

PD-L1/PD-1 Axis in Glioblastoma Multiforme

Subjects: Oncology

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Glioblastoma (GBM) is the most popular primary central nervous system cancer and has an extremely expansive course. Aggressive tumor growth correlates with short median overall survival (OS) oscillating between 14 and 17 months. The survival rate of patients in a three-year follow up oscillates around 10%. The interaction of the proteins programmed death-1 (PD-1) and programmed cell death ligand (PD-L1) creates an immunoregulatory axis promoting invasion of glioblastoma multiforme cells in the brain tissue. The PD-1 pathway maintains immunological homeostasis and protects against autoimmunity. PD-L1 expression on glioblastoma surface promotes PD-1 receptor activation in microglia, resulting in the negative regulation of T cell responses.

Keywords: PD1 ; PD1 ligand ; glioblastoma multiforme

1. Introduction

Glioblastoma (GBM) is the most common primary cancer in the central nervous system and has an extremely expansive course ^[1]. The standard approach to glioma treatment consists in the most extensive as possible surgical resection and in adjuvant radiation strengthened by temozolomide (TMZ) administration ^[2]. The median overall survival (OS) oscillates between 14 and 17 months ^{[3][4]}. The survival rate of patients in a three-year follow up oscillates around 10% ^[5]. GBM resistance to typical therapies requires verification. Glioblastoma cells interact with the surrounding environment, creating forceful interactions among heterogenous cell groups, various chemokines with cytogenetic effects, and extracellular proteins stimulating tumorigenesis, uncontrolled multifocal expansion, and immunological evasion ^[6].

The proteins programmed death-1 (PD-1) and programmed cell death ligand (PD-L1) interplay, creating an immunoregulatory axis promoting invasion of glioblastoma multiforme cells in the brain tissue ^[7]. Physiologically, the main function of PD-1 is to restrain T-cell anti-tumor activity and amplify Tregs activation, which limit T-cell reaction and protect against hyper immunity. The PD-L1/PD-1 axis maintains immunological homeostasis and protects against autoimmunity ^[8].

2. PD-L1 and PD-1 Structure

PD-L1 is encoded by the gene *PDCDL1*, localized on the 9th chromosome in p24.1 position. PD-L1 is also described as CD274 and B7-H1. This ligand was discovered and described in 1999 by Dong et al. as a member of the B7 protein family ^[9]. Seven exons encode the full-length protein PD-L1, consisting of 290 amino acids (40 kDa). PD-L1 as a type-I transmembrane complex of proteins includes single IgV and IgC domains on the external part, a transmembrane domain with hydrophobic properties, and a 30-amino acid cytoplasmatic tail as a signal transducer ^[10].

PD-L1 activity depends on binding to the PD-1 receptor encoded by the *PDCD1* gene. It is information transcribed from the second chromosome and consists of 288 amino acids (50–55 kDa). It contains an IgV domain in the extracellular domain and transmembrane region ^[11].

The intracellular region forms a tail composed of a tyrosine-based switch motif (ITSM)–inhibitory motif. This receptor was described by Ishida et al., who used subtractive hybridization to identify genes regulating programmed cell death ^[12]. PD-L1 is expressed and excreted by neoplastic cells, APCs, lymphocytes B, and parenchymal cells. It induces T-cell apoptosis or anergy and modulates inflammation in situ ^{[13][14]}.

The binding of PD-1 to the corresponding PD-1 receptor activates the protein tyrosine phosphatase SHP-2, which dephosphorylates Zap 70 (**Figure 1**). This process T cells proliferation and downregulates lymphocyte cytotoxic activity ^[15].

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- GLIOMA CELL**
- The diagram illustrates the complex signaling pathways within a glioma cell that regulate the expression of PD-L1. Key components and interactions include:
- External Stimuli:** VGF (Vascular Growth Factor) and EGFR (Epidermal Growth Factor Receptor) are shown as ligands. PAMPs (Pathogen-Associated Molecular Patterns) and Necrosis HSP (Heat Shock Protein) are also indicated.
 - Signaling Pathways:**
 - EGFR Pathway:** VGF binds to EGFR, activating PI3K, Akt, and mTOR, which in turn activates S6K1.
 - MyD88 Pathway:** TLR (Toll-Like Receptor) and IFNGR1/IFNGR2 (Interferon Gamma Receptors) activate MyD88, leading to the activation of MEK/ERK and TRAF6.
 - STAT-1 Pathway:** IFNGR1/IFNGR2 and IFN-γ (Interferon Gamma) activate STAT-1.
 - IRF-1 Pathway:** IFNGR1/IFNGR2 and IFN-γ activate IRF-1.
 - Gene Expression Regulation:**
 - PD-L1 Gene and Promoter:** These are the primary targets for regulation by NF-κB, IRF-1, and STAT-1.
 - MXA Gene:** The MXA gene is also regulated by IRF-1 and produces endogenous interferon.
 - mRNA Processing and Translation:**
 - PD-L1 mRNA:** Transcribed from the PD-L1 gene, it has a 5'UTR and 3'UTR.
 - MXA mRNA:** Transcribed from the MXA gene.
 - Endoplasmic Reticulum:** The site of protein synthesis for PD-L1 and Nonclassical MHC.
 - Nonclassical MHC:** A complex that presents PD-L1 on the cell surface.
 - Other Factors:**
 - miR-S13:** A microRNA that targets the PD-L1 mRNA.
 - Autocrine Stimulation:** IFN-α and IFN-β (Interferon Alpha and Beta) are shown as factors that can stimulate the cell.
 - Tumour Microenvironment:** The overall context in which these processes occur, including the presence of tumour antigens.

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