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Intrinsic Susceptibility MRI Identifies Tumors with $ALK^{F1174L}$ Mutation in Genetically-Engineered Murine Models of High-Risk Neuroblastoma

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Abstract

The early identification of children presenting $ALK^{F1174L}$-mutated neuroblastoma, which are associated with resistance to the promising $ALK$ inhibitor crizotinib and a marked poorer prognosis, has become a clinical priority. In comparing the radiology of the novel Th-$ALK^{F1174L}$/Th-$MYCN$ and the well-established Th-$MYCN$ genetically-engineered murine models of neuroblastoma using MRI, we have identified a marked $ALK^{F1174L}$-driven vascular phenotype. We demonstrate that quantitation of the transverse relaxation rate $R_2^*$ (s$^{-1}$) using intrinsic susceptibility-MRI under baseline conditions and during hyperoxia, can robustly discriminate this differential vascular phenotype, and identify $MYCN$-driven tumors harboring the $ALK^{F1174L}$ mutation with high specificity and selectivity. Intrinsic susceptibility-MRI could thus potentially provide a non-invasive and clinically-exploitable method to help identifying children with $MYCN$-driven neuroblastoma harboring the $ALK^{F1174L}$ mutation at the time of diagnosis.

Introduction

Neuroblastoma arises in the sympathetic nervous system during embryogenesis and is the most common extracranial solid tumor in children [1]. For the established subset of patients presenting with high-risk neuroblastoma, the current portfolio of therapeutic options has limited success, with five-year survival rates rarely exceeding 40%.

The poor clinical outcome and aggressive tumor phenotype of high-risk neuroblastoma strongly correlates with amplification of the proto-oncogene $MYCN$ and enhanced tumor angiogenesis [2]. Recently, mutations in the anaplastic lymphoma kinase ($ALK$) tyrosine kinase gene have been identified in ~8–10% of primary neuroblastoma, leading to constitutive activation of the $ALK$ protein. The most common and potent $ALK$ mutation, $ALK^{F1174L}$, is associated preferentially with $MYCN$ amplification, a markedly poorer prognosis, and confers resistance to the promising $ALK$ inhibitor crizotinib [3–5].

With crizotinib in pediatric phase I clinical trials, and other $ALK$ inhibitor studies in development, a current challenge is to rapidly identify upfront children with high-risk $ALK$ mutated or amplified neuroblastoma who may benefit from or become resistant to $ALK$-targeted therapy.

As most pediatric cancers, including neuroblastoma, originate from only a few genetic anomalies during development, they are amenable to genetically engineered mouse (GEM) modeling approaches. GEM models of neuroblastoma, such as the Th-$MYCN$ murine model, which develop spontaneous tumors mirroring the major pathophysiological, genetic and radiological features of high-risk $MYCN$-amplified childhood neuroblastoma, represent clinically-relevant tools for the study of neuroblastoma biology and response to novel therapeutics [6–8]. The development of GEM models co-expressing $ALK^{F1174L}$ and $MYCN$ to the neural crest, such as the Th-$ALK^{F1174L}$/Th-$MYCN$ model, have recently been used to demonstrate how the $ALK^{F1174L}$ mutation potentiates the oncogenic activity of $MYCN$, and that $ALK^{F1174L}$-acquired resistance to crizotinib can be overcome through inhibition of key cellular modulators of $n$-$myc$ [9–11].

In this study, we hypothesized that the presence of the $ALK^{F1174L}$ mutation results in a phenotypic difference in hemo-
dynamic vasculature in tumors in the Th-ALKF1174L/Th-MYCN model, and which can be evaluated using intrinsic susceptibility magnetic resonance imaging (MRE). The aim of our study was to demonstrate that quantitation of the transverse relaxation rate, \( R_2^* \), and changes in \( R_2^* \) induced by breathing 100% oxygen, \( \Delta R_2^* \) air-oxygen, could discriminate between tumors arising in Th-ALKF1174L/Th-MYCN and Th-MYCN mice, and that intrinsic susceptibility MRE could thus potentially provide a non-invasive and clinically-translatable method to help identify children, presenting MYCN-amplified neuroblastoma harboring the ALKF1174L mutation.

Materials and Methods

Ethics statement

All procedures involving animals were approved by the Institute of Cancer Research Animal Ethics Committee and the UK Home Office and carried out according to the United Kingdom National Cancer Research Institute guidelines for the welfare of animals in cancer research [12].

Animal models

The generation of the Th-ALKF1174L/Th-MYCN mice has been recently described [11]. Th-ALKF1174L/Th-MYCN and Th-MYCN mice were identified by analyzing DNA from mice tails using real-time quantitative reverse transcription polymerase chain reaction (qRT-PCR, Transnetyx Inc., Cordova, Tennessee), and mice with tumors were initially identified by palpation.

Magnetic Resonance Imaging

All the \(^1H\) MRI studies were performed on a 7T Bruker horizontal bore micro-imaging system (Bruker Instruments, Ettlingen, Germany) using a 3 cm birdcage coil. Anesthesia was induced by an intraperitoneal 0.1 ml injection of a combination of fentanyl citrate (0.315 mg/ml) plus fluanisone (10 mg/ml) (Hypnorm, Janssen Pharmaceutical, Oxford, UK) and midazolam (5 mg/ml) (Roche, Welwyn Garden City, UK) and water (1:1:2). A nose piece was positioned for oxygen delivery.

For all the mice, anatomical T2-weighted coronal and transverse images were acquired from twenty contiguous 1 mm-thick slices through the mouse abdomen, using a rapid acquisition with refocused echoes (RARE) sequence with 4 averages of 128 phase encoding steps over a 3 x 3 cm field of view, two echo times (TE) of 36 and 132 ms, a repetition time (TR) of 4.5 s and a RARE factor of 8. These images were used to determine tumor volumes, and for planning the intrinsic-susceptibility MRI measurements, which included optimization of the local field homogeneity. The baseline transverse relaxation rate \( R_2^* \), sensitive to the concentration of paramagnetic species, principally deoxygenemoglobin, was quantified in tumors from Th-ALKF1174L/Th-MYCN (n = 23) and Th-MYCN (n = 21) mice, using a multiple gradient echo (MGE) sequence. MGE images were acquired from three 1 mm thick transverse slices through each tumor, using 8 averages of 128 phase encoding steps over a 3 x 3 cm field of view, and an acquisition time of 3 min 20 s. Images were acquired with 8 echoes spaced 3 ms apart, an initial echo time of 6 ms, a flip angle 45° and a repetition time of 200 ms. Subsequently, 100% oxygen (BOC Ltd, Guildford, UK) was delivered at a rate of 2 L/min to Th-ALKF1174L/Th-MYCN (n = 10) and Th-MYCN (n = 12) mice. After a 5 minutes equilibrium period, identical MGE images were acquired whilst the mouse continued to inhale oxygen.

All the MGE data were fitted voxelwise using in-house software (ImageView, working under IDL, ITT, Boulder, Colorado, USA) with a robust Bayesian approach that provided estimates of \( R_2^* \) and \( \Delta R_2^* \) oxygen-air = \( R_2^* \) oxygen - \( R_2^* \) air [13].

Histological Assessment

Formalin fixed paraffin embedded sections from Th-ALKF1174L/Th-MYCN and Th-MYCN mice were stained with haematoxylin and eosin, and visualized under light microscopy. The extent of functional vasculature in tumors from both models was assessed using the perfusion marker Hoechst 33342, and quantified as fluorescent area fractions (%), as previously described [8].

Statistical analysis

Statistical analysis was performed using GraphPad Prism 5 (GraphPad Software Inc., La Jolla, USA). The mean values for tumor volume, mean of median values for \( R_2^* \) and \( \Delta R_2^* \) oxygen-air and the mean fluorescent area fractions were used for statistical analysis. \( R_2^* \) and \( \Delta R_2^* \) oxygen-air were assumed to be normally distributed, which was confirmed using the D’Agostino-Pearson omnibus K² normality test [14]. Any significant difference in tumor volume, \( R_2^* \), \( \Delta R_2^* \) oxygen-air and the fluorescent area fractions between Th-ALKF1174L/Th-MYCN and Th-MYCN mice were identified using Student’s 2-tailed unpaired t-test, with a 5% level of significance.

Results

ALKF1174L mutation induces a differential radiological presentation of neuroblastoma

Anatomical T2-weighted RARE MR images revealed solid masses within the retroperitoneum in peri-renal and para-spinal abdominal regions of both the Th-ALKF1174L/Th-MYCN and Th-MYCN mice, typical of the clinical distribution and radiological presentation of human neuroblastoma (Figure 1A) [15,16]. All the Th-ALKF1174L/Th-MYCN (n = 23) and Th-MYCN (n = 21) mice examined presented with abdominal tumors with wide range of volumes (525–3400 mm³ for tumors in Th-ALKF1174L/Th-MYCN mice and 335–2450 mm³ for tumors in Th-MYCN mice). Tumors in the Th-MYCN mice appeared heterogeneous, with 94% of tumors presenting with areas of hypointense signal, consistent with previous observations [8]. Tumors in the Th-ALKF1174L/Th-MYCN mice were generally hyperintense and significantly more homogeneous, with only 48% of the tumors having hypointense regions (p = 0.006, \( \chi^2 \)-test). On \( T_2 \)-weighted images (Figure 1B and C, and Figure 2A), tumors in the Th-MYCN mice were generally hypointense at longer echo times, whereas the signal from tumors in the Th-ALKF1174L/Th-MYCN remained hyperintense, providing a stark positive contrast with the surrounding organs for 74% of the Th-ALKF1174L/Th-MYCN mice (compared to 14% for the Th-MYCN mice, p<0.001, \( \chi^2 \)-test).

ALKF1174L mutation results in a slower tumor transverse relaxation rate, \( R_2^* \), in neuroblastoma

Parametric maps revealed a heterogeneous distribution of \( R_2^* \) in tumors in both models (Figure 2B). Quantitation of the transverse relaxation \( R_2^* \) revealed significantly slower rates in tumors in the Th-ALKF1174L/Th-MYCN cohort, compared to the faster \( R_2^* \) rate determined in tumors in the Th-MYCN mice (Figure 2D). Baseline \( R_2^* \) detected tumors harboring the ALKF1174L mutation with a sensitivity of 90% (95% CI: 66.9–98.2) and a specificity of 81% (95% CI: 57.4–93.7).
ALKF1174L mutation is associated with a stark differential BOLD MRI response to hyperoxic challenge in neuroblastoma

Continuous inhalation of 100% oxygen resulted in tumors in the Th-MYCN mice demonstrating a very strong, heterogeneous increase in T2*-weighted image signal intensity (blood oxygen level dependent (BOLD) effect) (Figure 2A). In contrast, tumors from Th-ALKF1174L/Th-MYCN mice showed a negligible BOLD effect on T2* weighted image intensity with hyperoxia. As with baseline R2*, parametric maps revealed a heterogeneous distribution of ΔR2*oxygen-air in tumors in both models (Figure 2B and C).

Tumor regions showing a relatively fast baseline R2* remained so during hyperoxia, whereas regions of relatively slower baseline R2* typically showed a more marked reduction in R2* with 100% O2. The significant difference in R2* between the two models was lost with 100% oxygen challenge (Figure 2E). As a consequence, tumors in Th-ALKF1174L/Th-MYCN mice demonstrated a significantly lower absolute value of ΔR2*oxygen-air than the tumors in Th-MYCN mice (Figure 2F). Hyperoxia-induced ΔR2*oxygen-air detected tumors harboring the ALKF1174L mutation with a sensitivity of 90% (95% CI: 54.1–99.5) and a specificity of 94.1% (95% CI: 69.2–99.7).

Figure 2. Identification of tumors harboring the ALKF1174L mutation in MYCN-driven transgenic mice with intrinsic susceptibility MRI.

A) Anatomical transverse T2*-weighted MR images acquired at TE = 6.8 and 15.6 ms from representative Th-ALKF1174L/Th-MYCN and Th-MYCN mice with abdominal neuroblastoma during initial air breathing, and at TE = 15.6 ms after 5 minutes of continuous inhalation of 100% oxygen. B) Corresponding parametric tumor transverse relaxation rate R2* maps calculated during initial air breathing and after 3 minutes of continuous inhalation of 100% oxygen. C) Resulting parametric tumor ΔR2*oxygen-air (R2*oxygen − R2*air) maps. D) Tumor R2* during initial air breathing. E) tumor R2* after 5 minutes of breathing 100% oxygen, and F) tumor ΔR2*oxygen-air (R2*oxygen − R2*air) were determined from Th-ALKF1174L/Th-MYCN and Th-MYCN mice with abdominal neuroblastoma. Individual data points represent the mean of the median values determined from all three imaging slices for each animal, as well as the mean ±1 s.e.m, p, Student’s 2-tailed unpaired t-test with a 5% level of significance.
No correlation between tumor $R_2^*$ and $\Delta R_2^*$, or between either $R_2^*$ or $\Delta R_2^*$ and tumor volume, was determined across the cohorts, indicating that the differences in intrinsic susceptibility MRI between tumors in the Th-ALKF1174L/Th-MYCN and Th-MYCN mice were independent of tumor size.

**ALKF1174L** mutation is associated with reduced functional vasculature in neuroblastoma

Gross examination of the tumors in situ, prior to excision, revealed an intense dark red coloration across the whole tumor in the Th-MYCN mice, in contrast to the pale appearance of tumors in the Th-ALKF1174L/Th-MYCN GEM model (Figure 3A). Histological examination with H&E staining revealed the presence of large hemorrhagic regions filled with stacked erythrocytes in 100% of the tumors from Th-MYCN mice (n = 10), but only in 10% the Th-ALKF1174L/Th-MYCN model (n = 10, p < 0.0001, $\chi^2$-test) (Figure 3B and C). Fluorescence microscopy of Hoechst 33342 uptake revealed homogeneously and well-vascularized tumors in both models (Figure 3D). However, quantitation of the fluorescent area fractions revealed that tumors in the Th-MYCN mice had significantly higher uptake, consistent with the presence of more perfused, functional vasculature (Figure 3E).

**Discussion**

In the current study we demonstrate that baseline $R_2^*$ and hyperoxia-induced $R_2^*$ can discriminate a differential hemodynamic tumor vascular phenotype between tumors arising in Th-ALKF1174L/Th-MYCN and Th-MYCN models of high-risk, MYCN over-expressing neuroblastoma. With a sensitivity of 90% and a specificity of 81% for baseline $R_2^*$, and a sensitivity of 90% and a specificity of 94% for 100% oxygen-induced $R_2^*$, intrinsic susceptibility MRI provides a robust method to discriminate and identify Th-MYCN transgenic mice harboring the ALKF1174L mutation.

The rapid and completely noninvasive quantitation of tumor $R_2^*$ is suitable for the scanning of young children. Furthermore, such pediatric imaging sessions are usually performed under general anesthesia, providing an opportunity to transiently perturb the oxygen content inhaled by the patient, and enabling quantification of tumor $R_2^*$ under more hyperoxic conditions (from 21 to 30–40% O$_2$). Upon successful translation, intrinsic susceptibility MRI could provide a rapid method to identify children with ALK-driven tumors, enabling the stratification of children with this ultra-high-risk neuroblastoma at the time of diagnosis.

The baseline transverse relaxation rate $R_2^*$ is sensitive to the concentration of paramagnetic deoxyhemoglobin in the vascular compartment in tissue (BOLD effect). Compared to most normal tissues, tumors exhibit relatively fast native $R_2^*$ values, a consequence of the high concentration of deoxygenated erythrocytes within the vascular compartment associated with immature, irregular and unstable microcirculation [17]. The fast baseline $R_2^*$ measured in the Th-MYCN model is consistent with the aggregation of deoxygenated erythrocytes, described as blood lakes [18], and which are characteristic of childhood neuroblastoma [19]. The significantly slower baseline $R_2^*$ and differential intrinsic susceptibility MRI presentation of tumors in the Th-ALKF1174L/Th-MYCN mice is consistent with the absence of such blood lakes.

Inhalation of high oxygen content gases results in the rapid re-oxygenation of hemoglobin, with paramagnetic deoxyhemoglobin within perfused tumor vessels being replaced by diamagnetic oxyhemoglobin, and a reduction in $R_2^*$ [20]. Hyperoxia resulted in an overall significant reduction in $R_2^*$ of tumors in the Th-MYCN mice, whereas the response in tumors in the Th-ALKF1174L/Th-MYCN mice was negligible, suggesting a clear difference in hemodynamic functional vasculature between the two GEM models. This was corroborated by the significant overall difference in Hoechst 33342 uptake. Interestingly, a spatially different hyperoxia $\Delta R_2^*$ response was apparent in tumors in the Th-MYCN mice. Tumor regions exhibiting relatively fast baseline $R_2^*$ showed a less pronounced response with 100% O$_2$ breathing, also consistent with the presence of blood lakes which are typically disconnected from the perfused vascular network [21]. In contrast, tumor regions with relatively slower baseline $R_2^*$ showed a less pronounced response to hyperoxia, indicative of functional vasculature.

Intriguingly, tumors from the Th-ALKF1174L/Th-MYCN mice showed no clear $\Delta R_2^*$ response to hyperoxia, despite Hoechst uptake significantly lower in the Th-ALKF1174L/Th-MYCN mice compared to the Th-MYCN mice (n = 5). Data are mean ± 1 s.e.m, p, Student’s 2-tailed unpaired t-test with a 5% level of significance.

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**Figure 3. Pathological comparison of tumors from Th-ALKF1174L/Th-MYCN and Th-MYCN mice with abdominal neuroblastoma.** A) Gross pathology, B) composite images, and C) high magnification (x200) images from hematoxylin and eosin stained sections. Note the presence of large hemorrhagic regions filled with aggregated erythrocytes (‘blood lakes) extravasated from blood vessels (arrowed) in the tumor from the Th-MYCN mouse. D) Composite fluorescence images of uptake of the perfusion marker Hoechst 33342 into tumors from Th-ALKF1174L/Th-MYCN and Th-MYCN mice with abdominal neuroblastoma. E) Quantitation of Hoechst 33342 uptake revealed significantly lower functionally perfused vasculature in tumors of Th-ALKF1174L/Th-MYCN mice (n = 5) compared with tumors in Th-MYCN mice (n = 5). Data are mean ± 1 s.e.m, p, Student's 2-tailed unpaired t-test with a 5% level of significance.
33342 uptake indicating the presence of perfused blood vessels in both GEM models. This suggests a difference in vascular architecture that precludes erythrocyte delivery, but not plasma perfusion, in the tumor vessels in the Th-ALKF1174L/Th-MYCN mice, or in oxygen consumption by the tumor cells [17,22]. The extensive hemorrhage and blood lakes present in the Th-MYCN model are indicative of vessel wall instability due to rapid endothelial cell proliferation and defective pericyte coverage [23]. This implies that the tumor vasculature in the Th-ALKF1174L/Th-MYCN mice may be more mature.

Given its relationship to blood oxygen saturation and partial pressure of oxygen in and around blood vessels, quantitation of baseline tumor R2* is also being investigated as an imaging biomarker of hypoxia [24]. For example, slow baseline R2* has been shown to correlate with increased hypoxia, determined by pimonidazole staining, in chemically-induced rat mammary tumors [25]. Conversely, fast baseline R2* is associated with hypoxia in prostate cancer [26]. In this context, we have recently shown that neuroblastoma from Th-MYCN mice exhibiting fast baseline R2*, induced by a hemorrhagic phenotype, are relatively oxic, as revealed by negligible pimonidazole staining [8]. R2* is first of all a marker of impaired hemodynamic function and as such shares a causal relationship to hypoxia. However this relationship is ambivalent as several differential hemodynamic phenotypes can in fact lead to hypoxia, while having an opposite effect on R2* values. In this study the suggested tumor vascular phenotype in the Th-ALKF1174L/Th-MYCN model, which hinders the delivery of erythrocytes and causes a significantly slower R2*, may result in increased tumor hypoxia in the Th-ALKF1174L/Th-MYCN model compared with the Th-MYCN model.

Recent reports have implicated a role of the ALKF1174L mutation in tumor angiogenesis in neuroblastoma. Selective targeted inhibition of ALK resulted in a significant reduction in vascular density in xenografts derived from MYCN non-amplified SH-SY5Y cells, which harbor the ALKF1174L mutation, accompanied with a decrease in vascular endothelial growth factor (VEGF) and matrix metalloproteinases (MMPs) [27]. The expression of MMP-9 by stromal cells has previously been shown to regulate the vascular architecture in a murine orthotopic MYCN amplified neuroblastoma xenograft model by promoting pericyte recruitment [26]. Collectively, these studies suggest that ALK and the ALKF1174L mutation contribute to tumor vasculogenesis and pericyte recruitment via the regulation of VEGF and MMP-9, leading to a dense vascular network with smaller, more stable vessels. The differential intrinsic susceptibility MRI vascular phenotype observed in tumors in the Th-MYCN and Th-ALKF1174L/Th-MYCN models reported herein demonstrates an important role for ALKF1174L in angiogenesis in MYCN-overexpressing neuroblastoma in vivo, strongly suggesting a role in vasculogenesis.

This study has some limitations. The Th-ALKF1174L mutation leads to the constitutive activation of the ALK protein in neuroblastoma [29–32], and the vascular phenotype associated with amplified or wild-type (WT) ALK expression is currently unknown. Amplification of ALK has also been shown to lead to activation of the ALK protein with, however, a 17-fold reduced kinase activity compared to ALKF1174L mutants [29]. In the absence of a Th-ALKF1174L GEM model, it is difficult to conclude if intrinsic susceptibility MRI would be solely able to identify ALKF1174L-mutated MYCN amplified neuroblastoma, or more generally ALK-driven MYCN amplified neuroblastoma.

There is a clear need to more deeply interrogate the role of the ALKF1174L mutation on vascular morphogenesis and architecture, which may be a major determinant of impaired drug delivery and a contributing factor to the poor prognosis of children with ALKF1174L-mutated MYCN-amplified neuroblastoma. The altered vascular phenotype may also impact on the response to anti-vascular therapies, including cediranib, currently being considered in clinical trials for the treatment of high-risk MYCN-amplified neuroblastoma [33]. Retrospective analysis of historical pathological samples for ALK mutations should provide sufficient statistical power to shed light on the vascular phenotype of neuroblastoma associated with ALK amplification, and each of the rare ALK mutations, including the lethal ALKF1174L mutation.

Coupled with our recent identification of quantitation of R2* as a biomarker of treatment response to cediranib in the Th-MYCN GEM model [9], the present study reinforces R2* as a biomarker of vasculogenesis and its response to therapy in these clinically relevant GEM models. Furthermore, it provides a strong rationale for the evaluation of intrinsic susceptibility MRI for assessing any anti-angiogenic effects resulting from successful targeted inhibition of ALK signalling in ALKF1174L-mutated neuroblastoma, which could ultimately used for the assessment of second generation ALK inhibitors [34,35]. To conclude, this study has provided a strong rationale for the immediate incorporation of intrinsic susceptibility MRI into forthcoming imaging-embedded clinical trials of next generation ALK inhibitors in ALK-mutated and -amplified neuroblastoma.

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Author Contributions

Conceived and designed the experiments: VJ LG RG LC SR. Performed the experiments: VJ LG AH. Analyzed the data: VJ LG AH DMK ADJP LC SR. Wrote the paper: VJ LG AH RG DMK ADJP LC SR.

References


