of Colorado Institutional Review Board (COMIRB) deemed this study exempt from IRB review given the retrospective nature of this study. Informed consent was given by patients to collect specimens as part of their routine medical care at the University of Colorado, Anschutz Medical Campus.

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