The microenvironment in sporadic and neurofibromatosis type II–related vestibular schwannoma: the same tumor or different? A comparative imaging and neuropathology study

Daniel Lewis, MRCS, MA,1,2 Carmine A. Donofrio, MD,1 Claire O’Leary, PhD,1,3 Ka-Joh Li, PhD,2 Xiaoping Zhu, MD, PhD,2 Ricky Williams,1 Ibrahim Djoukhadar, PhD, FRCR,1 Erjon Agushi, MBBS,2 Cathal J. Hannan, MRCS,1 Emma Stapleton, FRCS(ORL-HNS),4 Simon K. Lloyd, MPhil, FRCS(ORL-HNS),4 Simon R. Freeman, FRCS(ORL-HNS),4 Andrea Wadeson, BSc,1 Scott A. Rutherford, FRCS(SN),1 Charlotte Hammerbeck-Ward, PhD, FRCS(SN),1 D. Gareth Evans, MD, FRCP,5 Alan Jackson, PhD, FRCR,2 Omar N. Pathmanaban, PhD, FRCS(SN),1,6 Federico Roncaroli, MD,1,3 Andrew T. King, FRCS(SN),1,7 and David J. Coope, PhD, FRCS(SN)1,3

1Department of Neurosurgery, Manchester Centre for Clinical Neurosciences, Salford Royal NHS Foundation Trust, Manchester Academic Health Science Centre; 2Division of Informatics, Imaging and Data Sciences, Wolfson Molecular Imaging Centre (WMIC), University of Manchester; 3Division of Neuroscience and Experimental Psychology, School of Biological Sciences, Faculty of Biology, Medicine and Health, University of Manchester; 4Department of Otolaryngology, Salford Royal NHS Foundation Trust, Manchester Academic Health Science Centre; 5Department of Evolution and Genomic Sciences, School of Biological Sciences, Faculty of Biology, Medicine and Health, University of Manchester; 6Division of Cell Matrix Biology & Regenerative Medicine, School of Biological Sciences, Faculty of Biology, Medicine and Health, University of Manchester; and 7Division of Cardiovascular Sciences, School of Medical Sciences, Faculty of Biology, Medicine and Health, University of Manchester, United Kingdom

OBJECTIVE  Inflammation and angiogenesis may play a role in the growth of sporadic and neurofibromatosis type 2 (NF2)–related vestibular schwannoma (VS). The similarities in microvascular and inflammatory microenvironment have not been investigated. The authors sought to compare the tumor microenvironment (TME) in sporadic and NF2-related VSs using a combined imaging and tissue analysis approach.

METHODS  Diffusion MRI and high-temporal-resolution dynamic contrast-enhanced (DCE) MRI data sets were prospectively acquired in 20 NF2-related and 24 size-matched sporadic VSs. Diffusion metrics (mean diffusivity, fractional anisotropy) and DCE-MRI–derived microvascular biomarkers (transfer constant \(K_{\text{trans}}\), fractional plasma volume, tissue extravascular-extracellular space \(v_e\), longitudinal relaxation rate, tumoral blood flow) were compared across both VS groups, and regression analysis was used to evaluate the effect of tumor size, pretreatment tumor growth rate, and tumor NF2 status (sporadic vs NF2-related) on each imaging parameter. Tissues from 17 imaged sporadic VSs and a separate cohort of 12 NF2-related VSs were examined with immunohistochemistry markers for vessels (CD31), vessel permeability (fibrinogen), and macrophage density (Iba1). The expression of vascular endothelial growth factor (VEGF) and VEGF receptor 1 was evaluated using immunohistochemistry, Western blotting, and double immunofluorescence.

RESULTS  Imaging data demonstrated that DCE-MRI–derived microvascular characteristics were similar in sporadic and NF2-related VSs. \(K_{\text{trans}}\) \((p < 0.001)\), \(v_e\) \((p \leq 0.004)\), and tumoral free water content \((p \leq 0.003)\) increased with increasing tumor size and pretreatment tumor growth rate. Regression analysis demonstrated that with the exception of mean
Approximately 95% of vestibular schwannomas (VSs) occur as a unilateral, sporadic tumor. Bilateral VSs are, conversely, the hallmark of patients with the dominantly inherited tumor syndrome, neurofibromatosis type 2 (NF2).1 Sporadic and NF2-related VSs share biallelic inactivation of the NF2 gene,2–4 but they present with phenotypic differences.5–6 NF2-related VSs develop as multifocal tumors along cranial nerve VIII5–6 and are therefore difficult to remove. Sporadic VSs are challenging due to their often large size at presentation and their unpredictable growth rate.7

It is currently unknown if the tumor microenvironment (TME) differs in sporadic and NF2-related VSs. There is growing evidence that tumor-associated macrophages (TAMs) may play a role in the growth and progression of sporadic VSs.8–10 Previous studies examined the inflammatory microenvironment in NF2-related tumors,11,12 but TME data in both VS groups are limited as lesions are often resected at different time points in their growth trajectory.13–16 Angiogenesis has also received increased attention as a potential therapeutic target in both sporadic and NF2-related tumors.17–20 Tumor microvasculature and tumor microstructure can be investigated and monitored with advanced MRI modalities such as dynamic contrast-enhanced (DCE) MRI.21 DCE-MRI–derived microvascular parameters correlate well with tissue markers of permeability, microvessel surface area (SA), and cell density.22,23

In the present study we used both DCE-MRI and diffusion MRI to compare the microvascular and microstructural characteristics across a cohort of sporadic and NF2-related VSs and establish if their TMEs are similar. By studying the tissue of resected tumor specimens, we sought to establish to what extent DCE-MRI–derived imaging biomarkers reflect the inflammation and microvasculature in these tumors and understand the relationship between inflammation and angiogenesis in both VS groups.

**Methods**

**Study Design and Patient Population**

DCE-MRI data sets from two separate patient cohorts were used in this study. Between December 2015 and August 2018, 24 patients with sporadic VSs were recruited via the Manchester Skull Base Unit multidisciplinary team meeting at Salford Royal Hospital in Salford, United Kingdom. Fifteen of these patients were recruited under a combined [11C]-(R) PK-11195 PET and DCE-MRI study. A further 9 patients were recruited under a separate ethics protocol and underwent only the MRI acquisition. All patients underwent at least two MRI acquisitions to establish tumor growth, including the study scan in 6 patients. The study MRI scan and the results of previous MRI were reviewed collectively by the multidisciplinary team and tumors were classified as static/shrinking, growing, or sufficiently large to preclude monitoring of growth and mandate early resection. This classification was based upon clinical decision, with patients with growing VSs recommended to undergo either microsurgery or stereotactic radiosurgery (SRS), and patients with static or shrinking tumors offered a period of observation. The median length of follow-up from diagnosis was 29 months (IQR 23–49 months) for the static/shrinking cohort, 12 months (IQR 8–25 months) for growing tumors, and 3 months (IQR 2–6 months) for the large tumor cohort, reflecting the early decision to treat these patients. Volumetric measurements of tumor size were made for both the study MRI scan and the preceding clinical scan to confirm different growth behaviors between the tumor cohorts.8

In addition, pretreatment DCE-MRI data from 12 patients with NF2 who had been recruited as part of a previous study at our institution were retrospectively analyzed.24 These patients had been recruited through the NF2 clinic in Manchester, United Kingdom, and following imaging, all patients underwent treatment with the anti–vascular endothelial growth factor (VEGF) antibody bevacizumab. All patients had proven NF2 with at least 1 VS demonstrating a high growth rate, defined as an increase of 4 mm or greater in the maximal transverse diameter over a 12-month period.25 Similar to the sporadic VS cohort, volumetric measurements of tumor size were made for both the study MRI scan and the preceding clinical scan. For all VS studies, tumor extrameatal extension and the degree of tumor brainstem compression at the time of the study scan were reported using the Koos classification system.44

**Imaging Methods**

**MRI Acquisition**

For all patients, MRI data were acquired on a 1.5-T whole-body scanner using a dedicated head coil (Philips Achieva, Philips Medical Systems). A high-resolution 3D T1-weighted gradient echo sequence with full brain coverage (TE 3.2 msec, TR 8.6 msec, slice thickness 1.2 mm) both before and after contrast was obtained for tumor delineation. Diffusion-weighted imaging (DWI) was
obtained using 6 directions and 3 B values (0, 400, and 800 sec/mm$^2$), and DCE-MRI data were collected using a previously described dual-injection, dual-temporal-resolution technique. The details of which can be found in the Supplementary Methods.

**MRI Data Analysis**

Acquired DWI data for both the sporadic and NF2 patient cohorts were processed using the FSL 4.1 Diffusion Toolbox (http://www.fmrib.ox.ac.uk/fsl/) through a standard multistep procedure that incorporated both eddy current correction and brain extraction. Through this approach, voxel-wise maps of mean diffusivity (MD) and fractional anisotropy (FA) were generated. Voxel-wise maps of microvascular kinetic parameters and absolute cerebral blood flow (CBF$_{ET}$) estimates were derived from the low-dose, high-temporal-resolution DCE-MRI data sets using the extended Tofts model and a previously developed T1-weighted early time points method, respectively. Individual tumors were manually delineated on the coregistered T1-weighted postcontrast image to create a tumor object mask. These object masks were then applied to the CBF$_{ET}$ map, kinetic parameter maps, and diffusion parameter maps (MD, FA) to allow derivation of whole tumor blood flow, mean kinetic parameter estimates (transfer constant [Ktrans]), tissue extravascular-extracellular space ($v_v$), fractional plasma volume ($v_p$), longitudinal relaxation rate [R1s]), and diffusion metrics (MD, FA). Individual tumor volumes were determined on postcontrast T1-weighted imaging. Further details are provided in the Supplementary Methods.

**Tissue Analysis**

Tissues from 17 sporadic tumors that had undergone DCE-MRI and a separate cohort of 12 previously resected NF2-related VSs were analyzed. Ethical approval was given for all tissue analyses. Frozen samples were collected for 10 patients (6 sporadic and 4 NF2-related VSs). Serial 5-µm sections were cut from each paraffin block and used for H & E staining, immunoperoxidase immunohistochemistry, and immunofluorescence. Tissue sections were assessed quantitatively for TAM density (Iba1), microvessel area (CD31), vascular permeability (fibrinogen), and cellular proliferation (anti–Ki-67) using immunoperoxidase immunohistochemistry. Semiquantitative evaluation of VEGF and VEGF receptor 1 (VEGFR-1) was also performed. Western blot analysis on the 10 frozen specimens was used to confirm VEGF and VEGFR-1 protein expression. Double immunofluorescence staining for Iba1 and VEGF was performed on 8 paraffin sections (4 sporadic and 4 NF2-related VSs) to evaluate the VEGF expression in macrophages. Detailed protocols are described in Supplementary Methods.

For quantitative analysis, 20 representative images were taken at magnification ×20 from the H & E–stained sections and sections stained with Iba1, CD31, fibrinogen, VEGF, and VEGFR-1 for each case using a 3DHistech Pannoramic-250 microscope slide-scanner (3DHistech Ltd.). All images were analyzed using ImageJ software (https://imagej.nih.gov/ij/). Quantification was performed by two independent observers (D.L. and C.A.D.); interobserver agreement was greater than 95% across all measurements, and results were further validated by an experienced neuropathologist (F.R.) by reviewing and quantifying random images from each case. Negative controls were performed by substituting the primary antibody with phosphate-buffered saline. Normal tonsil and appendix tissues served as positive controls for all primary antibodies tested (Supplementary Fig. S1). Detailed tissue analysis protocols are shown in the Supplementary Methods.

**Statistical Analysis**

The statistical program Stata (version 11, StataCorp LLC) was used for all statistical analyses. Normality and homogeneity of variance for all individual data parameters were assessed using the Shapiro-Wilk and Levene tests, respectively. For comparison of parametric diffusion and DCE-MRI–derived values between sporadic and NF2-related VSs, a 2-tailed t-test was used. For nonparametric data (patient age, tumor size, annual adjusted growth rate) the Mann-Whitney U-test was used. The intetumor correlations between tumor size, imaging-derived parameters, and tissue-derived parameters are reported as Pearson’s product-moment correlation coefficient ($r$), or Spearman’s rho in the case of nonlinear associations. For linear associations, the results of linear regression are reported as adjusted $R^2$ estimates on included figures.

A 2-tailed t-test was used for comparison of parametric tissue–derived parameters (CD31 microvessel SA, fibrinogen optical density [OD], Ki-67 labeling index) between sporadic and NF2-related VSs and between high and low intratumoral Iba1$^+$ areas. For comparison between sporadic VS growth groups (static, growing, large) a 1-way ANOVA with Bonferroni correction was used. Semiquantitative (VEGF, VEGFR-1) and categorical variables were compared using either Pearson’s chi-square or Fisher’s exact test. To evaluate the effect of tumor size, annual adjusted growth rate, and tumor NF2 status on imaging and tissue-derived parameters, simple univariate and multiple linear regression models, with tumor NF2 status as an interaction term, were used. Nonlinear associations were addressed through use of the cube root transform of tumor size and tumor annual adjusted growth rate in the regression models. Residual normality was analyzed using the Jarque-Bera test and by assessing the linearity of quantile-quantile plots. Heteroscedasticity was evaluated using White’s test and by producing scatterplots of residuals versus predicted values. Multicollinearity between predictor variables was assessed through postregression testing and collinear variables were excluded.

**Results**

**Patient Population**

Details of the demographics of the 24 patients with sporadic VS and 12 NF2 patients included in the imaging study are presented in Table 1. The median age across the sporadic VS cohort was 55.3 years (IQR 42.2–62.1 years). Seventeen patients underwent resection, 2 were treated with SRS, and 5 had no treatment and are undergoing long-term follow-up. Eight tumors were classified as...
TABLE 1. Comparative table of patient demographics, DCE-MRI–derived microvascular parameters, and diffusion metrics between sporadic and NF2-related VSs

<table>
<thead>
<tr>
<th>Variable</th>
<th>Sporadic VS (n = 24)</th>
<th>NF2-Related VS (n = 20)</th>
<th>Mean Difference (95% CI)</th>
<th>p Value*</th>
</tr>
</thead>
<tbody>
<tr>
<td>Median age (IQR), yrs</td>
<td>55.3 (42.2–62.1)</td>
<td>24.0 (20.5–29.0)</td>
<td>−24.2 (−32.6 to −15.7)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Koos grade, n (%)</td>
<td></td>
<td></td>
<td></td>
<td>0.89†</td>
</tr>
<tr>
<td>II</td>
<td>7 (29)</td>
<td>6 (30)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>III</td>
<td>10 (42)</td>
<td>7 (35)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>IV</td>
<td>7 (29)</td>
<td>7 (35)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Median VS size (range), cm³</td>
<td>1.82 (0.34–18.67)</td>
<td>2.51 (0.20–17.87)</td>
<td>0.20 (−2.66 to 3.06)</td>
<td>0.65</td>
</tr>
<tr>
<td>Median annual adjusted growth rate (IQR), cm³/yr</td>
<td>0.62 (0.09–1.29)</td>
<td>0.75 (0.23–2.24)</td>
<td>0.16 (−0.78 to 1.11)</td>
<td>0.48</td>
</tr>
<tr>
<td>Mean tumor Ktrans (SD), min⁻¹</td>
<td>0.12 (0.05)</td>
<td>0.11 (0.04)</td>
<td>−0.01 (−0.04 to 0.02)</td>
<td>0.49</td>
</tr>
<tr>
<td>Mean tumor vₑ (SD)</td>
<td>0.05 (0.02)</td>
<td>0.05 (0.02)</td>
<td>−0.0006 (−0.01 to 0.01)</td>
<td>0.91</td>
</tr>
<tr>
<td>Mean tumor vₑ (SD)</td>
<td>0.37 (0.12)</td>
<td>0.35 (0.08)</td>
<td>−0.02 (−0.09 to 0.04)</td>
<td>0.49</td>
</tr>
<tr>
<td>Mean TBF (SD), ml/100 ml/min</td>
<td>93.3 (31.1)</td>
<td>89.5 (23.2)</td>
<td>−3.78 (−20.7 to 13.2)</td>
<td>0.65</td>
</tr>
<tr>
<td>Mean tumor R₁N (SD), sec⁻¹</td>
<td>0.68 (0.12)</td>
<td>0.76 (0.17)</td>
<td>0.07 (−0.01 to 0.16)</td>
<td>0.08</td>
</tr>
<tr>
<td>MD (SD), mm²/sec × 10⁻³</td>
<td>0.67 (0.09)</td>
<td>0.80 (0.12)</td>
<td>0.14 (0.07 to 0.20)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>FA (SD)</td>
<td>0.39 (0.06)</td>
<td>0.36 (0.05)</td>
<td>−0.03 (−0.07 to −0.000003)</td>
<td>0.05</td>
</tr>
</tbody>
</table>

Boldface type indicates statistical significance.

* The p value was calculated using a 2-tailed t-test, or Mann-Whitney U-test in the case of nonparametric data.
† Differences in tumor Koos grade determined using Pearson’s chi-square test.

static or shrinking (n = 1), 12 as growing, and 4 tumors as sufficiently large to preclude monitoring of growth and mandate early resection. Confirmatory volumetric measurements demonstrated that the growing tumors were significantly larger (2.28 vs 0.62 cm³, p < 0.001) and displayed a significantly higher annual adjusted growth rate compared with the static/shrinking group (0.77 vs 0.02 cm³/year, p < 0.001). The median size among the 4 large sporadic tumors was 13.0 cm³ (IQR 10.7–16.6 cm³), and while they were primarily treated due to large size at presentation, volumetric measurements confirmed growth in these tumors within the short follow-up period (median annual adjusted growth rate = 2.94 cm³/year, IQR 1.70–5.25 cm³/year). The NF2 patient cohort was significantly younger (median age was 24 years [IQR 20.5–29.0 years]). Of the 12 NF2 patients, 4 had undergone previous resection of a contralateral VS. Median tumor size (2.51 vs 1.82 cm³) and median annual adjusted growth rate (0.75 vs 0.62 cm³/year) were higher among NF2-related VSs compared to sporadic VSs, but these differences were not statistically significant (Table 1).

Features of the 17 sporadic VS and 12 NF2 patients whose tissue was used for analysis are detailed in Table 2. The NF2 patient cohort demonstrated significantly higher tumor volumes at the time of resection (median volume 8.43 vs 2.51 cm³, 95% CI mean difference = 0.53–17.9 cm³) and a nonsignificantly longer median period of radiological follow-up prior to resection (28 vs 13 months, 95% CI mean difference = −2 to 37 months). Among resected sporadic tumors, 3 were classified as static, 10 as growing, and 4 tumors as sufficiently large to preclude monitoring of growth and mandate early resection. Across the resected NF2-related VS cohort there was a range of pretreatment tumor growth rates, and these rates were comparable to those of the resected sporadic VS cohort (median annual adjusted growth rate = 0.80 vs 0.77 cm³/year, 95% CI mean difference = −1.17 to 1.65 cm³/year).

**Imaging Data**

Comparative DCE-MRI and diffusion MR-derived imaging metrics between the sporadic and NF2-related VS cohorts are shown in Table 1. NF2-related VSs displayed significantly higher tumor MD values (p < 0.001) and lower FA values (p = 0.05) than sporadic tumors. There was no significant difference in any of the DCE-MRI–derived microvascular metrics between the two tumor groups (p > 0.05).

Mean tumor Ktrans increased with increasing tumor size in both sporadic (rho = 0.83, p < 0.001) and NF2-related (rho = 0.77, p < 0.001) VSs (Fig. 1A). There were size-dependent increases in tumor vₑ in both tumor groups (p < 0.001), but in sporadic tumors this increase stopped once tumor volume exceeded 10 cm³ (Fig. 1B). Mean tumor R₁N decreased with increasing tumor size in both sporadic (rho = −0.58, p = 0.003) and NF2-related (rho = −0.86, p < 0.001) VSs (Fig. 1C). As shown in Fig. 1D, both sporadic and NF2-related VSs also demonstrated increases in tumor MD with increasing tumor size (p < 0.001), and there was a significant inverse linear relationship between tumor MD and FA across both tumor cohorts (p ≤ 0.002; Supplementary Fig. S3). Representative images from a patient with a large sporadic tumor and a patient with a large NF2-related VS are shown in Fig. 1E and F, respectively.

Univariate regression analysis (Supplementary Table S1) demonstrated that the cube root transform of tumor size and annual adjusted tumor growth rate were significant independent predictors of mean tumor Ktrans (p < 0.001), mean tumor vₑ (p ≤ 0.02), mean tumor R₁N (p < 0.001), mean tumor diffusivity (p ≤ 0.001), and mean...
tumor FA (p ≤ 0.003). Univariate and multiple regression analysis (Supplementary Table S2) demonstrated that with the exception of MD (p < 0.001), tumor NF2 status had no statistically significant effect on any of the imaging parameters or the observed relationship between the derived imaging parameters and tumor size (p > 0.05). Intertumor correlation analyses between different imaging metrics across each VS cohort are shown in Supplementary Tables S3 and S4.

### Imaging and Pathology Correlation Analysis

Correlation indices between imaging and tissue analysis in the 17 sporadic VSs are shown in Supplementary Table S5. Cell density correlated negatively with mean

<table>
<thead>
<tr>
<th>Variable</th>
<th>Sporadic VS (n = 17)</th>
<th>NF2-Related VS (n = 12)</th>
<th>Mean Difference (95% CI)</th>
<th>p Value*</th>
</tr>
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<tbody>
<tr>
<td>Median age at operation (IQR), yrs</td>
<td>49.4 (41.3–55.8)</td>
<td>32.8 (27.1–42.8)</td>
<td>−12.2 (−21.3 to −3.1)</td>
<td>0.01</td>
</tr>
<tr>
<td>Sex, n (%)</td>
<td>Male 7 (41.2)</td>
<td>6 (50)</td>
<td></td>
<td>0.64</td>
</tr>
<tr>
<td></td>
<td>Female 10 (59)</td>
<td>6 (50)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Tumor laterality, n (%)</td>
<td>Lt 10 (59)</td>
<td>5 (42)</td>
<td></td>
<td>0.36</td>
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<tr>
<td></td>
<td>Rt 7 (41.2)</td>
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<tr>
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<td>Median tumor volume (IQR), cm³</td>
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<td>Median annual adjusted growth rate (IQR), cm³/yr</td>
<td>0.77 (0.54–2.71)</td>
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<td>0.23 (−1.17 to 1.65)</td>
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<td>Median length of radiological follow-up (IQR), mos</td>
<td>13 (8–24)</td>
<td>28 (6–67)</td>
<td>18 (−2 to 37)</td>
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**FIG. 1.** Comparison of DCE-MRI–derived microvascular parameters and diffusion metrics between sporadic and NF2-related VSs. A: Intertumor scatterplot analysis of mean tumor $K_{\text{trans}}$ against VS size (cm³). B: Intertumor scatterplot analysis of mean tumor $v_e$ against VS size (cm³). C: Intertumor scatterplot analysis of mean tumor $R_1N$ against VS size (cm³). D: Intertumor scatterplot analysis of tumor MD (mm²·s⁻¹ × 10⁻³) against VS size (cm³). E and F: Parametric maps from a patient with a large sporadic tumor and a patient with a large NF2-related VS are shown in E and F, respectively. Mean parameter values for the VSs displayed in E and F are outlined in the scatterplot shown in panel A through the use of a blue and red box, respectively. Spearman’s rho reported for the intertumor correlation across all 44 VSs (A–D). EES = extravascular-extracellular space; $T1W+C$ = T1-weighted postcontrast.

*The p value was calculated using the Mann-Whitney U-test, or Pearson’s chi-square test in the case of tumor laterality, sex, and tumor Koos grade.

**TABLE 2.** Comparative table of patient demographics, tumor laterality, and tumor size for the patients included in the pathology study

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Boldface type indicates statistical significance.

* The p value was calculated using the Mann-Whitney U-test, or Pearson’s chi-square test in the case of tumor laterality, sex, and tumor Koos grade.
tumor \( v_c \) (\( \rho = -0.61, p = 0.02 \)). Tumor MD correlated positively with fibrinogen OD (\( \rho = 0.70, p = 0.002 \)), but there was no correlation observed between cell density and either diffusion metric. Tumor \( K^{\text{trans}} \) correlated positively with both mean fibrinogen OD (\( r = 0.62, p = 0.008 \)) and microvessel SA (\( r = 0.74, p < 0.001 \)). Derived \( v_c \) estimates correlated strongly with microvessel SA (\( r = 0.89, p < 0.001 \); Supplementary Fig. S4).

The 10 rapidly growing VSs demonstrated higher TAM density compared to both the static (\( p < 0.001 \)) and large (\( p = 0.003 \)) VS groups (Supplementary Table S6). There was a nonsignificant increase in TAM infiltrates with increasing tumor \( K^{\text{trans}} \) across all tumors (\( \rho = 0.23, p = 0.38 \); Fig. 2A). The 4 large VSs demonstrated marked heterogeneity in TAM density, and after exclusion of these tumors the relationship between \( v_c \) cell percentage and \( K^{\text{trans}} \) was statistically significant (\( \rho = 0.76, p = 0.002 \)). Compared to static tumors, growing VSs displayed significantly greater overall microvessel SA (\( p = 0.011 \)). Image sections from a growing VS with high \( K^{\text{trans}} \) values are shown in Fig. 2B alongside comparative immunostains demonstrating the high vascular permeability and dense TAM infiltrates seen within this tumor (Fig. 2C). Across all VS tissue specimens there was a positive correlation between microvessel SA and \( v_c \) TAM density (\( r = 0.55, p = 0.002 \); Fig. 2D).

Representative tissue sections from both a slowly growing and rapidly growing NF2-related VS are shown in Fig. 2E and F, respectively, and demonstrate the close correspondence between tumor vascularity and TAM infiltrates.

**Sporadic and NF2-Related VS Pathology Comparison**

As shown in Table 3, NF2-related VSs demonstrated significantly lower fibrinogen OD values (\( p < 0.001 \)) compared to growing sporadic VS; there was no significant difference in either TAM density (\( p = 0.65 \)) or microvessel SA (\( p = 0.16 \)). Univariate linear regression analysis to evaluate the effect of tumor size, tumor growth rate, and tumor NF2 status on tissue-derived parameters is shown in Supplementary Table S7. Whereas both tumor size and tumor growth were significant independent predictors of mean \( v_c \) cell percentage (\( p \leq 0.04 \)) and mean CD31 percentage microvessel SA (\( p \leq 0.01 \)), with the exception of mean tissue fibrinogen OD (\( p = 0.001 \)), tumor NF2 status
Relationship Between Microvessel SA, TAM Density, and Cellular Proliferation Indices

Figure 3 demonstrates the relationship between TAM density, microvessel SA, and Ki-67 labeling index in the 4 large sporadic VSs. Both microvessel SA (p < 0.001; Fig. 3A) and fibrinogen OD (p < 0.001) were significantly elevated in areas of high TAM density compared to low-TAM-density regions (Supplementary Table S8). Across all tumor areas, TAM density correlated with fibrinogen OD (rho = 0.71, p < 0.001; Fig. 3B) and microvessel SA (rho = 0.52, p < 0.001). The Ki-67 labeling index was significantly greater in regions of high TAM density compared to low-density regions (p < 0.001; Fig. 3C).

TABLE 3. Comparative tissue-derived parameters between sporadic and NF2-related VSs

<table>
<thead>
<tr>
<th>Variable</th>
<th>Sporadic VS, All (n = 17)</th>
<th>Sporadic VS, Growing (n = 10)</th>
<th>NF2-Related VS (n = 12)</th>
<th>Mean Difference (95% CI)</th>
<th>p Value</th>
<th>Mean Difference (95% CI)</th>
<th>p Value*</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mean Iba1+ cell %</td>
<td>46.5 ± 20.2</td>
<td>59.9 ± 9.75</td>
<td>56.3 ± 23.2</td>
<td>9.73 (−6.88 to 26.34)</td>
<td>0.24</td>
<td>-3.62 (−20.05 to 12.81)</td>
<td>0.65</td>
</tr>
<tr>
<td>Mean CD31 % microvessel SA</td>
<td>2.63 ± 1.26</td>
<td>2.81 ± 0.96</td>
<td>2.20 ± 1.02</td>
<td>-0.43 (−1.33 to 0.47)</td>
<td>0.33</td>
<td>-0.62 (−1.50 to 0.27)</td>
<td>0.16</td>
</tr>
<tr>
<td>Cell density (H &amp; E cell nuclei/20 HPF)</td>
<td>2823 ± 906</td>
<td>2401 ± 865</td>
<td>2272 ± 621</td>
<td>-550 (-1170 to 70)</td>
<td>0.08</td>
<td>-129 (-790 to 533)</td>
<td>0.68</td>
</tr>
<tr>
<td>Mean fibrinogen OD (%)</td>
<td>0.37 ± 0.13</td>
<td>0.37 ± 0.08</td>
<td>0.24 ± 0.03</td>
<td>-0.12 (−0.20 to −0.04)</td>
<td><strong>0.004</strong></td>
<td>-0.12 (−0.18 to −0.07)</td>
<td>&lt;0.001</td>
</tr>
</tbody>
</table>

HPF = high-power field.

Data for A, B, and C are given as mean ± SD. Boldface type indicates statistical significance.

* The p value was calculated using 1-way ANOVA with Bonferroni correction.

had no statistically significant effect on any of the tissue parameters.
section from a large sporadic VS with corresponding regions of both high and low TAM density, microvessel SA, and vascular leak is shown in Fig. 3D–F.

Tumor VEGF and VEGFR-1 Expression

VEGF expression was present in all sporadic and NF2-related VSs (Fig. 4A). Western blot results confirmed the immunohistochemistry data. The molecular weight of VEGF was approximately 20 KDa. VEGF expression varied across individual cases, but there were no statistically significant differences between sporadic and NF2-related VSs \( (p = 0.21, \text{Fishers exact test}) \). VEGF-1 was expressed in all cases. Western blot analysis confirmed the expression (Fig. 4B) and showed a molecular weight of approximately 65 KDa, consistent with the soluble isoform of the VEGFR-1.27–29 Sporadic and NF2-related VSs showed spatial correspondence of areas of high TAM density and VEGF expression (Fig. 5A and B). Double immunofluorescence demonstrated that VEGF was expressed mostly by TAMs (Fig. 5C), although the intensity of VEGF staining varied from area to area. We were unable to confirm the expression of VEGFR-1 in TAMs due to the low signal of VEGFR-1 in immunofluorescence.

Discussion

This study expanded on our previous observation showing that an increase in inflammation and vascular permeability correlates with growth in sporadic VSs.8 The present findings demonstrate that sporadic and NF2-related VSs show similar microvasculature and a similar relationship between neovascularization and intratumoral inflammation, and that VEGF-expressing TAMs are instrumental to angiogenesis. Our results also suggest that inflammation and angiogenesis can represent potential therapeutic targets in both tumor groups.

To our knowledge, this is the first study that compared DCE-MRI and diffusion MRI-derived microvascular and microstructural metrics in sporadic and NF2-related VSs. Alongside increases in tumor \( K^\text{trans} \), both sporadic and NF2-related VSs demonstrated concomitant increases in the extravascular-extracellular space fraction \( (\nu_e) \) and reductions in \( R_1\text{N} \), an inverse correlate of tumoral free water content\(^\text{11} \) with increasing tumor size. Derived diffusion metrics mirrored these changes and supported previous evidence\(^\text{10} \) that larger tumors display both higher MD and lower FA, a DTI-derived diffusion directionality measure sensitive to changes in tumor organization and microstructure\(^\text{31,32} \). Our imaging findings suggest that as both sporadic and NF2-related VSs grow and enlarge, there are not only increases in the capillary permeability SA product, but also concomitant increases in the extravascular-extracellular space and the degree of interstitial tumoral free water, and a reduction in the overall density of the tissue microstructural architecture.

Our results demonstrated a relationship between tumor vascularity and TAM infiltration and showed that DCE-MRI–derived vascular biomarkers such as \( K^\text{trans} \) can be used as indirect measures of inflammatory response, although a direct comparison between imaging data and tissue analysis was not feasible in the NF2 cohort. Nonetheless, analysis of a separate cohort of resected growing NF2-related VSs showed a strong correlation between TAMs and microvessel SA, similar to what was observed in sporadic VS.

The 4 large sporadic VSs in our cohort demonstrated disproportionately high \( K^\text{trans} \) values compared to the average TAM tissue density. The discrepancy between imaging and tissue data may result from limitations of our DCE-MRI technique when applied to highly vascular

FIG. 4. Expression of VEGF and VEGFR-1 in VS. A: Western blot demonstrating VEGF protein expression in a cohort of sporadic and NF2-related VSs, with β actin as a control. B: Western blot demonstrating VEGFR-1 protein expression in a cohort of sporadic and NF2-related VSs with β actin as a control. C: Immunostains (VEGFR-1 = brown, Iba1 = red) from a growing sporadic VS demonstrating spatial colocalization of VEGFR-1 and Iba1 immunostaining (left = immunoperoxidase—whole section, right = immunoperoxidase, x20 HPF). Figure is available in color online only.
tumors, but may also reflect the dramatic differences in spatial resolution between imaging and tissue-based techniques. Across these tumors, there was considerable heterogeneity in TAM density, with regions of both high and low vascularity and TAM infiltration, respectively, and the spatial resolution of our DCE-MRI–derived metrics may limit the ability to fully detect such heterogeneity in vivo. In a previous study it was demonstrated that within growing sporadic VSs, macrophages rather than Schwann cells accounted for the majority of proliferating cells.8 Our demonstration of elevated cellular proliferation rates within intratumoral regions of high TAM density further supports this finding, and suggests that even within individual tumors, growth occurs within regions of high vascularity and high TAM infiltration.

Previous studies have documented the proangiogenic cytokine VEGF and VEGFR-1 in VS tissue,17,18,33 but the cellular origin of these proteins and their relationship to intratumoral inflammation were not investigated. We demonstrated that intratumoral macrophages contribute significantly to VEGF production, and we showed predominant expression of the soluble form of the VEGFR-1.27,29,34 Alongside its roles in promoting vasodilatation, increasing vascular permeability, and inducing angiogenesis,17,35 VEGF isoforms act as a chemoattractant for circulating VEGFR-1–expressing macrophages.35–38 Combined imaging and pathology results presented here further support our previous results of increased inflammation and vascular permeability in growing sporadic VSs,10 but also suggest that VEGF/VEGFR-1 signaling drives chemotraction of VEGFR-1–expressing macrophages into the TME,36,37 in addition to driving angiogenesis. Further work is required to better evaluate the role that VEGFR-1–expressing macrophages play in VS angiogenesis and progression.

Among NF2 patients, the management of VS presents the biggest treatment challenge. Their bilateral and progressive nature causes considerable patient morbidity and makes treatment of these lesions through surgery or radiotherapy difficult.6,16 To date, the anti-VEGF antibody bevacizumab is the only targeted molecular therapy for these tumors,19,20 but concerns have been raised about its widespread usage due to cardiovascular and renal complications and the duration of response.39,40 Our evidence of a spatial relationship between TAM and angiogenesis in sporadic and NF2-related VSs suggests that treatment targeting VEGF may lead to a concomitant reduction in TAM and therefore further impact tumor growth. The fact that TAM rather than tumor cells proliferates in VS8 further supports the idea of targeting macrophages along with vascular supply. Recent studies have documented upregulation of other proinflammatory cytokines/chemokines such as interleukin-1β, interleukin-6, tumor necrosis factor-α, and macrophage colony-stimulating factor in sporadic VS.41–43 A number of safe, effective, and well-tolerated immunomodulatory agents are already in clinical usage against these molecules, and our findings raise the possibility that these agents could be repurposed to help control tumor progression and growth in both VS groups.

**FIG. 5.** Immunostains demonstrating Iba1 and VEGF coexpression. A: Immunostains from a rapidly growing sporadic VS demonstrating macroscopic spatial colocalization between Iba1 and VEGF: upper left = Iba1 (red), immunoperoxidase—whole section; upper right = VEGF (brown), immunoperoxidase—whole section; lower = immunofluorescence demonstrating colocalization between Iba1 (red channel) and VEGF (green channel). B: Higher magnification (×20 HPF) of the area framed in the whole sections (A; immunoperoxidase, ×20) demonstrates a region of high Iba1+ TAM density (left) with corresponding immunopositivity for VEGF (right). C: Higher magnification (×20 HPF) immunofluorescence images of the area framed in the whole sections demonstrates areas of cellular colocalization between Iba1 and VEGF (yellow): DAPI (blue), Iba1 (red), VEGF (green), and Iba1-VEGF (red-green; immunofluorescence, ×20). Figure is available in color online only.
Conclusions

This is the largest comparative study of MRI-derived microstructural and microvascular characteristics in both sporadic and NF2-related VSs and the first such study to compare, in tissue, the relationship between tumor vascularity and inflammatory cell infiltration in both these tumor groups. Our imaging and tissue findings have demonstrated that both sporadic and NF2-related VSs show marked similarities regarding their microvascular metrics, and that in both tumor groups inflammation may be a relevant therapeutic target.

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References


**Disclosures**

Dr. Evans reports being a consultant to AstraZeneca.

**Author Contributions**

Conception and design: Lewis, Jackson, King, Coope. Acquisition of data: Lewis, Donofrio, O’Leary, Williams, Djoukhadar, Agushi, Stapleton, Lloyd, Freeman, Rutherford, Hammerbeck-Ward, Evans, Jackson, Pathmanaban. Analysis and interpretation of data: Lewis, Donofrio, O’Leary, Li, Zhu, Williams, Agushi, Hannan, Roncaroli, Coope. Drafting the article: Lewis, Donofrio, O’Leary, Hannan, Roncaroli, King, Coope. Critically revising the article: Lewis, Donofrio, O’Leary, Li, Zhu, Williams, Hannan, Lloyd, Freeman, Wadeson, Rutherford, Hammerbeck-Ward, Evans, Jackson, Pathmanaban, Roncaroli, King, Coope. Reviewed submitted version of manuscript: all authors. Approved the final version of the manuscript on behalf of all authors: Lewis. Statistical analysis: Lewis, Donofrio, O’Leary. Administrative/technical/material support: Wadeson. Study supervision: Jackson, King, Coope.

**Supplemental Information**

Online-Only Content

Supplemental material is available with the online version of the article. [Supplementary Methods, Tables, and Figures.](https://thejns.org/doi/suppl/10.3171/2020.3.JNS193230).

**Correspondence**

Daniel Lewis: Manchester Centre for Clinical Neurosciences, Salford Royal NHS Foundation Trust, Manchester Academic Health Science Centre, Salford, Greater Manchester, UK. dan.lewis1112@gmail.com.