Presurgical Mapping of Brain Tumors Using Statistical Probabilistic Anatomical Maps

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Abstract

Purpose: Multi-tracer neuroimaging is widely used for the diagnosis, radiotherapy, and neurosurgery of brain gliomas. Anatomical and functional information is important to avoid having neurological deficits induced during the resection or radiation therapy of brain gliomas. The aim of this study was to investigate presurgical anatomical labeling of brain gliomas on FLT and FET using statistical probabilistic anatomic maps (SPAM), which are images of cerebral cortical, cerebellar, and subcortical volumes of interest (VOIs). Methods: FDG, FLT, and FET PET scans were acquired. FLT and FET PET images were coregistered to the FDG PET images, which were then spatially normalized onto the target brain. An inverse spatial normalization parameter was calculated and applied to SPAM. For the anatomical labeling of brain glioma regions, the volumes of brain gliomason FLT and FET images were extracted using segmentation. Probabilistic information of the glioma region was then calculated using SPAM and the segmented glioma volumes. SPM and an in-house program were used for image processing. Results: The probability of SPAM labeling a brain glioma region could be extracted using the inverse normalized SPAM and segmented glioma regions. In a sample case, the probabilistic anatomical region of the glioma included 21% of the postcentral gyrus, 12% of the superior parietal gyrus, and 6% of the angular gyrus. Conclusion: Anatomical information about brain gliomas could be extracted using SPAM. This proposed method would be optional for presurgical mapping, to avoid an additional functional mapping study that might otherwise be necessary to avoid producing neurological deficits.

Keywords

SPAM, Presurgical Mapping, Brain Tumor

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1. Introduction

Brain tumors are the most common solid tumors, and are typically evaluated by $^{18}$F FDG PET (FDG) or by combined FDG and MR imaging [1]. The hybrid PET/MR allows for molecular, anatomical, and functional imaging such as ASL, proton-spectroscopy, $^{11}$C methionine PET, and $^{68}$Ga DOTATOC PET [2]. $^{3'}$-deoxy-$^{3'}$-$[^{18}$F] fluorothymidine (FLT) PET has been found useful for the noninvasive grading of gliomas, their prognostic assessment, and evaluation of response to malignant glioma treatments [3]. O-$(2-[^{18}$F]fluoroethyl)-L-tyrosine (FET) has gained ground for the diagnostic work-up of brain tumors, because it overcomes the limitations of FDG PET by providing a much higher tumor to non-tumor contrast [4].

Presurgical evaluation of language in brain tumor patients is important. For example, functional magnetic resonance imaging is used to assess laterality in preoperative brain tumor cases [5]-[7]. In addition, navigated transcranial magnetic stimulation (nTMS) is increasingly used in presurgical brain mapping. Preoperative nTMS results correlate well with direct cortical stimulation (DCS) data in the identification of the primary motor cortex [8].

A statistical probabilistic anatomic map (SPAM) is an atlas-based volume of interest (VOI) that is used to obtain the regional counts from spatially normalized individual images into standardized brain templates. For example, the SPAM images of the Montreal Neurological Institute, consisting of 98 VOIs, were used to objectively calculate the PET or SPECT counts in neocortical, cerebellar, and subcortical VOIs. Each image consisted of probabilities belonging to specific regions. The SPAM probability-weighted mean counts of frontal cortex, temporal cortex, parietal cortex, occipital cortex, whole cerebral cortex, cerebellar cortex, basal ganglia, and thalamus for each hemisphere were obtained from the 98VOIs using the probabilistic brain atlas. Counts were calculated by dividing the probability-weighted sum of all voxels in the VOI by the sum of the probability sum of the VOI and the number of voxels [9] [10]. SPAM was used to quantify brain perfusion SPECT [10], surgical prognosis of temporal lobe epilepsy [11], and perfusion and glucose metabolism of epileptogenic zones in temporal lobe epilepsy [12].

In this present study, we developed a SPAM-assisted presurgical evaluation of brain tumors using the inverse spatial normalization mapping technique provided in the SPM software’s toolbox. Our method as developed herein could provide functional mapping without the need for any additional data acquisition by imaging techniques such as fMRI, nTMS, or DCS.

2. Material and Methods

2.1. Patients

The method developed for this study was retrospectively applied to already acquired PET data in our institute [3]. The patients who were previously diagnosed with glioma and who were treated with multiple therapies (such as operation, chemotherapy and radiotherapy) within a 7-year period from 2003 to 2009. Our previous study was approved by the institutional review board of the Korean Cancer Center Hospital, and written informed consent was obtained from patients. Patients (13 men and 7 women, age range: 31 - 72 years, mean age: 51.7 ± 12.6 years) with newly diagnosed gliomas who underwent MRI, FLT, and FET PET were included in our study. All patients had been referred to our institute for initial diagnosis. A pathological diagnosis was obtained by stereotactic biopsy or open surgery. Grading of the tumor was performed according to the World Health Organization (WHO) grading system for gliomas. Three patients had grade II, two had grade III, and fifteen had grade IV gliomas.

2.2. PET Study

Briefly, the PET acquisition process was described as follows: Twenty patients underwent FDG, FLT, and FET PET within 3 days. The images of the brain were obtained thirty minutes after intravenous injection of approximately 370 MBq of FDG, FLT, or FET. An ECAT HR+ PET scanner (Siemens Medical System, Knoxville, TN) was used. A 20-min emission scan (128 $\times$ 128 matrix) and a 10-min post-emission transmission scan using a $^{68}$Ge source were performed. The attenuation-corrected images were reconstructed with an ordered subset expectation maximization (OSEM) algorithm (2 iterations and 16 subsets) with segmented attenuation correction.

2.3. Coregistration

In this study, performance of 4 registration methods including mutual information (MI), normalized mutual
2.4. Glioma Region Mapping Using Statistical Probabilistic Anatomic Map

In this study, we used the Korean version of SPAM, which was developed by the Seoul National University group [13]. SPAM images of the Seoul National University consisting of 152 VOIs were used to objectively calculate the PET counts in the neocortical, cerebellar, and subcortical VOIs. Each image consisted of the probabilities that belonged to its specific regions.

Individual FET PET or FLT PET data was coregistered to FDG PET data using the mutual information method provided in the SPM 8 software. The FDG PET image was then spatially normalized onto Korean standard templates [13] by both linear and nonlinear transformations. The inverse spatial normalization parameter was calculated and applied to SPAM. For the anatomical labeling of brain glioma regions, the volume of brain gliomas on FLT and FET images was extracted using segmentation. The probabilistic information of the glioma region was then calculated using SPAM and the segmented glioma volume.

3. Results

Figure 1 shows the result of co-registration. Image registration of FDG & FLT, FDG & FET, and FDG & MR was successfully performed using MI, NMI, and ECC methods. However, image registration of FDG & FLT and FDG & MR could not be performed using NCC method. Yellow arrow indicates the mismatch of coregistration.

Figure 2(a) shows the individual FDG PET, FLT PET, and FET PET scans. Figure 2(b) demonstrates the
coregistration of FDG PET and FET PET, and of FDG PET and FLT PET data.  

Figure 3 shows the preprocessing step for anatomical mapping using SPAM. The FDG PET image was spatially normalized onto the target brain and the inverse spatial normalization parameter was calculated. Inverse spatial normalization was applied to SPAM onto coregistered FDG PET, FLT PET, and FET PET scans.

Figure 4 shows the application of inverse normalized SPAM in regions such as the postcentral gyrus, precentral gyrus, and superior parietal gyrus onto FLT PET imaging data. The probability of a SPAM-labeled brain glioma region could be extracted using inverse normalized SPAM and the segmented glioma region. In the sample case, the probabilistic anatomical region of the glioma included 8.9% of the precentral gyrus, 36.8% of the postcentral gyrus, 21.7% of the superior parietal gyrus, 10.9% of the supramarginal gyrus, 6.6% of the angular gyrus, and 14.8% of Brodmann’s area 44.
4. Discussion

In our present study, we developed a SPAM-assisted presurgical evaluation of brain tumors using an image processing step involving SPM, but without any additional presurgical data acquisition. For the successful implementation of inverse spatial normalization, spatial normalization onto the template is important [14]. In our present study, we applied the SPAM-assisted presurgical evaluation of brain tumors to FLT PET data because FLT PET data lack specific anatomical information. Due to high contrast in the tumor to non-tumor ratio, we used FDG PET for spatial normalization. The FLT image was coregistered to an FDG PET image. In our previous study, we compared 4 different coregistration methods: Mutual information, normalized mutual information, normalized cross correlation, and entropy correlation coefficient. We found that image registration of FDG and FLT, FDG and PET, and FDG and MR was successfully performed using mutual information, normalized mutual information, and entropy correlation coefficient methods, but not via the normalized cross correlation method [15]. We found that inverse spatial normalization of SPAM onto coregistered FLT PET information could provide enhanced anatomical information on the native spatial space of FLT PET data.

Many studies on the feasibility of fMRI in presurgical mapping for brain tumors have been introduced [5]-[7][16]. Because the fMRI study of cancer patients is difficult, SPAM-assisted presurgical mapping for brain tumor patients would be a more feasible option for clinics.

Presurgical mapping is also important for epilepsy surgery evaluations in which the epileptic focus is to be located. Typically, the presurgical evaluation was performed using MRI, SPECT, PET, fMRI, or MEG [17]-[19]. Although developed in brain glioma cases, our SPAM-assisted presurgical mapping technique could apply to presurgical mapping for epilepsy surgery as well.

Taken together, the data in the present study demonstrate the feasibility of SPAM-assisted presurgical mapping for brain tumors. However, because there would be individual differences in brain function, and anatomical changes due to the tumor’s volume, our SPAM-assisted mapping method could best be used as an instant mapping method just prior to an fMRI study, or in cases when presurgical mapping using fMRI or other methods is not available.

Our proposed SPAM-assisted presurgical mapping could provide efficient tool for avoiding of producing neurological deficits without additional acquisition of functional MR. In addition, our proposed SPAM-assisted presurgical mapping method would be beneficial when there was a lack of anatomical information.

5. Conclusion

Anatomical information about brain gliomas could be successfully extracted using SPAM. This proposed method would be an option for presurgical mapping, that could help to prevent induction of neurological deficits during surgery without the need for additional functional mapping studies.

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References


