CHAPTER 6

Multiregional tumor drug-uptake imaging by PET and microvascular morphology in end-stage diffuse intrinsic pontine glioma


* shared authorship

J Nucl Med. 2017 Jul 26
ABSTRACT

INTRODUCTION Inadequate tumor uptake of the vascular endothelial growth factor antibody bevacizumab could explain lack of effect in diffuse intrinsic pontine glioma. METHODS By combining data from a PET imaging study using $^{89}$Zr-labeled bevacizumab and an autopsy study, a 1-on-1 analysis of multiregional in vivo and ex vivo $^{89}$Zr-bevacizumab uptake, tumor histology, and vascular morphology in a DIPG patient was performed. RESULTS In vivo $^{89}$Zr-bevacizumab measurements showed heterogeneity between lesions. Additional ex vivo measurements and immunohistochemistry of cervicomedullary metastasis samples showed uptake to be highest in the area with marked microvascular proliferation. In the primary pontine tumor, all samples showed similar vascular morphology. Other histologic features were similar between the samples studied. DISCUSSION In vivo $^{89}$Zr-bevacizumab PET serves to identify heterogeneous uptake between tumor lesions, whereas subcentimeter intralesional heterogeneity could be identified only by ex vivo measurements. $^{89}$Zr-bevacizumab uptake is enhanced by vascular proliferation, although our results suggest it is not the only determinant of intralesional uptake heterogeneity.
INTRODUCTION
End-stage diffuse intrinsic pontine glioma (DIPG) shows glioblastoma histology [1] and overexpression of proangiogenic factors, including vascular endothelial growth factor (VEGF) [1,2]. However, whether DIPG patients benefit from anti-VEGF treatment, such as the monoclonal antibody bevacizumab, is still unclear [3–5]. So far, the results of trials with bevacizumab have been disappointing [6,7]. A recent drug imaging study assessing tumor uptake of $^{89}$Zr-labeled bevacizumab in DIPG showed inter- and intrapatient heterogeneity, but factors determining uptake are currently unknown [8]. Therefore, in this study we performed a 1-on-1 analysis of multiregional in vivo and ex vivo $^{89}$Zr-bevacizumab uptake, tumor histology, and vascular morphology in a DIPG patient.

METHODS
A 12-y-old girl presented with paralysis of the left abducens nerve. Brain MR imaging revealed a T1-weighted hypointense, T2-weighted hyperintense lesion infiltrating 50% of the pons [9], suggestive of DIPG. Biopsy demonstrated high-grade diffuse glioma features, and Sanger sequencing of H3F3A and HIST1H3B revealed a wild-type status [10]. Whole-exome sequencing showed no mutations in HIST1H3C or HIST2H3C either. A diagnosis of histone H3 wild-type DIPG was made [11]. The patient was enrolled in phase A of the VUMc-01 DIPG study (Dutch Trial Register: NTR2391) and treated with radiotherapy (54 Gy [30 treatments of 1.8-Gy each]) and weekly gemcitabine (175 mg/m²/wk) for 6 wk. Progression-free survival was 3.8 mo. After disease progression, the patient participated in a molecular imaging study (NTR3518), including an immuno-PET scan at 145 h after injection of a 0.1 mg/kg dose of bevacizumab labeled with $^{89}$Zr (0.9 MBq/kg), as previously described [8]. Four days afterward, the patient died (death was unrelated to study participation) and participated in an autopsy study. All 3 studies were approved by the Institutional Review Board of VUMc and performed in accordance with the Declaration of Helsinki. For each study, both parents gave written informed consent [8,12,13].

In vivo $^{89}$Zr-bevacizumab uptake measurements
In vivo $^{89}$Zr-bevacizumab uptake was quantified as decay-corrected maximum activity concentration (in Bq/mL) in manually delineated volumes of interest, that is, tumor areas with visually enhanced uptake. Activity concentrations were also converted into SUVs, that is, decay-corrected maximum activity concentration normalized to injected dose per body weight.
Ex vivo $^{89}$Zr-bevacizumab uptake measurements

Brain autopsy was performed 2 h after death. Multiple 0.5-cm$^3$ tumor and control samples (macroscopically nonaffected brain, cerebrospinal fluid, and blood) were obtained for ex vivo $^{89}$Zr-radioactivity measurements and histologic analysis. Ex vivo radioactivity concentrations were measured with a g-well counter (1480 Wizard; Wallac). The in vivo PET scanner (Ingenuity TF-128 PET/CT scanner; Philips Healthcare) was cross-calibrated with the well counter [14,15]. Ex vivo $^{89}$Zr-radioactivity concentrations were also quantified as corrected maximum activity concentration and converted into SUV.

Immunohistochemistry

Four-micrometer-thick formalin-fixed, paraffin-embedded tissue sections were stained with hematoxylin and eosin and immunostained as previously validated and described [16,17] against glial fibrillary acidic protein (1:400; Dako); MIB-1 (1:40; Dako); glucose transporter 1 (1:200; Thermo Scientific), smooth muscle actin (1:200; Dako), collagen-IV (1:50; Dako), CD34 (1:50; Dako), and VEGF (1:50; Pharmagen). Specific stainings were chosen to determine differences in proliferation grade in the vascular endothelial cells (MIB-1) of the tumor; to highlight vascular morphology by identifying endothelial cells (CD34), vascular smooth muscle cells, activated pericytes (smooth muscle actin), and the vascular basal membrane (collagen-IV); to visualize tumor morphology and resident astrocytes (glial fibrillary acidic protein); to identify blood–brain barrier integrity and areas of increased hypoxia (glucose transporter 1); and to determine VEGF expression. Immunopositivity was detected with 3,3’-diaminobenzidine. Omitting primary antibodies yielded no significant staining. Sections were photographed using a Leica DM6000B microscope.

RESULTS

In vivo $^{89}$Zr-bevacizumab uptake

The T1-weighted postgadolinium MR images obtained 8 d before death showed a 3.6 x 3.3 cm primary pontine tumor, with strong contrast enhancement but no apparent necrosis. Furthermore, extensive meningeal and subependymal metastases in the right ventricular trigone (VTM) and cervicomedullary junction (CMM) (Fig. 1), as well as in the right frontal lobe, left Sylvian fissure, left lateral ventricle posterior horn, and along the spinal cord, were observed (Supplementary Figs. 1A–1F). At 145 h, the most intense $^{89}$Zr-bevacizumab uptake positivity was seen in the CMM (corrected maximum activity concentration, 7,800 Bq/mL; SUV, 9.9; volume, 2.0 mL), followed by the primary pontine tumor (5,607 Bq/mL; SUV, 7.1; volume, 13.1 mL) and the VTM (2,537 Bq/mL; SUV, 3.2; volume, 1.2 mL).
FIGURE 1 | Gadolinium-enhanced T1-weighted MR images obtained 8 d before death, and corresponding PET slices obtained 4 d before death (145 h after $^{89}$Zr-bevacizumab injection), showing primary pontine tumor and metastases in VTM and CMM.
Corr. Acmax = corrected maximum activity concentration; VTM = ventricular trigone metastasis; CMM = cervicomedullary metastasis. Arrows represent areas of disease.

**Ex vivo $^{89}$Zr-bevacizumab uptake**

Ex vivo radioactivity measurements at 242 h were obtained for multiple tumor samples (Fig. 2A), including the VTM, primary tumor, CMM, and a small 0.02-g fragment of dural metastasis that was previously undetected by MRI and PET (Supplementary Fig. 2), and macroscopically normal areas of the pons, cerebellum, and dura. Tracer uptake was highest in the dural metastasis (22,652 Bq/mL; SUV, 28.7) and CMM samples, with a considerable difference between the cranial (27,152 Bq/mL; SUV, 34.4) and caudal (16,184 Bq/mL; SUV, 20.5) parts. Differences in uptake were also observed between the cranial (2,949 Bq/mL; SUV, 3.7) and caudal (11,343 Bq/mL; SUV, 14.4) parts of the primary tumor. Radioactivity was low in normal tissue samples (<1,844 Bq/mL; SUV, <2.3), cerebrospinal fluid (62 Bq/mL; SUV, 0.1), and blood (716 Bq/mL; SUV, 0.9).

**Correlation between in vivo and ex vivo $^{89}$Zr-bevacizumab uptake**

Figure 2B shows the ratio between the in vivo and ex vivo $^{89}$Zr-bevacizumab uptake of the primary tumor, VTM, and CMM.
FIGURE 2 | (A) Ex vivo $^{89}$Zr-bevacizumab uptake (242 h after injection) in tumor samples and macroscopically normal tissue samples, cerebrospinal fluid, and blood. (B) Correlation with in vivo $^{89}$Zr-bevacizumab uptake (arrows, 145 h after injection).

VTM = ventricular trigone metastasis; CMM = cervicomedullary metastasis; DM = dural metastasis; CSF = cerebrospinal fluid; Corr. Acmax = corrected maximum activity concentration. *Numerator for ex vivo uptake of primary tumor and CMM is average of cranial and caudal corr. ACmax.

Correlation between ex vivo $^{89}$Zr-bevacizumab uptake and histology

After ex vivo radioactivity measurements, multiple samples from the primary tumor and CMM were investigated with histochemistry and immunohistochemistry to explore differences in histology as a possible explanation for tracer uptake heterogeneity. Hematoxylin and eosin staining showed a diffuse growth pattern, high cellularity, and variable cytonuclear atypia (Figs. 3A and 3B) in all tumor samples and confirmed the absence of tumor in macroscopically normal tissue. Necrosis and florid glomeruloid microvascular proliferation were present only in the cranial CMM (with the highest ex vivo radioactivity) (Fig. 3A). In this sample, MIB-1 staining was positive in up to 50% of tumor cells and in numerous endothelial cells in the microvascular walls (Fig. 3B).
FIGURE 3 | Histology and immunohistochemistry of samples from cranial and caudal parts of primary pontine tumor and CMM, respectively: hematoxylin and eosin (A); MIB-1 (for detection of proliferating cells) (B); astrocyte marker glial fibrillary acidic protein (C); CD34 (endothelial cell marker) and Glut-1 (glucose transporter expressed by brain endothelial cells as part of blood–brain barrier, and often deregulated in areas with hypoxia) (D); smooth muscle actin (marker of vascular smooth muscle cells and activated pericytes) and collagen-IV (component of vascular basal membrane) (E); and VEGF staining (F).

All images are 200x original magnification. Diffuse growth pattern is seen in all samples. Extensive necrosis and glomerular microvascular proliferation is seen with multiple MIB-1–positive (endothelial) cells in microvascular walls in caudal CMM. Partial loss of CD-34 is seen in caudal pons. Similar staining against glial fibrillary acidic protein, glucose transporter 1, and VEGF is seen in all samples. All vessels and microvessels are highlighted in stain against collagen-IV and smooth muscle actin.
In the remaining samples, microvascular proliferation was less pronounced: here, 30\% (in both primary tumor samples) to 50\% (in the caudal CMM sample) of tumor cell nuclei were MIB-1–positive, whereas endothelial cells were MIB-1–negative (Fig. 3B). Glial fibrillary acidic protein was expressed in tumor cells in the primary tumor and CMM, as well as in residual nonneoplastic astrocytes (Fig. 3C). Staining of CD34 showed partial loss of microvascular immunoreactivity in the caudal part of the primary tumor. In the other samples, CD34 immunopositivity was maintained (Fig. 3D). Immunoreactivity for glucose transporter 1 was similar in all samples (Fig. 3E). Staining of collagen-IV and smooth muscle actin highlighted the vascular basal lamina and tunica media, respectively, in all samples without obvious differences (Fig. 3E). Staining of VEGF was similar among all samples (Fig. 3F).

**DISCUSSION**

This study directly correlated multiregional $^{89}$Zr-bevacizumab uptake with tumor histology and vascular morphology in a patient with histone H3 wild-type DIPG [12]. Observed differences between in vivo and ex vivo radioactivity measurements reflected tracer pharmacokinetics (e.g., ongoing tracer clearance from blood over time) in combination with spatially or temporally heterogeneously increasing $^{89}$Zr-bevacizumab tissue deposition. Moreover, in vivo PET underestimates the true activity in small lesions (in this case, the CMM and VTM) because of partial-volume effects. Because of this effect [14], the actual radioactivity concentration as assessed by imaging at 145 h after $^{89}$Zr-bevacizumab injection was likely underestimated by approximately 70\%. Finally, as opposed to ex vivo measurements, at 145 h intravascular $^{89}$Zr-bevacizumab still contributes to the in vivo PET tumor signal (blood pool SUV, $\sim$4) [8]. Thus, whereas in vivo PET serves to identify whole-body tracer distribution and heterogeneity between lesions, the subcentimeter intralesional heterogeneity of tracer distribution can be defined only with ex vivo measurements.

Interestingly, hematoxylin and eosin and MIB-1 staining showed prominent microvascular proliferation, with higher $^{89}$Zr-bevacizumab uptake, in the cranial part of the CMM versus the caudal part, with lower uptake. No other histologic differences were found between the two samples, such as pattern of tumor growth, extent of necrosis, or difference in immunoreactivity for endothelial markers. In the two samples from the pons, intralesional uptake heterogeneity was also observed, although here both areas showed similar vascular morphology, tumor growth, and extent of necrosis. These findings suggest that vascular proliferation is an important, yet not the only, determinant of intralesional heterogeneity in $^{89}$Zr-bevacizumab uptake.
The presence of microvascular proliferation and disorganized vessels [18] in World Health Organization grade IV glioma is considered a consequence of hypoxia-induced overexpression of proangiogenic factors, including VEGF [1,2]. In the cranial CMM sample, glomeruloid microvascular structures with increased endothelial cell turnover (as determined by MIB-1-immunoreactivity) were prominent. On this basis, VEGF was expected to be overexpressed. VEGF immunoreactivity, however, showed mostly aspecific staining and did not differ between samples. This may be explained by the fact that the very short half-life of the VEGF protein (15–20 min [19]) was incompatible with its immunohistochemical detection 2 h after death. Glucose transporter 1 staining was similar in all samples analyzed, suggesting a partially intact blood–brain barrier. However, our histologic analysis did not provide definitive clues to determine the role of blood–brain barrier integrity in $^{89}$Zr-bevacizumab uptake in this case.

**CONCLUSION**

Although concerning a single case, we conclude that in vivo PET is capable of detecting heterogeneity in $^{89}$Zr-bevacizumab uptake between lesions, which correlates well with ex vivo measurements. However, PET cannot detect subcentimeter intralesional uptake heterogeneity. Furthermore, our results suggest that targeting of $^{89}$Zr-bevacizumab is enhanced in areas with vascular proliferation. However, because significant inter- and intralesional heterogeneity was also observed in areas that did not show differences in vascular proliferation, other factors present in the DIPG microenvironment likely also play a role. Because the DIPG microenvironment is heterogeneous and dynamic [3], patients are likely to fail treatment if timing and patient selection are not optimized. The results of this study underline the potential of immuno-PET studies, especially when combined with biopsy or autopsy studies, in the quest for optimal selection and timing of treatment schedules.
SUPPLEMENTARY DATA

A

Primary pontine tumor
MRI + / PET +

B

Cervicomedullary junction
MRI + / PET +

C

Spinal cord
- MRI +
- PET +
SUPPLEMENTARY FIGURE 1 | Gadolinium-enhanced T1-MRI, PET and macroscopy (if available) images of (A) the primary pontine tumor and metastases at the (B) cervicomedullary junction, (C) spinal cord, (D) posterior horn of the left lateral ventricle, (E) right frontal lobe and (F) left Sylvian fissure.

White arrows represent areas of $^{89}$Zr-bevacizumab uptake and/or disease.
SUPPLEMENTARY FIGURE 2 | Macroscopic picture of the dural metastases (white arrows), which were detected during autopsy but were undetected by MRI and PET. The dotted arrow indicates the primary pontine tumor.
REFERENCES


