Role of diffusional kurtosis imaging in grading of brain gliomas: a protocol for systematic review and meta-analysis

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ABSTRACT

Introduction Central nervous system (CNS) gliomas are the most common primary intra-axial brain tumours and pose variable treatment response according to their grade, therefore, precise staging is mandatory. Histopathological analysis of surgical tumour samples is still deemed as the state-of-the-art staging technique for gliomas due to the moderate specificity of the available non-invasive imaging modalities. A recently evolved analysis of the tissue water diffusion properties, known as diffusional kurtosis imaging (DKI), is a dimensionless metric, which quantifies water molecules’ degree of non-Gaussian diffusion, hence reflects tissue microenvironment’s complexity by means of non-invasive diffusion-weighted MRI acquisitions. The objective of this systematic review and meta-analysis is to explore the performance of DKI in the presurgical grading of gliomas, both regarding the differentiation between high-grade and low-grade gliomas as well as the discrimination between gliomas and other intra-axial brain tumours.

Methods and analysis We will search PubMed, Medline via Ovid, Embase and Scopus in July 2018 for research studies published between January 1990 and June 2018 with no language restrictions, which have reported on the performance of DKI in diagnosing CNS gliomas. Robust inclusion/exclusion criteria will be applied for selection of eligible articles. Two authors will separately perform quality assessment according to the quality assessment of diagnostic accuracy studies-2 tool. Data will be extracted in a predesigned spreadsheet. A meta-analysis will be held using a random-effects model if substantial statistical heterogeneity is expected. The heterogeneity of studies will be evaluated, and sensitivity analyses will be conducted according to individual study quality.

Ethics and dissemination This work will be based on published studies; hence, it does not require institutional review board approval or ethics clearance. The results will be published in peer-reviewed journals.

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INTRODUCTION

Gliomas are the most common primary intra-axial brain tumours. Precise and robust diagnosis and staging are crucial for implementing successful management strategies and predicting their outcome. However, there is noticeable variation in individual tumour response even if they are attributed the same histological grade. This could be explained by the divergent proliferative patterns, cellular complexity and neovascularisation among these tumours. WHO 2007 classification used to classify brain gliomas into four subgroups depending on their histopathological features such as cellular atypia, mitotic activity and anaplasia. Meanwhile, efforts have been made in order to further stratify gliomas based on their molecular features aiming to better explain the tumours’ behaviour and determine better treatment strategies. The revised fourth edition of the updated WHO classification of central nervous system tumours in 2016 incorporates both molecular and histopathological features of gliomas. As previously, histopathological examination after neurosurgical sampling is still considered as the state-of-the-art staging method of gliomas; owing to the
evolved to quantify water molecules’ degree of non-Gaussian imaging (DKI) is a dimensionless metric that has recently emerged as presurgical imaging tools and as tools for evaluating treatment response to chemotherapy and radiotherapy regimens in patients with gliomas. However, DWI and DTI quantify water molecules’ mobility on the assumption of unrestricted—but possibly hindered—random diffusion. Accordingly, the likelihood of certain proton diffusing from one location to another in a given time (known as the probability distribution function, PDF) is thought to be Gaussian related. However, the paramount differences inside the brain tissue cytoarchitecture due to cell membranes, organelles and discrete compartments will drift the diffusion of water molecules from the normal Gaussian distribution. Therefore, the real PDF will be more soared up in contrast to the Gaussian PDF. The widely used metric of diffusion attenuation in tumours, namely the apparent diffusion coefficient, is limited in detecting this deviation from the normal Gaussian behaviour. A novel diffusion model known as diffusional kurtosis imaging (DKI) is a dimensionless metric that has recently evolved to quantify water molecules’ degree of non-Gaussianity. Thus, DKI can provide a more realistic biomarker that reflects the brain microenvironment’s complexity. It is simply considered a continuation of the DTI model and at least 2 non-zero diffusion gradient factors (b-values) in more than 15 non-linear diffusion directions are applied to acquire both the kurtosis metrics (radial kurtosis, axial kurtosis and mean kurtosis (Mk)) and the diffusion tensor metrics (mean diffusivity and fractional anisotropy simultaneously). It is therefore important to understand the diagnostic performance of DKI, both regarding its ability to stage the gliomas in subgroups and to differentiate gliomas from other intra-axial brain tumours.

Recent studies have shown promising results. In April 2018, a systematic review and meta-analysis of the diagnostic accuracy of DKI for glioma grading revealed that the pooled area under the curve for Mk in differentiating high-grade from low-grade gliomas was 0.94. This review included only 10 studies, whereas in the current work, we believe that more studies about DKI have been held in the interim. Moreover, we aim to include studies that compared DKI between gliomas and other intra-axial brain tumours.

**OBJECTIVE**

This review aims to investigate the diagnostic performance of DKI in the grading of gliomas and the differentiation between gliomas and other intra-axial brain tumours.

**REVIEW QUESTIONS**

How accurate is DKI in differentiating between high-grade and low-grade gliomas and in differentiating brain gliomas from other intra-axial tumours?

**METHODS**

This review protocol was developed according to the Preferred Reporting Items for Systematic Reviews and Meta-Analyses-Protocols (PRISMA) statement and guidance from the Joanna Briggs Institute Reviewers’ Manual for the systematic review of studies of diagnostic test accuracy. The subsequent full systematic review will be prepared according to the PRISMA-Diagnostic Test Accuracy Studies checklist.

**Patient and public involvement**

This work will be based on published studies; hence, there will not be patient or public involvement. The results will be published in peer-reviewed journals.

**Inclusion and exclusion criteria**

We will include all studies that have investigated DKI in tumours of glial cell origin, either primary or recurrent gliomas, in adult patients, using either WHO 2007 or WHO 2016 classifications and compared the DKI features in gliomas with other non-glial tumours. Exclusion criteria will comprise paediatric groups, non-Original research articles (reviews, commentaries, erratum, books, editorial and conference abstracts), animal studies, non-imaging studies, non-MRI studies, non-DKI
DWI-MRI studies, non-neoplastic conditions, non-glial tumours only, non-cerebral tumours and studies written in languages other than English, French or German.

Search strategy for identification of studies
A systematic literature search will be developed in July 2018 in four databases: PubMed, Medline via Ovid, Scopus and Embase using the keywords ‘glioma’ and ‘diffusional kurtosis’ by both medical subject heading and text words, without language restrictions. The suggested search syntax on PubMed and Medline is summarised in table 1 and the flow chart is illustrated in figure 1.

Study selection for inclusion in the review
Initially, studies identified by the literature search will be independently screened for primary eligibility by two authors, based on title and abstract. Full text of the primarily eligible studies will be independently assessed for final inclusion in the systematic review and meta-analysis by the same raters. Any disagreements will be resolved through discussions. Reasons for study exclusion will be stated clearly.

QUALITY ASSESSMENT
Eligible studies will be independently assessed according to the revised tool of quality assessment of diagnostic accuracy studies-2 tool14 by two coauthors. Any disagreements will be resolved by consensus. The tool comprises four key domains which are: patient selection, index test, reference standard and patients’ flow in the study and timing of the reference standard and the index test. Under each domain, risk of bias will be assessed through answers to signalling questions given in the tool, taking into account the review questions. Regarding the patient selection domain, retrospective studies will be considered to have high risk of bias, meanwhile low risk will be attributed to prospective studies. Regarding the index test domain, whether the neuroradiologist was blinded to the pathology during the image processing and region of interest drawing or not will determine the risk of bias; the neuroradiologists being blinded indicates low risk of bias. Histopathological results will serve as the reference standard. In the domain of flow and timing, unclear risk of bias will be ascertained if the interval between the

Figure 1 Flow diagram for search strategy. CNS, central nervous system; DKI, diffusional kurtosis imaging.
index test and the reference standard is not mentioned in the study. Additionally, unclearness will be considered also if any patient was excluded from the analysis without relevant reasons. Concerns regarding applicability will be assessed in the first three domains only.

**DATA EXTRACTION AND MANAGEMENT**

Data extraction will be performed by two coauthors in a predesigned standardised sheet. Extracted data will include the following: first author name, publication year, type of study, details about the patient population, data acquisition techniques, image processing, postprocessing software, reference standard and diagnostic test accuracy results (true-positive, false-positive, true-negative and false-negative values). Any missing data will be requested from the related study authors.

**DATA SYNTHESIS AND ANALYSES**

We will construct 2×2 tables using reported number of true-positive, true-negative, false-positive and false-negative cases to calculate different indicators of diagnostic performance. A narrative synthesis will summarise the available evidence. Paired forest plots illustrating sensitivity and specificity with their 95% CIs will be built using the Review Manager software. We will use a bivariate random-effect model for data synthesis based on the assumption that studies are of sufficient heterogeneity in terms of study populations and study methodology. We will also report the number of true positives, false positives, true negatives and false negatives. A random-effect meta-analysis as well as aggregation of data using the hierarchical summary receiver operator characteristics will be pursued.

**REFERENCES**