

RMCA-LSA: A Method of Monkey Brain Extraction

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Abstract: The traditional level set algorithm selects the position of the initial contour randomly and lacks the processing of edge information. Therefore, it cannot accurately extract the edge of the brain tissue. In order to solve this problem, this paper proposes a level set algorithm that fuses partition and Canny function. Firstly, the idea of partition is fused, and the initial contour position is selected by combining the morphological information of each region, so that the initial contour contains more brain tissue regions, and the efficiency of brain tissue extraction is improved. Secondly, the canny operator is fused in the energy functional, which improves the accuracy of edge detection of rhesus monkey brain tissue while retaining the advantage of the traditional level set algorithm in processing an uneven gray image. Experimental results show that the algorithm can accurately extract the brain tissue of rhesus monkeys with an accuracy of up to 86%.

Keywords: Monkey brain extraction; magnetic resonance (MR); level set

1 Introduction

Most brain research is based on MRI images [1]. However, the intensity of the magnetic field during brain research generally does not exceed 1.5 Tesla, the imaging of the cerebral limbic area is fuzzy and the structural information of fine brain tissues such as hippocampus and amygdala cannot be obtained [2]. For this reason, the researchers propose to use the brain of macaque monkeys [3,4] and compare the results with the human brain across species, so as to better advance the study of the human brain. This paper focuses on the research on the brain tissue extraction of macaques. The rhesus monkey brain tissue is separated from the non-brain tissue in magnetic resonance imaging to accurately extract the rhesus monkey brain tissue, to ensure the validity of the follow-up research results. Therefore, based on the National Natural Science Foundation of China and the Natural Science Foundation of Shanxi Province, this paper aims to propose an automated algorithm for accurately extracting macaque brain tissue.

At present, brain extraction methods are divided into three categories: region-based method, graphbased method, and deep learning-based method. The method based on the region [5-8] is to conduct curve evolution and complete segmentation according to the intensity information of the target image. However, because only the gray level information of the image is considered, the uneven gray level of



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medical images cannot be accurately segmented. The method based on atlas [9-14] refers to making brain probability atlas for different macaque species based on prior information, then performs brain extraction through a specific registration method. This method requires a large amount of manpower and time to make the probability atlas of different macaque species, which is time-consuming and cannot achieve the purpose of rapid brain tissue extraction. Methods based on deep learning [15-19] mainly use the Bayesian convolutional neural network, Unet network model and other deep convolutional models to achieve the extraction of brain tissue. However, deep learning relies heavily on model training and parameter selection, which requires a lot of time and data to debug the model.

To sum up, facing the existing problems in the current research results, this paper uses the level set method [20-24] to extract the monkey brain. The level set method does not need a lot of data set and has a good effect on medical image processing. But the traditional level set method for the selection of the initial contour position has randomness, to a certain extent, reduce the rate of brain tissue extract, due to lack of edge information processing algorithms at the same time, leads to extract a monkey can't realize the brain tissue extract on the edge of the fine [25-29].

Therefore, the paper, with a view to the study of the extraction of the monkey brain, gives the Level set of fused Regional morphology and the Level set of the Canny functional function (RMCA-Level Set). Based on the level set algorithm, the construction method of its initial contour and energy function is improved. The main contents are as follows:

- 1. To solve the problem of location selection of the initial contour, the idea of partition is introduced and the contour building model combining regional morphological information is proposed. The initial contour construction is completed by combining the morphological information of each region, which can speed up the iteration speed of the initial contour and improve the accuracy of monkey brain tissue extraction.
- 2. To solve the problem that the effect of edge segmentation of brain tissue is not ideal, a level set algorithm integrating Canny energy functional is proposed. In LBF energy functional, Canny edge detection operator is integrated to enhance the segmentation of edge regions, which not only extends the advantages of traditional level set algorithm in processing gray-scale inhomogeneous images, but also strengthens the detection of target edge, so as to achieve the purpose of accurately extracting the brain tissue of macaque monkeys.

2 RMCA-Level Set Algorithm

The level set algorithm is a method to solve the curve evolution. The idea is that the low-dimensional curve is taken as the zero level set of the high-dimensional surface, and the initial contour is segmenting through the evolution iteration of the curve, that is, the initial curve where the level set function starts to iterate. The more target regions the initial contour contains, the shorter the iteration time is, and the more accurate the result is [30,31].

As shown in Fig. 1, the model in this paper consists of two modules. The first module is an improvement of the initial contour construction algorithm. Based on the idea of partition, statistical analysis is performed on the morphological information of each region to establish the equation, and the initial contour is constructed to include the brain tissue of the macaque monkey to the maximum extent. The second module is to increase the processing of the edge information of brain tissue. Canny edge detection operator is fused in the LBF energy functional to achieve accurate detection of the edge of monkey brain tissue.



Figure 1: The overall architecture of our proposed method

2.1 Initial Contour Construction Algorithm

2.1.1 Image Binaryzation

Image binarization is the process of changing the gray value of pixel point to 0 or 255 according to the set threshold. The threshold *th* is assumed to exist, and all pixels of the image are divided into two categories, one is greater than the threshold *th*, and the other is less than the threshold *th*. The mean values of a1, a2 and the mean value of the whole image are calculated respectively. The optimal threshold for the best segmentation effect is calculated according to the following formula.

$$\sigma^2 = \omega 1 \times (a - a1)^2 + \omega 2 \times (a - a2)^2 \tag{1}$$

where, $\omega 1$ = number of background pixels/total pixels, $\omega 2$ = number of foreground pixels/total pixels.

2.1.2 Delete Small Connected Areas

The connected area of the image was counted, and the small non-brain tissue area was deleted by threshold setting. To compare the area of the small connected area of the rhesus monkey image and accurately select the threshold, Fig. 2 selected the statistical data of the 300, 310, 320 and 330 layers of the rhesus monkey brain of 32127 for display, and cut off the data with the area of the connected area greater than 1000. It can be found that the Area of small connected Area is concentrated around 200, so the threshold value of Area is preliminarily set as 200.

At the same time, the area of the connected area of the layers 230, 260, 280, 290, 310, 320, 330, 340, 350, 360, 370, 380, 390 and 410 in no.32126 and no.32127 rhesus macaque MRI images were calculated. In Fig. 3, the statistical data with the connected area greater than 1000 are cut off, which better shows the statistical results of the data of small areas. It can be found from the box diagram that the box bodies of the connected area of the images of no.32126 and no.32127 macaques at different layers are mostly concentrated below 200. Therefore, in the step of deleting the small area connected area, the threshold is set as 200.



Figure 2: Scatter plots of connected areas in different layers of the same macaque



Figure 3: Box diagram of connected areas of different macaques. (a) 32126- Statistics of the small connected area of macaques and (b) 32127- Statistics of the small connected area of macaques

Tab. 1 shows the comparison results before and after deleting small connected regions. This step removes tissue such as the muscles around the eye and simplifies the calculation.

	5 8	
Before delete		
After delete		

Table 1: Remove small area connectivity region results

2.1.3 Partition

To make the initial contour contain more target areas and improve the accuracy of monkey brain extraction, the image needs to be partitioned. Because the four local areas can maximize the coverage of the target area with the least amount of calculation, and the more target regions are included, the faster and more accurate the curve evolves. For any image A, divide it into four rectangular local areas as the seed areas for constructing the initial contour. The target image A(N * M) is partitioned according to the following formula:

$$A = \begin{cases} A1(1:N/2, 1:M/2) \\ A2(N/2+1:N, 1:M/2) \\ A3(1:N/2, M/2+1:M) \\ A4(N/2+1:N, M/2+1:M) \end{cases}$$
(2)

The 300th layer image of rhesus monkey no. 32127 was divided into four local areas, and the centroid points of each connected region in each local area were calculated respectively. The results were shown in Fig. 4.



Figure 4: Centroid of each connected region in each rectangular region. (a) upper left, (b) upper right, (c) left lower and (d) right lower

2.1.4 Initial Contour Construction

According to the morphological characteristics of the images, the monkey brain focusses on the image center and around the brain tissue adhesion. Therefore, the Euclidean distance between each centroid point and the partition point is calculated based on the centroid points of each connected region. In each local area, the centroid point with the smallest Euclidean distance is selected as the seed point to construct the initial contour, and the construction of the initial contour is completed [32,33]. Tab. 2 shows the construction results of the initial contours (red curve) of the monkey brain at different levels randomly selected from different macaques.



 Table 2: Results of constructing the initial contour

2.2 Fusing Canny's Energy Functional

To improve the efficiency of monkey brain edge detection, the edge detection operator was fused in the traditional level set LBF energy functional [34–36]. Tab. 3 compares the experimental results of various edge detection operators.



 Table 3: Comparison of edge detection results

Tab. 3 compares the detection results of the first-order edge detection operator (first row) and the secondorder edge detection operator (second row) on the same image, it can be found based on the second derivative test results of edge detection operator is better than the first derivative edge detection operator. The Canny operator adds a non-maximum value suppression operation and a dual-threshold filtering process to make the edge curve continuous and the lines thinner, making the detection result more accurate. Therefore, the Canny operator and the level set energy function are selected for fusion to better detect the boundary of monkey brain tissue.

The horizontal set function shall conform to the characteristics of the symbolic distance function, which is defined as follows:

$$\begin{cases} \Phi(x,y) > 0 & (x,y) \in \Omega_2 \\ \Phi(x,y) = 0 & (x,y) \in C \\ \Phi(x,y) < 0 & (x,y) \in \Omega_1 \end{cases}$$
(3)

Among, $\Omega 1 = outside(C)$, $\Omega 2 = inside(C)$.

Introduce the level set function into the energy functional, and define the following LBF energy function:

$$E^{LBF}(C, f_1(x), f_2(x)) = \lambda_1 \int_{\Omega} \left(\int_{outside(C)} K(x - y) |I(y) - f_1(x)|^2 H_{\varepsilon}(\Phi) dy \right) dx$$

$$+ \lambda_2 \int_{\Omega} \left(\int_{inside(C)} K(x - y) |I(y) - f_2(x)|^2 (1 - H_{\varepsilon}(\Phi)) dy \right) dx$$
(4)

Among, $l_1 = l_2 = 1.K(x) = K(|x|)$ is a non - negative monotone kernel function that depends on the image gradient. f_1 and f_2 are the gray fitting value of the inner and outer regions of the image near point x. $H_{\varepsilon}(x)$ is Heaviside function, $\delta_{\varepsilon}(x)$ is the derivative of the Dirac delta function. The definition of $H_{\varepsilon}(x)$ and $\delta_{\varepsilon}(x)$ is:

$$\begin{cases} H_{\epsilon}(x) = \frac{1}{2} \left(1 + \frac{2}{\pi} \arctan(\frac{x}{\epsilon})\right) \\ \delta_{\epsilon}(x) = \frac{d}{dx} H_{\epsilon}(x) = \frac{1}{\pi} \frac{\epsilon}{\epsilon^2 + x^2} \end{cases}$$
(5)

When canny detects the edge, the noise is first removed by the Gaussian filter. The generation equation of the Gaussian filter with the size of: $(2k + 1) \times (2k + 1)$

$$G(\mathbf{x}, \mathbf{y}) = \frac{1}{2\pi\sigma^2} \exp(-\frac{\mathbf{i} - (\mathbf{k}+1))^2 + (\mathbf{j} - (\mathbf{k}+1))^2}{2\sigma^2}); 1 \le \mathbf{i}, \mathbf{j} \le (2\mathbf{k}+1)$$
(6)

After the filtering operation is completed, the convolution operation is carried out, that is, all pixels in the image should be weighted and summed with their surrounding pixels. The gradient amplitude and direction of all pixels in the image are obtained according to the following formula.

$$M(x,y) = (P_x^2(x,y) + P_y^2(x,y))^{\frac{1}{2}}$$

$$\theta(x,y) = \arctan\left[\frac{P_Y(X,Y)}{p_x(x,y)}\right]$$
(7)

where, $P_x(i,j)$ and $P_y(i,j)$ are partial derivatives in the x and y directions, respectively. In order to prevent the thick curve from causing inaccurate segmentation results, it needs to set the high and low two threshold parameters. When the gradient of a point is greater than the high threshold, it will be marked as a strong edge point. When the gradient is less than the low threshold, it will be marked as a weak edge point. If it is between the two, it is necessary to determine the pixel points in the 8 areas around the point, when there is a strong edge point, the point is marked as a strong edge point, and vice versa. Therefore, for a certain point I(x, y) in the image, the canny energy functional can be defined as follows:

$$E^{CA} = \iint_{\Omega} C(x, y) H(\Phi) dx dy$$

=
$$\iint_{\Omega} \left[\frac{G^2(x, y)}{\partial x^2} + \frac{G^2(x, y)}{\partial y^2} \right] \times I(x, y) H(\Phi) dx dy$$
 (8)

where, * is the convolution operation, $H(\Phi)$ is The heisner function, G(x, y) is the Gaussian kernel function, it solves the second partial derivative concerning x and y respectively.

Also, the length term constraint function L(F) is added to constrain the properties of the curve so that the curve can be kept smooth.

$$L(\Phi) = \int \delta(\Phi) |\nabla \Phi(x)| dx$$
(9)

In the evolution process, in order to prevent the horizontal set function from losing the characteristic of sign distance function and causing instability in the evolution process, the distance penalty term function was added in the energy functional to shorten the reinitialization process of the level set, so that the evolution process could be carried out smoothly.

$$P(\Phi) = \frac{1}{2} (|\nabla \Phi(x)| - 1)^2 dx$$
(10)

Therefore, the total energy functional of the model in this paper is:

$$E^{CE} = E^{LBF} + E^{CA} + \mu P(\Phi) + \nu L(\Phi)$$
(11)

where, μ and v are constants.

The standard gradient descent flow is used to minimize the energy functional so that the $f_1(x)$ and $f_2(x)$ of the energy functional to the minimum value are:

$$\begin{cases} f_1(x) = \frac{\int_{\Omega} K(x-y) [H_{\varepsilon}(\Phi(y)) \times I(y)] dy}{\int_{\Omega} K(x-y) H(\Phi(y)) dy} \\ f_2(x) = \frac{\int_{\Omega} K(x-y) [(1-H_{\varepsilon}(\Phi(y))) \times I(y)] dy}{\int_{\Omega} K(x-y) (1-H(\Phi(y))) dy} \end{cases}$$
(12)

The gradient descent flow equation obtained by keeping $f_1(x)$ and $f_2(x)$ certain is:

$$\frac{\partial\varphi}{\partial t} = -\delta_{\varepsilon}(\Phi)(\lambda 1e1 - \lambda 2e2) + \delta_{\varepsilon}(\Phi) \times C + v\delta_{\varepsilon}(\Phi)div(\frac{\nabla\Phi}{|\nabla\Phi|}) + \mu(\nabla^{2}\Phi - div(\frac{\nabla\Phi}{|\nabla\Phi|}))$$
(13)

where,

$$\begin{cases} e_1(x) = \int_{\Omega} K(x-y) |I(x) - f_1(x)|^2 dy \\ e_2(x) = \int_{\Omega} K(x-y) |I(x) - f_2(x)|^2 dy \end{cases}$$
(14)

The process of solving the minimum value of the horizontal set energy functional is the process of solving the minimum value of the partial differential equation. This process meets the requirements of using the finite-difference method. The image space belongs to the equipartition grid, so the finite-difference method is generally used for calculation. It should be noted that in the case of fixed space step h, the time step Δt should meet the following conditions CFL(Courant - Friedrichs - Levy):

$$F_{MAX} \times \Delta t < h \tag{15}$$

where F_{MAX} represents the maximum moving speed of all points in the image.

The evolution equation of the level set function $\Phi_{i,j}^n$ of the point (i,j) in the image coordinate space at time N is:

$$\frac{\Phi_{i,j}^{n+1} - \Phi_{i,j}^{n}}{\Delta t} = L(\Phi_{i,j}^{n})$$
(16)

where, Δt is the time step, and $L(\Phi_{ij}^n)$ is the numerical approximation of the evolution of the horizontal set.

3 Analysis of Experimental Results

3.1 Dataset

1. UC Davis Dataset. The data set collected data from 19 rhesus monkeys using the Siemens Skyra3T scanner. The data included structures T1 and T2, as well as task-state fMRI and dMRI. In this experiment, NMR data with structures T1 were used. All the 19 monkeys were females, ranging in age from 18.5 to 22.5 years. The weight distribution was 7.28-14.95kg. Scanning sequence parameters: voxel resolution = $0.3 \times 0.3 \times 0.3$ mm, TE = 6.93 ms, TR = 15 ms, TI = 1100 ms, Flip Angle = 8°.

- 2. Mountsinai-P Dataset. Data of 9 macaques were collected by the Philips 3T scanner. The data included structures T1, T2, and dMRI. In this experiment, NMR data with structures T1 were used. The data set included 8 males and 1 female, with an age distribution of 3.4-8 years and a weight distribution of 4.7-7.42kg. The scanning sequence parameters were as follows: voxel resolution = $0.5 \times 0.5 \times 0.5$ mm, TE = 6.93 ms, TR = 15 ms, TI = 1100 ms, Flip Angle = 8°.
- 3. Princeton Dataset. The data set used Simens Prisma VE11C 3T scanner to collect data from two rhesus monkeys. The data included structures T1, T2, dMRI, and task-state fMRI. In this experiment, NMR data with structures T1 were used. All the two macaques in the data set were male, with an age distribution of 3 years and a weight distribution of 4.7-5.5kg. Scanning sequence parameters were as follows: voxel resolution = $0.5 \times 0.5 \times 0.5$ mm, TE = 2.32 ms, TR = 2700 ms, TI = 850 ms, Flip Angle = 9°.
- 4. Uminn Dataset. The data set was used to collect the data of 2 rhesus monkeys with Simens 7T scanner. The data included structures T1, T2, dMRI, and task-state fMRI. In this experiment, NMR data with structures T1 were used. All the two macaques in the data set were female, and their age distribution was over 10 years old. The scanning sequence parameters were as follows: voxel resolution = $0.3 \times 0.3 \times 0.3$ mm, TE = 3.65 ms, TR = 2500 ms, TI = 1100 ms, Flip Angle = 7°.
- Tab. 4 below shows some data of the above four experimental data sets.



 Table 4: Experimental data set

3.2 Self Contrast Experiment

Experiment one: The ce-LevelSet of the model in this paper was applied to UC Davis, Mountsinai-P, Princeton and Uminn, respectively, to verify the feasibility of the model in this paper. The default parameters of the model in this paper are: time step $\Delta t = 0.02$, space step h = 1, The parameter of Heaviside function and Dirac function is $\varepsilon = 1$. The length penalty coefficient is $0.001 \times 255 \times 255$, $\sigma = 8$.

Tab. 5 shows the results of monkey brain tissue extraction after the model was applied to four sets of UC Davis, Mountsinai-P, Princeton and Uminn, and it can be found that the model applied to four different data sets in this paper can achieve complete extraction of monkey brain tissue, with good extraction effect.



 Table 5: Brain extraction results

Tab. 6 shows the average value of DSC and JS of rhesus monkey brain tissue obtained after the model is applied to four data sets. From Tab. 6, it can be found that the monkey brain extraction results obtained after the model is applied to different data sets are good, and the similarity of DSC and JS with the standard brain tissue can reach about 0.75, and the performance is stable.

Dataset	DSC mean value	JS mean value
UC Davis	0.745	0.752
Mountsinai-P	0.734	0.745
Princeton	0.768	0.758
Uminn	0.753	0.763

Table 6: The average value of DSC and JS values of the brain extracted by the model in different data sets

3.3 Contrast Experiment

Experiment two: BET algorithm [37-39], watershed algorithm [40,41], LBF model and the model in this paper were used to extract the 300th, 320th, 340th and 360th layers of the brains of 32127 macaques in the UC Davis data set. A macaque brain image is cut into 512 layers according to the Z-axis [42], namely 512 images, each pixel size of 480×512 pixels. The number of iterations was set to 300 times for experimental comparison in this paper.

As can be seen from Tab. 7, the BET algorithm, LBF model, and Unet model cannot correctly peel off the fat and other non-brain tissues around the eyes, and their extraction effect on the edge part of brain tissues is poor. The Watershed algorithm cannot correctly extract brain tissue. In this paper, the model can accurately segment the eye and the temporal region and other muscle tissues, and the segmentation effect of the marginal region is better.

BET, watershed model, shown in Tab. 8 LBF model, Unet, this paper model the 300th floor of the monkey brain, 300, 320, 340 layers in the brain extract results as well as standard brain mask as a result, The red areas are the missing part of the mask extracted by the algorithm model compared with the standard mask, the yellow areas are the redundant part of the mask extracted by the algorithm model

compared with the standard mask, and the green areas are the overlap between the mask extracted from the algorithm model and the standard mask. It can be seen from the table that the extraction results from BET, LBF, and Unet models include non-brain tissues such as muscles around the eyes, and the segmentation of marginal tissues such as the temporal region is inaccurate. The results of the watershed model include more non-brain tissue, such as skull, and more brain tissue regions, which are not accurately segmented. The results of the model extraction in this paper are accurate for the edge segmentation of brain tissue and clean for the separation of non-brain tissue.

Model	Extraction results		
	Slice-300	Slice-320	Slice-340
BET			
Watershed algorithm	14055	*~)	₫^n
LBF model			
Unet			
Model of the paper			

 Table 7: Extraction results of different models

Table 8: Extracted masks for different model	ls
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Model -	Extraction results		
WIOUCI	Slice-300	Slice-320	Slice-340
BET			
Watershed algorithm		***	6
LBF model	100		
Unet			
Model of the paper			

3.4 Assessment and Analysis

In order to more accurately compare the differences between the extraction results of different models, DSC and Jaccard similarity coefficient are used to carry out quantitative analysis on the experimental results. The DSC and Jaccard similarity coefficients range from 0 to 1 The closer the value is to 1, the higher the similarity is and the more accurate the segmentation result of the model is. On the contrary, the closer the similarity coefficient value is to 0, the lower the similarity is, and the worse the segmentation effect of the model is.

Tab. 9 shows the DSC and JS comparison results of the BET algorithm, watershed algorithm, LBF model, Unet, and the model in this paper after extracting the brain of 32127 rhesus monkeys. It can be found that the average value of DSC value and JS value of this model are 0.745 and 0.752, all take to its highest level in five models, the results show that this model of extracting similarity is higher than other models, at the same time, DSC and JS value standard deviation was 0.064 and 0.063, respectively, in the five models at the minimum, indicating that this model is more stable than other models, the results of the monkey brain images are extracted difference is smaller. The mean values of DSC and JS values of the other four models are all below 0.7, indicating that the monkey brain extraction results of these four models are poor. Meanwhile, the standard difference between DSC values and JS values can reach 0.1 or above, which is greater than the standard deviation of the model in this paper, indicating that these four models are less stable than the model in this paper.

	DSC		JS	
	Mean value	Standard deviation	Mean value	Standard deviation
BET	0.651	0.192	0.659	0.225
Watershed algorithm	0.402	0.233	0.413	0.233
LBF model	0.645	0.119	0.579	0.107
Unet	0.689	0.103	0.648	0.093
Model of the paper	0.745	0.064	0.752	0.063

Table 9: Mean and standard deviation of DSC and JS values extracted by different models for the brain 32127

Fig. 5 shows that the median of the DSC value of the model in this paper is 0.78, which is the highest among the five models, the box body part between 0.82 0.67, The median of the DSC values of the LBF, Unet, and BET models are all in near 0.7, the LBF and Unet box parts are narrow, the BET model box part is between 0.5 0.75, the median of the DSC value of watershed algorithm is 0.42, and the box is between 0.23-0.48. we can get the brain tissue extract results of this model are the best.

Fig. 6 shows that the median of the JS value of the model in this paper is 0.78, which is the highest among the five models. the box body part between 0.81 and 0.66, LBF, the median of the DSC values of the LBF, Unet, and BET models are all in around 0.6, the LBF and Unet box parts are narrow, the box part of the BET model is between 0.52-0.77, we can get the brain tissue extract results of this model are the best.

Meanwhile, in this paper, the brain tissue of Monkey No. 32125 to 32136 was extracted using the above five models, and the DSC value and JS value of the extracted results were shown in Figs. 7 and 8.



Figure 5: Dice similarity coefficient



Figure 6: Jaccard similarity coefficient



Figure 7: Comparison of DSC values of different models



Figure 8: Comparison of JS values of different models

As can be seen from Figs. 7 and 8, DSC value and JS value in the brain tissue extraction results of different rhesus monkeys in the model presented in this paper perform well, at the highest value, and the model is relatively stable, with little difference in the extraction results for different rhesus monkeys. The extraction results of LBF, Unet, and BET models were significantly different and the models were unstable, while the extraction results of watershed models were poor and very unstable. In conclusion, compared with the other four models, DSC value and JS value in the experimental results of this model are close to 1, with a higher similarity coefficient and more accurate extraction results.

4 Conclusions

The automatic extraction of the macaque brain is the primary problem encountered in the research of the macaque brain. Therefore, this paper puts forward the RMCA-Level Set algorithm. First, the idea of partition is fused, and the initial contour is constructed by combining the mentality information of each region. The initial contour is constructed by binarization, deletion of small connected regions, and calculation of the center of mass by regions, etc., so as to improve the efficiency of brain tissue extraction. Secondly, to solve the problem of low edge extraction accuracy, a canny operator was fused into LBF energy functional to improve the extraction accuracy of the edge of brain tissue in this paper. Numerous experiments show that this model can be used to extract the brain tissue of the rhesus monkey.

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