

IMAge/enGINE: a freely available software for rapid computation of high-dimensional quantification

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Background: High-dimensional image data including diffusion weighted imaging, diffusion tensor imaging and dynamic imaging are important in exploring the connectivity, cellularity, pharmacokinetic and blood supply. IMAge/enGINE is software especially designed for high-dimensional medical image computing.

Methods: IMAge/enGINE is implemented based on open-source and cross-platform tools such as Qt, ITK and VTK. It processes the high-dimensional image data in a slice-by-slice computation mechanism. For computational efficiency, C++ is used for implementing IMAge/enGINE and multi-thread computing is handled in the scale of voxels. The architecture of IMAge/enGINE is modularized for easier extension.

Results: IMAge/enGINE has following features: (I) IMAge/enGINE is free for research use; (II) it has an easy-to-use graphic user interface designed for clinical users without programming or engineering background; (III) its frame work is open-source and extensible. Developers can implement algorithms as modules and integrate them into IMAge/enGINE or generate their own application.

Conclusions: The source of IMAge/enGINE is hosted at https://github.com/VusionMed/IMAgeenGINE. Multiple diffusion and perfusion models are implemented and integrated into IMAge/enGINE and its binaries can be downloaded freely at http://www.vusion.com.cn/?page_id=14971.

Keywords: Diffusion weighted imaging; medical image computing; dynamic contrast-enhanced MRI

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Introduction

Routine modalities of MR imaging such as diffusion weighted imaging (DWI) and dynamic contrast-enhanced MRI (DCE-MRI) generate high dimensional image data which require processing at a further step for interpreting.

DWI enables the identification of the structure and organization of tissue at the microscopic level (1,2). And it has been widely performed on the brain, prostate, liver, or kidney (3,4). DWI enables converting MR imaging into lots of quantified models, such as apparent diffusion coefficient (ADC) (5), intra-voxel incoherent motion (IVIM) (6), diffusional kurtosis imaging (DKI) (7) and neurite orientation dispersion and density imaging (NODDI) (8). The response of tumors (9,10) and brain development can be assessed by these quantified models (8).

Dynamic susceptibility enhanced (DSC) (11) and dynamic contrast enhanced (DCE) (10) MRI are often used to identify the pathological tissues by dynamically monitoring the change in image after using the contrast agent (12). Contrast agents, like gadolinium, with good detective sensitivity can be detected with routine acquisition protocols. Blood flow, permeability and other parameters could be computed by kinetic analysis of these high dimensional data.

Indeed, the application of these techniques holds great

potential to improve clinical outcomes and to potentially guide patient managements. However, with several challenges such as technical complexity and methodological challenges, it is challenging for a researcher with clinical background to compute these quantified datasets. In order to overcome this limitation, several integrated software such as commercial workstations, OsiriX (13) and ImageJ (14) are designed to view and process medical image. The problem is a trade-off between being easy to use and extensive for the software.

IMAge/enGINE is free and extensible software for highdimensional medical image computing. Comparing to other free software, IMAge/enGINE is equipped with a graphic user interface which is designed specifically for users without engineering or programming skills. Most of the operations are simplified as clicking buttons and dragging sliders. In addition, IMAge/enGINE offers an open-source and extensible framework. Developers with proper C++ programming skills can integrate and implement their own algorithms into an application.

IMAge/enGINE was released in 2017 on a freely available basis. Since then it has been downloaded by more than 300 unique users from approximately 100 different hospitals and institutions.

Methods

IMAge/enGINE is integrated out-of-box software with easy-to-use graphic user interface. Its binary package of WINDOW platform can be downloaded freely at http:// www.vusion.com.cn/?page_id=14971. The usage for research only will be verified during a simple sign-up process. Afterwards, an account will be created for logging into this software and accessing most of its functions. The framework part of IMAge/enGINE is open-source and archived at https://github.com/VusionMed/IMAgeenGINE.

We chose C++ as the primary language for implementing IMAge/enGINE to integrate with the existing open source libraries and tools. IMAge/enGINE was built upon Qt (15), DICOM toolkit (DCMTK) (16), visualization toolkit (VTK) (17), insight toolkit (ITK) (18), common toolkit (CTK) (19), CMake (20) and TEEM (21). Among them, Qt provides a cross-platform GUI front. DICOM toolkit is used for reading and writing DICOM objects. The DICOM widget of CTK is used for managing the DICOM files. ITK is used for image registration, segmentation and implementing algorithms of computing blocks. VTK is

used for image visualization and interaction. TEEM is used for providing support for files with a suffix of *nrrd*.

Overview

IMAge/enGINE consists of 7 main functional blocks and three of them contain calculation modules. An overview of how these blocks communicate and relate to each other is given in *Figure 1*. Though the calculation modules are not included in the source code, a developer can implement his own calculation module by inheriting the sample class.

DICOM import

IMAge/enGINE employs DICOM browser widget in CTK to manage the DICOM database. User can select and load DICOM data. After data loading, the DICOM data will be read into memory as a VTKImageData with 4 dimensions, X-Y-Z and components (C). The C dimension is used for indicating the b-value and diffusion direction for Diffusion data and time frame for dynamic data. Because most of the MRI scanner manufactures keep the b-value information into the private tag of DICOM metadata, it is difficult to generally support the DICOM from all manufactures. For GE, Philips and Siemens DICOM data, current DicomHelper class can parse the b-value and gradient direction correctly. But for other manufactures data, users have to provide a .bvec and a .bval text files as defined by Chris Rorden's DCM2Nii tool (22) to describe the b-value and gradient direction of loading data. After the data are correctly loaded, users can apply rigid motion correction on the C dimension.

Display

There are two types of window widgets to render the image. Both window widgets can render two layers of images. One is the background image layer which is used for displaying the medical imaging and the other one is the label image layer which is used for drawing and displaying the labelling region of interest. The label image layer is also stored in memory using VTKImageData class and has the same matrix size with its corresponding background image. The background image layer of MasterWindowWidget holds the loaded images whose dimension is 3D+C. The background image layer of SlaveWindowWidget is only used for holding the computed image such as ADC map, IVIM D-star Map which is 2D image of the current slice



Figure 1 Overview of main modules of IMAge/enGINE.

displaying in MasterWindowWidget. Similarly, the label layer of MasterWindowWidget holds 3D image and SlaveWindowWidget only holds 2D image. There is only one MasterWindowWidget, while SlaveWindowWidget can be as many as the available calculation types of the loaded images.

Calculation

When the users trigger an update event such as adding a new type of computed image, scrolling a new slice or applying a new mask, a QT signal is sent to calculation module with the slice number of currently displaying in the MasterWindowWidget. Then the calculation module responds by calculating the required model to generate the computed images (2D) including an R^2 map which indicates the fitting error of each pixel from the informed slice. After the computed images are generated, they will be sent to the Display Core to either update the current SlaveWindowWidget or add a new SlaveWindowWidget. In our design, the calculation will not be performed for slices that are not displayed, and hence largely reduce the computation time. As the only exception, all slices will be calculated automatically when users save the results to local disk. This design is one of the major highlights of IMAge/ enGINE.

Fitting algorithms

Linear

One of the most used functions in calculation mode is fitting data to linear models, such as ADC, DTI/DKI. Linear formulation for ADC and DTI mode is simply obtained by taking logarithms on diffusion signals, and the fitting results are ADC values and tensor values respectively. Widely used method least-square regression is used for these two calculation modes. DKI Mode is somewhat complicated; it's converted to the constrained Linear fitting formulation (23), and tensor values and kurtosis values are combined to the fitting target. A standard quadratic programming algorithm active set is used for this constrained linear fitting. DSC calculation is a deconvolution problem, which is often performed using Singular Value Decomposition (24). The result of DSC deconvolution is the residue function which is used to calculate perfusion parameters.

Non-linear

For models including IVIM, body DKI which compute Kapp

(sDKI in our software), and NODDI, the signal intensity curve follows a non-linear decay. Mathematical fitting was decomposed into two steps: (I) exhaustive grid search which was first applied to find the initial guesses of the parameters to be fitted. (II) Nonlinear optimization via the publicly available ALGLIB library (http://www.alglib.net).

Segmentation

The segmentation module allows users drawing the region of interest (ROI) on displaying image. In our current build, nine ROIs can be created at the same time. The ROI drawing is actually filling the image data with different scalar values. Zero indicates background and one to nine indicate the nine ROI labels. The drawing operation can only happen in the MasterWindowWidget which actually modifies the 3D image data in its label image layer. The draw event triggers the update of statistical data in the statistic view. The statistical data include mean, standard deviation, minimum and maximum values of all the ROIs. The label image layer of SlaveWindowWidget only stores the 2D image which is a copy of label image in MasterWindowWidget with current slice. When an update event is triggered, the new slice of label image is copied and sent to the SlaveWindowWidget. The 3D label image can be saved to hard disk as an .nrrd file for later loading.

Data exporting

The data exporting is designed in two ways. One is for writing computing images to local hard disk in the format of DICOM which is handled by calculation module. A QT signal will be emitted by the GUI when users click the saving button, the calculation module will answer the signal accordingly. The other is to upload the computed image within selected ROI to server for further investigate, e.g., radiomics feature extraction as www.vusionmed. com providing. A developer can easily implement the client-server communication part by using the http tools providing by Qt.

Results

The graphic user interface (GUI) of IMAge/enGINE

The GUI of IMAge/enGINE is shown in *Figure 2*. The MasterWindowWidget shows the source 3D+C image data. And if a button in calculation modules is toggled on,

a related SlaveWindowWidget is added into Display Core layout loaded with related 2D computed Image.

Support of DICOM

DICOM images can be loaded into the software for displaying the diffusion and dynamic MRI data of GE, Siemens and Philips can be loaded into the software for computing. However, the mosaic image of DTI data from Siemens scanner will be treated as one single slice as *Figure 3*.

Image display

The basic interactions of image display are implemented:

- (I) Dragging left-button: window/level adjust;
- (II) Dragging middle-button: pan image;
- (III) Dragging right-button: zoom image;
- (IV) Wheel: scroll up/down image.

And the buttons on MasterWindowWidget and SlaveWindowWidget provide functions such as image screenshot, applying a full-rainbow sugar color encoding and mouse probing.

Calculation

So far three modules of quantitative image computing were listed in *Table 1*. We tested the diffusion module on two computers A and B with different configurations. One is equipped with a 4th gen i5 CPU which has a relatively lower performance than most of the using CPUs, while the other equipped with a high-performance Xeon E5 CPU. The calculation speed is listed in *Table 2*. All the algorithms are implemented using C++ as ITK filter pipeline. For time consuming algorithms: DKI and NODDI, multi-thread computing is employed. In comparison, we used computer A to run a DKI algorithm of DIPY on the same dataset and it costs 30 seconds which was roughly 10 times longer.

In the GitHub repository of IMAge/enGINE, the source code of calculating ADC is given as an example. Developers can implement their own algorithm as a new calculation module to process the high-dimensional data. For modules developed in C++, the interface of the example ADC module can be directly copied. If other languages such as MATLAB, Python and Java are used, the calculation module should be built into an executable file (or packed into a Docker image) and be called by IMAge/ enGINE framework. Due to the slice-by-slice updating scheme of IMAge/enGINE, the computation will only be

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Figure 2 Modules in GUI of IMAge/enGINE.



Figure 3 Loaded siemens DTI mosaic images. Left: source DTI images. Right: Calculated Fractional Anisotropic images.

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Table 1	Quantitative	image	computing	modules
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Modules	Computed image	Author	Available for free?
Diffusion module	Apparent Diffusion Coefficient (ADC)	Vusion tech	Yes
	Exponential ADC (eADC)		
	Computed DWI (cDWI)		
	Intra-voxel incoherent motion (IVIM) (6)		
	Simplified diffusion kurtosis imaging (sDKI) (25)		
	Diffusion tensor imaging: fractional anisotropy (FA) with and without direction color encoding; axial diffusion coefficient (AD); radial diffusion coefficient (RD)		
	Diffusion kurtosis imaging: axial kurtosis (AK); mean kurtosis (MK); radial kurtosis (RK) (7)		
	Neurite orientation dispersion and density imaging (NODDI) (8)		
Dynamic contrast module	Time-to-peak (TTP); mean-transit-time (MTT); wash-in (WIN); wash-out (WO)	Vusion tech	Yes
Dynamic susceptibility module	Cerebral blood volume (CBV); cerebral blood flow (CBF); time-to-peak (TTP); mean-transit-time (MTT); time-of-arrival (T0)	Vusion tech	No

Table 2 Calculation speed of diffusion parameters on two computers with different configurations

Coloulation type	In-plane image size	Number of Divelues and directions	Computing time for one slice	
Calculation type		Number of B-values and directions —	Computer A	Computer B
ADC/eADC/cDWI	228 228	2 B-values	~2 ms	~2 ms
IVIM	256 256	20 B-values	~720 ms	~580 ms
sDKI	256 256	20 B-values	~350 ms	~300 ms
FA/cFA/RD/MD*	128 128	1 B-value * 16 directions	~120 ms	~50 ms
AK/RK/MK*	256 256	2 B-values * 25 directions	~3.14 s	~1.36 s
NODDI*	256 256	2 B-values * 25 directions	~260 s	~118 s

*Modules implemented with multi-thread computing. Computer A: CPU Intel i5-4570 CPU, 4 Cores, 3.20 Ghz; Computer B: CPU Intel Xeon E5-1650 *2, 12 Cores in total, 3.50 Ghz. ADC, apparent diffusion coefficient; eADC, exponential ADC; cDWI, computed DWI; IVIM, intra-voxel incoherent motion; sDKI, simplified diffusion kurtosis imaging; FA, fractional anisotropy; RD, radial diffusion coefficient; MD, mean diffusivity; AK, axial kurtosis; RK, radial kurtosis; MK, mean kurtosis; NODDI, neurite orientation dispersion and density imaging.

applied on the relevant slices. The computing efficiency of calculation modules largely depends on the algorithm and the language used.

Segmentation

Two classic ways of drawing ROI are provided. One is to click on the image to add control points to form a closed contour which labels the area inside the contour with ROI index. The other is to directly paint on the image with input brush size and ROI index. Since the ROIs are stored in MasterWindowWidget as different values in a 3D label map, drawing on different slice of source image with the same ROI index forms a 3D ROI. The 3D label map is written as a .nrrd file onto local hard disk by clicking the save button. The coordinates of the 3D label map are handled to keep its geometry consistent with the source image, shown in *Figure 4*.



Figure 4 Example of the 3D label map and the source image. (A) Drawing ROI on the right eye of using IMAge/enGINE (only 3 slices are shown) and saving to disk. (B) Load the saved ROI to 3D slicer and the source image shows consistent geometry.

Discussion

Prospect of IMAge/enGINE will continue to be used by radiology research groups with or without professional engineers. For researchers with pure clinical background, the current software will provide them an efficient tool to complete diffusion, DCE model calculation and the ROI statistics measuring. For research groups with engineering backgrounds, they can write home-made plugins in python or C++ and integrate them into IMAge/enGINE, then deliver to clinical researchers to practice.

Comparision with other free softwares

There is a trade-off between being easy to use and extensive for medical imaging processing softwares. The phrase "easy-to-use" in this article indicates that even a researcher with clinical but without programming or engineering background still can have a friendly and convenient using experience. *Figure* 5 lists some medical image softwares. Most of the well-integrated softwares such as commercial workstations optimize with usability and usually provide a dedicated workflow for anatomy and modality. Though

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Figure 5 A list of medical imaging processing softwares ranking by easy-to-use and easy-to-extend.

some of these workstations provide research interface for adopting home-made algorithm, e.g., FUNC2 of GE and PRIDE of Philips, their flexibility of extension is limited. OsiriX is a well-integrated DICOM viewer focusing on image fusion, rendering and annotation while being partial open-source and providing a plug-in mechanic for other developers to add their own tool. 3D-Slicer and Image-J are open-source and broaden their extensibility to provide multiple extension interfaces for plugins developed by other research groups. However, clinicians and researchers may feel confused about the amount of parameters exposed in the GUI. It usually requires user to be able to attend workshop, find proper plugins and/or carefully read documentations to be able to complete their research tasks. Toolkits with GUI such as MITK provide greater flexibility to allow users to combine the various command line tools and libraries they provide to generate their own task-oriented tool for handling images. However, without proper programming knowledge, it is unlikely to use this type of software for the research group without engineering background members. There are also toolkits without GUI such as DIPY 19 which is popular among research groups with developing abilities.

In terms of balancing easy-to-use and easy-to-extend, IMAge/enGINE is similar to OsiriX by providing documented plug-in mechanism for extensibility. But IMAge/enGINE is freely available, ad-free, more opensource and focused on image computation instead of image fusion as well as render. Moreover, MRI is vastly used in brain research and there are many software focused on processing neuro imaging such as FSL (26), MRTrix (27), SPM (28), PANDA (29) and DSI studio (30). Though IMAge/enGINE provides similar functions such as DTI, DKI, NODDI calculation, IMAge/enGINE is not designed for any specific anatomy parts but for high dimensional data. So far the majority of the users of IMAge/enGINE have been using it to process body MRI data, such as breast, prostate and liver.

Terms of license

IMAge/enGINE is freely available for academic research use (studies paid by government-derived funds or donations) provided that the original research publications relevant to the software are cited. The software is also free for educational purposes. Note that the license terms do not generally include trials paid by pharmaceutical companies. For commercial use, IMAge/enGINE is sold and supported by the company Vusion Tech, Hefei, China. Individuals or organizations are not allowed to compile software products derived from IMAge/enGINE that are to be sold commercially or shipped together with other commercial products.

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Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

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