GPR133 Promotes Glioblastoma Growth in Hypoxia

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lioblastoma (GBM) is an incurable brain malignancy in dire need of novel therapeutic targets.¹ A critical component of GBM's resistance to conventional therapies is a stem-like tumor cell population that employs cell-intrinsic and microenvironment-mediated mechanisms to support tumor progression.²⁻⁷ Among microenvironmental factors that promote GBM stem cell (GSC) phenotypes is intratumoral hypoxia,^{8,9} a naturally occurring consequence of vascular thrombosis in these tumors. While many of the effects of hypoxia on tumor progression are mediated by hypoxiainducible transcription factors (Hif),^{9,10} no effective Hif inhibitors have been developed. In our effort to identify novel targetable mediators of hypoxia-driven tumor progression, we recently published on the role that GPR133 (ADGRD1),^{11,12} an orphan member of the adhesion family of G protein-coupled receptors (aGPCRs),¹³⁻¹⁷ plays in GBM progression. Here, we describe evidence that GPR133 is selectively expressed in hypoxic regions of GBM via direct transcriptional upregulation by Hif1 α . Furthermore, we show that GPR133 knockdown abrogates tumor initiation. This compelling evidence suggests that GPR133 plays an important protumorigenic role in GBM, particularly in the context of hypoxia, and argues that it represents a novel therapeutic target.

This research was approved by NYU School of Medicine's IRB (Protocol 12–01130) and IACUC (protocol 160503).

ABBREVIATIONS: aGPCR, adhesion family of G protein-coupled receptor; ChIP, chromatin immunoprecipitation; GBM, glioblastoma; GSC, glioblastoma stem cell; Hif, hypoxia inducible transcription factor; PPN, pseudopalisading necrosis

FINDINGS

GPR133 is Expressed Within Hypoxic Regions of GBM

Our immunohistochemical analysis indicated that the GSC marker CD133 (PROM1) is enriched in the hypoxic regions of pseudopalisading necrosis (PPN) within human GBM biospecimens (Figure 1A). In order to identify novel genes involved in hypoxia-driven growth of GBM, we performed an RNA-seq comparison of FACS-sorted CD133+ and CD133- cells from a patient-derived human GBM culture (Figures 1B and 1C). One of the top genes enriched in the CD133+ tumor population was *GPR133* (*ADGRD1*), an orphan member of the aGPCR family (Figure 1D).

To test the expression pattern of GPR133 in human GBM, we analyzed histological specimens from 9 GBM patients and observed that all tumors expressed GPR133. No staining was observed in normal human brain specimens (Figure 2A). Within each tumor, GPR133 immunostaining was specific to areas of PPN, which also expressed the hypoxia marker Hif1 α , an Hif (Figures 2B and 2C).¹⁸

This observation raised the possibility that GPR133 expression is regulated by hypoxia and Hif1 α . To investigate this hypothesis, we subjected 8 patient-derived human GBM cultures to hypoxia (1% O₂) in vitro for 24 h. Six of 8 primary GBM cultures showed upregulation of GPR133 mRNA after hypoxia (Figure 2D). Furthermore, knockdown of Hif1 α with lentiviral shRNA reduced GPR133 mRNA (Figure 2E) and chromatin immunoprecipitation (ChIP)-PCR analysis demonstrated direct binding of Hif1 α to the GPR133 promoter (Figure 2F). These findings indicated that GPR133 is predominantly expressed in hypoxic areas of human GBM via direct transcriptional upregulation by transcription factor Hifl α .



GPR133 Promotes Tumor Growth In Vitro and In Vivo

Our next question was whether GPR133 promotes or suppresses tumor growth. To answer this question, we employed lentiviral shRNA-mediated knockdown of GPR133 in patientderived GBM cultures. Tumor cells with GPR133 knockdown (GPR133-KD) showed reduced proliferation compared to scrambled shRNA control, using Ki67 immunofluorescence staining (Figures 3A and 3B). In addition, the ability of GPR133-KD cells to form spheres was significantly reduced relative to controls (Figure 3C). These in vitro findings were reproduced with 2 different shRNA constructs, suggesting lack of nonspecific effects.

We then tested the effects of GPR133 knockdown on in vivo tumorigenicity. Human GBM cells bearing either scrambled or GPR133-KD shRNA were injected into the brains of immunocompromised NOD.SCID mice. We found that GPR133 knockdown prevented tumor initiation and, therefore, death in implanted mice (Figures 3D and 3E). This robust phenotype indicated that GPR133 is critical for tumorigenicity.

GPR133 Expression Correlates With Poor Prognosis in GBM

In order to understand the clinical relevance of our findings, we used the UCSC Cancer Genome Browser to analyze existing RNA-seq data from 160 GBM samples in the TCGA database.¹⁹ After ranking the samples according to expression levels of *GPR133*, we divided the samples into 2 groups: GPR133 low (percentile 1-50, blue) and GPR133 high (percentile 51-100, red; Figure 4). Then, we analyzed the survival of these 2 groups using Kaplan–Meier survival curves. We found that, in agreement with our mouse data, high *GPR133* expression correlates with poor prognosis and reduced survival (Figure 4). These data confirm that GPR133 has pro-tumorigenic function and suggest that its inhibition represents an attractive and novel therapeutic approach.

DISCUSSION

Understanding the molecular mechanisms that underlie the response of cancer cells to hypoxia is critical for the discovery



of new therapeutic targets, especially in tumors such as GBM, which exhibits extensive hypoxia. Previous literature demonstrated that low oxygen tension evokes stem-like phenotypes that lead to tumor progression.^{8,9,20-23} Furthermore, several lines of evidence have suggested that the reason for failure of antiangiogenic therapy in GBM lies in induction of hypoxia by vascular regression.^{24,25} The role of hypoxia-inducible transcription factors Hifl α and Hif2 α in the regulation of the response to hypoxia has been well documented.^{9,10,26} However, pharmacologic targeting of these transcription factors has been elusive to date.

Our experiments provide compelling evidence that GPR133 inhibition represents an appealing strategy for arresting hypoxiadriven tumor progression in GBM. First, its expression within hypoxic PPN areas in tumor biospecimens coincides with that of Hif1 α , because, as we demonstrated, the *GPR133* gene is a direct transcriptional target of Hif1 α . Second, GPR133 is not expressed in normal cerebral hemisphere tissue, indicating a favorable therapeutic window. Finally, genetic knockdown of GPR133 abrogates orthotopic tumor initiation and prevents death of implanted immunocompromised mice. This finding is in agreement with TCGA data that demonstrate an inverse correlation between the amount of tumoral *GPR133* mRNA and survival in GBM patients.

Our encouraging data lay the foundation for further translational development of GPR133 as a treatment target. Indeed, we are currently undertaking experiments toward further target validation, as well as toward development of GPR133 inhibitors. Importantly, we are also assessing whether GPR133 inhibition may be beneficial in other extracranial malignancies, in which hypoxia may contribute to disease progression.

Disclosures

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inversely correlates with survival. GBM tumors in the TCGA database (n = 160) were ranked by GPR133 mRNA expression and dichotomized into 2 groups: high and low. Kaplan–Meier survival curves indicated a significant difference in survival between the high and low groups (logrank P < .001). Adapted from reference 12, published OA-CC-BY-4.0.

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