

Gastric carcinoma with lymphoid stroma in the era of the immune context and immunotherapies

I. Gullo^{1,2,3}, G. Gonçalves⁴, M. Athellogou⁵, M.L. Pinto^{3,6,7}, G.M. Almeida³, C. Oliveira^{2,3}, F. Carneiro^{1,2,3}

¹Department of Pathology, Centro Hospitalar de São João, Porto, Portugal; ²Faculty of Medicine of the University of Porto (FMUP), Porto, Portugal; ³Institute of Molecular Pathology and Immunology of the University of Porto (Ipatimup), Porto, Portugal and Instituto de Investigação e Inovação em Saúde, University of Porto, Porto, Portugal; ⁴Department of Biomedical Sciences and Medicine, University of Algarve, Faro, Portugal; ⁵Definiens AG, Bernhard-Wicki Str 5, 80636 Munich, Germany; ⁶INEB-Institute of Biomedical Engineering, University of Porto, Porto, Portugal; ⁷ICBAS-Institute of Biomedical Sciences Abel Salazar, University of Porto, Porto, Portugal.

Background and aims

Novel cancer immunotherapies can selectively block the cancer evasion of immune surveillance. Phase II/III clinical trials with antibodies directed against CTLA-4 and PD-1/PD-L1 immune inhibitory checkpoints are currently ongoing in GC and are new attractive therapeutic strategies for GC patients. Recently, The Cancer Genome Atlas (TCGA) network proposed a four-tiered molecular classification of GC [1], that could help to guide optimal selection of therapy. EBV+ and MSI- high GC are two molecular subtypes in which the PD-1/PD-L1 blockade may be particularly beneficial [1,2]. Moreover, a morphological subtype of GC has been associated to MSI-high status and EBV infection, that is Gastric Carcinoma with Lymphoid Stroma (GCLS) [3,4,5]. Hence, the abundant immune infiltrate and the molecular features of GCLS offer an attractive landscape to study tumour immune micro-environment, immune inhibitory checkpoints and their relationship with GC cells. The aim of this study was to analyse the clinico-pathological features, EBV infection, MSI, PD-L1 status and tumour immune microenvironment in GCLS.

Material and methods

Twenty-four GCLSs, selected from a series of 1088 surgically resected GC patients, were analysed by: RNA *in situ* hybridisation (EBER) for EBV, PCR/fragment analysis for MSI, and immunohistochemistry (IHC) for cytokeratin (CK) AE1/AE3, CD3, CD8 and PD-L1. PD-L1 immunoreactivity was evaluated separately for tumour immune and epithelial cells. The Immunoreactivity Scoring System (IRS) recently described by Böger *et al* was applied [6]. Double immunofluorescence for CK/PD-L1 and CD68/PD-L1 was performed in selected cases. CD3+, CD8+ T cell densities and CD8/CD3 ratio (CD8/CD3R) were calculated both in the tumour centre (TC) and at the invasive front (IF) by digital analysis (Definiens®). A tissue microarray (TMA) was constructed from a control group of 54 non-GCLSs and analysed by IHC for PD-L1 and by EBER.

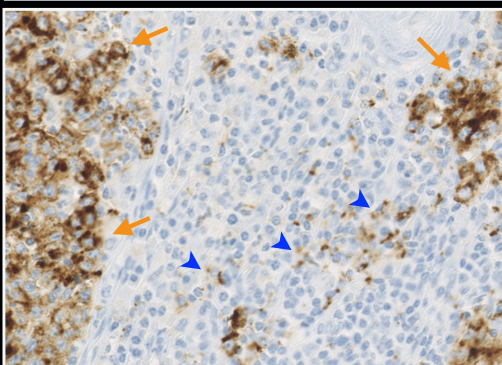
Results

Regarding EBV and MSI status, 3 groups were identified: EBV+/microsatellite stable (MSS) (n=16), EBV-/MSI-high (n=4), and EBV-/MSS (n=4). IRS>2 was restricted to EBV+ (6/16; 37.5%) and MSI-high (2/4; 50%) GCLSs.

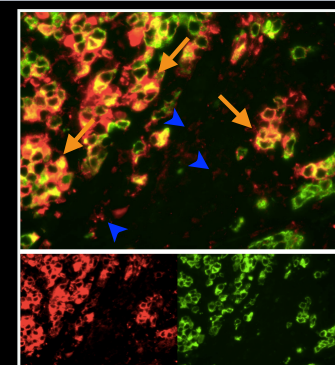
Overall, CD8/CD3R at the IF exceeded CD8/CD3R in the TC ($p<0.001$). CD8/CD3R was significantly higher ($p=0.008$) in EBV+ (n=16) than in EBV- (n=8) GCLSs.

By comparison with non-GCLSs, GCLSs were significantly associated with EBV infection (66.7% versus 5.56%, $p<0.001$) and PD-L1 protein expression (33.3% versus 13.0%, $p=0.04$). Moreover, GCLSs harbour distinctive clinico- pathological features: younger age ($p=0.03$), proximal location ($p=0.04$), indeterminate group of Lauren's classification ($p<0.001$), lower lymphatic invasion ($p=0.02$), lower pTNM stage ($p=0.002$) and better overall survival ($p=0.01$).

Tumour epithelial cells: negative (IRS≤2) Tumour immune cells: negative (<1%)	Tumour epithelial cells: positive (IRS>2) Tumour immune cells: positive (≥1%)	Tumour epithelial cells: negative (IRS≤2) Tumour immune cells: positive (≥1%)
n=2 (8.3%)	n=8 (33.3%)	n=14 (58.3%)



PD-L1 expression. Membranous, linear and strong PD-L1 expression was observed in tumour epithelial cells (arrows), whereas cytoplasmic dotted/granular pattern was observed in stromal immune cells (arrowheads). This GCLS case harboured PD-L1 amplification.



Conclusions

GCLSs are characterised by distinctive clinico-pathological features, EBV infection (66.7%), PD-L1 expression (33.3%) and high CD8/CD3R (at the IF and in EBV+ cases). In keeping with the recent molecular data, PD-L1 expression (IRS>2) was restricted to EBV+ and MSI-high cases, reinforcing the potential implications of immunotherapy in these molecular subtypes of GCLS.