Regional Ischemic Vulnerability of the Brain to Hypoperfusion
The Need for Location Specific Computed Tomography Perfusion Thresholds in Acute Stroke Patients

Seydomehdi Payabvash, MD; Leticia C.S. Souza, MD; Yifei Wang, BSc; Pamela W. Schaefer, MD; Karen L. Furie, MD; Elkan F. Halpern, PhD; R. Gilberto Gonzalez, MD, PhD; Michael H. Lev, MD

Background and Purpose—To characterize the spatial pattern of cerebral ischemic vulnerability to hypoperfusion in stroke patients.

Methods—We included 90 patients who underwent admission CT perfusion and MRI within 12 hours of ischemic stroke onset. Infarcted brain lesions (“core”) were segmented from admission diffusion-weighted imaging and, along with the CT perfusion parameter maps, coregistered onto MNI-152 brain space, which was parcellated into 125 mirror cortical and subcortical regions per hemisphere. We tested the hypothesis that the percent infarction increment per unit of relative cerebral blood flow (rCBF) reduction differs statistically between regions using regression analysis to assess the interaction between regional rCBF and region variables. Next, for each patient, a “vulnerability index” map was constructed with voxel values equaling the product of that voxel’s rCBF and infarction probability (derived from the MNI-152–transformed, binary, segmented, diffusion-weighted imaging lesions). Voxel-based rCBF threshold for core was determined within the upper 20th percentile of vulnerability index map voxel values.

Results—Different regions had different percent infarction increase per unit rCBF reduction (P=0.001). The caudate body, putamen, insular ribbon, paracentral lobule, and precentral, middle, and inferior frontal gyri had the highest ischemic vulnerability to hypoperfusion. A voxel-based rCBF threshold of <0.42 optimally distinguished infarct core in the highly-vulnerable regions, whereas rCBF <0.16 distinguished core in the remainder of the brain.

Conclusions—We demonstrated regional ischemic vulnerability of the brain to hypoperfusion in acute stroke patients. Location-specific, rather than whole-brain, rCBF thresholds may provide a more accurate metric for estimating infarct core using CT perfusion maps. (Stroke. 2011;42:1255-1260.)

Key Words: computed tomography ■ magnetic resonance imaging ■ stroke
variable ischemic sensitivity. Next, we constructed a voxel-based “vulnerability map” to visualize this spatial heterogeneity and to determine the voxel-based CTP blood flow thresholds that optimally correlate with infarct core.

Materials and Methods

Patients

Records of all stroke patients who underwent admission CTP and MRI at our center between May 2008 and June 2009 were reviewed. Inclusion criteria were: unilateral first-ever ischemic stroke; admission CTP and MR-DWI scans acquired within 12 hours of symptom onset and within 3 hours of each other; and the absence of any previous brain abnormalities based on admission MRI and clinical history. Our study received Institutional Review Board approval and was compliant with the Health Insurance Portability and Accountability Act.

Image Acquisition

All CT scans were obtained with a multidetector helical scanner (Light Speed; GE Medical Systems). CTP followed noncontrast CT and CTA, comprising a 90-second shuttle-mode acquisition. 1 image per slice every 3 seconds, after intravenous administration of 35 mL nonionic iodinated contrast (7 mL/s). Acquisition parameters were 80 kVp and 200 mAs, covering an 8-cm axial section of 16 adjacent 5-mm slices. Total radiation dose was <450 mSv, which is less than the 500-mSv Food and Drug Administration recommended upper limit.\(^2\) CTP source images were transferred to a GE Advantage workstation for postprocessing using deconvolution-based commercial software (CT Perfusion 3; General Healthcare) without application of vessel-suppression algorithms. A single reference arterial input function was selected semiautomatically as described previously.\(^3\)

MRI was performed on a 1.5-T Signa scanner (GE Medical Systems). Our standard stroke MR protocol includes a DWI sequence with two 180-degree pulses to reduce eddy–current distortions. Repetition time was 5000 ms; echo time was minimal. Axial images were acquired with 5-mm slice thickness and 1 mm interslice gap.

Image Analysis

For MRI, we manually segmented infarct core on admission DWI and developed a binary imaging dataset in which all voxels inside the infarct core were assigned a value of “1” and voxels outside of the core were assigned a value of “0.” These binary DWI lesion maps, along with the CTP parameter maps, were automatically coregistered to the MNI-152 brain space using FLIRT 5.5 (FMRIB Linear Image Registration Tool).

Two series of analyses were performed next: “region-based” and “voxel-based.”\(^*\) The former was used to detect differences in regional percent infarction increment per unit reduction of blood flow to establish the presence of regionally variable ischemic sensitivity in the brain. For the region-based analyses, the CTP and binary DWI lesion images were automatically parcellated into 125 paired parcellated brain regions. Next, the percent infarction and relative CBF (rCBF) were calculated across all patients. Then, multivariate regression analysis was applied to the region-based dataset to test the hypothesis that percent infarction increment per unit rCBF reduction differs statistically between the 125 paired parcellated brain regions. A regression model was constructed correlating percent infarct volume within each brain region with the following input variables: (1) regional rCBF; (2) an arbitrary categorical variable representing each region; and (3) an interaction term between the first 2 variables. A significant probability value in the interaction term of the model (rCBF×region) would support the hypothesis that percent infarct volume increment per unit rCBF reduction differs statistically between the brain regions.

We determined the brain regions with highest ischemic vulnerability using simple linear regression. For each region, the linear regression equation correlating regional percent infarction with regional rCBF was calculated. The slope of the regression line, B, reflects the increase in regional percent infarction per unit reduction in blood flow (higher slopes [|B|] suggest greater ischemic vulnerability).

For the voxel-based analyses, we determined the optimal pooled voxel-based rCBF thresholds that could distinguish infarct core from noninfarcted brain on the segmented admission DWI scans using receiver-operating characteristics curve analysis. First, we determined the optimal threshold at the operating point of the receiver-operating characteristics curve for the pooled voxels located within the upper 20\(^{th}\) percentile of the VI voxel values on the mean vulnerability map. Next, we determined the threshold for the pooled voxels in the remainder of the brain.

All values were expressed as either percentages or means±SD. Statistical analyses were performed using STATA 10.

Results

We included 90 patients with acute first-ever unilateral stroke. Of these, 51 (57\%) had left hemispheric stroke and 54 (60\%) were male. Based on the admission CTA imaging reports, 72 (80\%) patients had anterior circulation arterial occlusion (7 anterior cerebral artery, remainder middle cerebral artery), 6 (7\%) had posterior circulation occlusion, and 12 (13\%) had no visible occlusion. The majority of patients in our study had infarction in the middle cerebral artery territory (Figure 1). Admission CTP scans were performed 0.5 to 8.5 hours after stroke onset (mean, 3.7±2.0), and MRI scans followed CTP within 0.2 to 2.7 hours (mean, 0.4±0.3). There was a significant linear relationship between regional percent infarction and rCBF across all regions and patients ($R^2=0.35; P<0.001$).

The multivariate regression model showed a strongly significant interaction between rCBF and the region vari-
ables, confirming that regional percent infarction increment per unit rCBF reduction differs statistically between parcelled regions \((P/0.001; \text{Table 1})\) and, hence, that ischemic vulnerability varies between locations.

This regional variability was also shown by simple linear regression for each of the 125 paired parcellated brain regions. Overall, regional percent infarction increased with decreasing rCBF. Table 2 lists those regions with highest slope of the regression line corresponding to greatest ischemic vulnerability \((|B| = \text{slope, increase in regional percent infarction per unit reduction in blood flow}).\) The corpus callosum had the lowest \(|B|\) values \((|B|=0.10; R^2=0.23 \text{ left}; |B|=0.03; R^2=0.30 \text{ right})\), whereas the insula, precentral gyrus, and basal ganglia had the highest \(|B|\) values.

**Table 2. Brain Regions With Highest Ischemic Vulnerability to Hypoperfusion**

<table>
<thead>
<tr>
<th>Regions</th>
<th>Left Hemisphere</th>
<th>Right Hemisphere</th>
</tr>
</thead>
<tbody>
<tr>
<td>Caudate body</td>
<td>0.94</td>
<td>0.84</td>
</tr>
<tr>
<td>Putamen nucleus</td>
<td>0.79</td>
<td>0.85</td>
</tr>
<tr>
<td>Insular ribbon</td>
<td>1.30</td>
<td>1.15</td>
</tr>
<tr>
<td>Middle frontal gyrus</td>
<td>0.82</td>
<td>0.95</td>
</tr>
<tr>
<td>Frontal lobe subcortical matter</td>
<td>0.78</td>
<td>0.91</td>
</tr>
<tr>
<td>Precentral gyrus</td>
<td>1.30</td>
<td>1.15</td>
</tr>
<tr>
<td>Frontal lobe paracentral lobule</td>
<td>0.74</td>
<td>1.04</td>
</tr>
</tbody>
</table>

For each of the 125 paired parcellated brain regions, the linear regression equation correlating regional percent infarction with regional relative cerebral blood flow was calculated. This Table lists those regions with highest slope of the regression line corresponding to greatest ischemic vulnerability \((|B| = \text{slope, increase in regional percent infarction per unit reduction in blood flow}).\)

Voxel-based analysis provided visual corroboration of the region-based results (Figure 2). On the mean vulnerability map, the basal ganglia, insula, precentral gyrus, paracentral lobule, middle frontal, and inferior frontal gyri appear highly vulnerable bilaterally. Based on operating point receiver-

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**Table 1. Multivariate Regression Analysis to Test the Hypothesis That Percent Infarction Increment per Unit Relative Cerebral Blood Flow Reduction Differs Statistically Between the 125 Paired Parcellated Brain Regions**

<table>
<thead>
<tr>
<th></th>
<th>Left Hemisphere</th>
<th>Right Hemisphere</th>
</tr>
</thead>
<tbody>
<tr>
<td>B</td>
<td>SE</td>
<td>(\beta)</td>
</tr>
<tr>
<td>rCBF</td>
<td>0.535</td>
<td>0.487</td>
</tr>
<tr>
<td>Region</td>
<td>0.063</td>
<td>0.003</td>
</tr>
<tr>
<td>rCBF×region</td>
<td>-0.057</td>
<td>0.004</td>
</tr>
<tr>
<td>Constant</td>
<td>2.96</td>
<td>0.426</td>
</tr>
</tbody>
</table>

A regression model was constructed correlating percent infarct volume within each brain region with the 3 variables listed in the first column. The significant interaction term \((rCBF×\text{region})\) supports the hypothesis that percent infarct volume increment per unit rCBF reduction does differ statistically between parcellated brain regions (ie, ischemic vulnerability of brain tissue varies between locations).

\(B\) indicates regression coefficient; \(\beta\), standardized regression coefficient; \(P\), value of the input variable; rCBF, relative cerebral blood flow; SE, standard error of the B coefficient.
operating characteristics curve analysis of those voxels with VI values within the upper 20th percentile (ie, highly vulnerable locations; Figure 3), a voxel-based relative cerebral blood flow (rCBF) threshold of <0.42 optimally distinguished infarct core in the highly vulnerable regions (voxels within the upper 20th percentile of values on the vulnerability map; area under receive-operating characteristics curve [AUC], 0.72; 54% sensitivity, 80% specificity; P<0.001; bottom left), whereas an rCBF threshold of <0.16 distinguished core in the remainder of the brain, with equal sensitivity and specificity (AUC, 0.73; 54% sensitivity; 79% specificity; P<0.001; bottom right).

**Discussion**

We have shown that ischemic vulnerability varies across brain regions, and that the caudate, putamen, insula, precentral gyrus, inferior frontal, and middle frontal gyri are among the locations most highly sensitive to reductions in CBF. We have quantified that, for our particular CTP acquisition protocol and postprocessing software, ≈60% reduction in rCBF in these highly vulnerable locations distinguishes infarct core, whereas ≈85% reduction in rCBF distinguishes infarct core in the remainder of the brain.

The literature supports our findings. Previous studies in rats have shown higher frequency of DWI changes in the hippocampus, cortex, and caudate/putamen, and greater degree of selective neuronal loss in the caudate/putamen after unilateral hypoxia-ischemia. Cheng et al also have reported higher probabilities of infarct growth in the striatocapsular region of stroke patients with middle cerebral artery stem occlusion. Interestingly, the spatial pattern of sensitivity to hypoperfusion in our patients (Figures 2 and 3) resembles the topographical distribution of early DWI hyperintensities reported in patients with hypoxic-is-
chemic encephalopathy.\textsuperscript{7,8} This suggests the possibility that similar pathophysiological mechanisms of regionally selective neuronal loss may underlie cerebral injury in patients with embolic-occlusive and hypoxic-ischemic stroke.

There also are well-documented differences in the neurochemical response to ischemia of white matter versus GM, likely attributable to greater metabolic demands of GM.\textsuperscript{9} Within GM, certain cortical regions in our study appeared more vulnerable than others, including the insular, precentral, and inferior frontal gyri. The specific cortical areas with the highest ischemic sensitivity to hypoperfusion displayed on our mean vulnerability map (Figure 2) and regional regression results (Table 2) could, in part, be attributed to the high volume of convoluted GM at these locations. Our results also may reflect selective neurophysiologic vulnerability of these regions. In support of this latter hypothesis, Woo et al\textsuperscript{10} found selective cerebral GM loss in the frontal and insular cortices of patients with long-term heart failure, presumably attributable to ischemia accompanying perfusion deficits.

The vulnerability map of the brain reveals subtle topographical asymmetry (eg, of the frontal paracentral lobules; Figure 2 and 3) that may be artifactual; however, there may be other explanations for these findings. The inhomogeneous distribution of infarction in our patients (Figure 1) and/or the presence of outliers could contribute to this asymmetry, or it might reflect true asymmetrical vulnerability of these regions.

In agreement with previous studies, our rCBF threshold for infarct core was much higher (<0.42) in vulnerable cerebral areas versus the rest of the brain (<0.16). Arakawa et al\textsuperscript{11} reported CBF thresholds of 34.6 mL/100 g/min for GM and 20.8 mL/100 g/min for white matter in an MR perfusion study. Bristow et al\textsuperscript{12} found CBF thresholds of 20.0 mL/100 g/min and 12.3 mL/100 g/min for infarction in GM and white matter, respectively. Optimal CTP parameter thresholds for infarct core can vary significantly between vendors and even between different software from the same vendor.\textsuperscript{13} The optimal rCBF threshold (0.16) that we found in this study for less vulnerable brain voxels is in agreement with thresholds previously reported for an independent cohort who underwent a >65-second acquisition with postprocessing by the same software using the same default parameters (CT Perfusion 3).\textsuperscript{13}

Given the variability in absolute quantification with different software, we chose to report relative, rather than absolute, values.\textsuperscript{14} Although rCBV as a surrogate for infarct core is well-established, in our study we sought to study ischemic vulnerability based on the degree of hypoperfusion. The intrinsic physiological variability of CBV changes with ischemia, which include such phenomenon as luxury perfusion and elevated CBV attributable to autoregulatory mechanisms, make CBV a less desirable parameter given the aims of our study.\textsuperscript{15} Previous work also has suggested that CBV measurements are likely more sensitive to subtle differences in acquisition and postprocessing protocols than are CBF measurements.

Our findings may have clinical implications. Although we did not include time after ictus in our models, it is reasonable to hypothesize that less vulnerable areas may have a longer therapeutic time window for reperfusion treatment. This also may suggest that for those patients with ischemia at the most highly vulnerable brain locations, even early robust recanalization might be more effective if accompanied by neuroprotective therapies.

There are a number of limitations to our study. Our results are limited by the spatial distribution of stroke lesions in our cohort (Figure 1). We could not evaluate voxels that had few or no infarctions, most notably in the posterior circulation. The variable time between stroke onset, admission CTP, and admission DWI for different patients also may have distorted our results. Moreover, because CTP quantification is not yet standardized between different acquisition and postprocessing protocols from different vendors, the rCBF thresholds we report could vary slightly between different imaging platforms.\textsuperscript{13} Although the choice of a 20% cut-off in Figure 3 was arbitrary, this allowed us to visually threshold the “most” highly vulnerable regions for demonstration purposes.

Finally, the difference between our reported CBF thresholds and those of the literature is likely attributable to both differences in CTP acquisition length (45 seconds versus >65 seconds in current generation protocols), and that we did not apply vessel exclusion postprocessing algorithms to be consistent with our current clinical defaults and to maximize contrast-to-noise ratio. This suggests that the specific thresholds that we report may have limited generalizability to other acquisition and postprocessing platforms.

Conclusions
In conclusion, we have shown regional differences in ischemic susceptibility of the brain to hypoperfusion. Of the territories with infarction in our cohort, we found that the caudate and putamen were highly vulnerable, as were specific cortical areas, including the insula, precentral gyrus, and middle and inferior frontal gyri. Our findings support the hypothesis that location-specific thresholds may be more accurate than whole-brain thresholds for estimating the likelihood of infarction with CTP and, hence, have the potential to be of value in clinical management.

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Disclosures
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References


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背景和目的：描述脑卒中患者低灌注区脑组织局部缺血性易损性的空间分布情况。

方法：我们选择了90名在起病12小时内行CT灌注成像(CTP)及MRI检查的脑卒中患者。在弥散加权成像(DWI)相上对梗塞灶进行分割，同CT灌注成像的参数图，一起标记到MNI-150脑空间图上，图中双侧半球均被分作125对皮层及皮层下的区域。通过回归分析评估区域相对脑血流量(rCBF)和区域变量的关系，我们验证了脑内不同区域随每单位rCBF的减少梗塞百分率增加也不同这一假设。其次，对每个患者来说，“易损性的指标图”是由体素(体素的rCBF)和梗塞概率构建的(源于MNI-152转换的、二元、分段的DWI损伤)。评估梗死核心区的基于体素的rCBF阈值取决于第20百分位上的易损性指标图体素值。

结果：不同部位脑组织，随着单位rCBF的减少，梗塞百分率的增加不同(P=0.001)。尾状核体、硬脑膜、岛带、旁小叶外侧和中央前回、顶中回及额下回对低灌注的缺血易损性最高。基于体素的rCBF阈值<0.42最适宜于区别高易损区的梗死核心区，而rCBF阈值<0.16适于区别其他脑组织的梗死核心区。

结论：我们证实了急性脑卒中患者脑组织低灌注区的局部缺血易损性。CT灌注成像的rCBF阈值为我们提供了一个更为精确的度量标准评估梗死核心区，但它的使用有一定的区域特异性，并非适用于全脑。

关键词：计算机断层扫描，核磁共振成像，卒中

(Stroke. 2011;42:1255-1260. 吉林大学第一医院神经内科 孙莉 译 杨弋 吴江 校)

有多种因素能决定脑卒中后低灌注脑组织的命运，诸如脑血流减少的严重程度，侧枝循环建立的情况、发病的时程、局部脑组织对低灌注的敏感性。一些部位的脑组织基础代谢率高，如海马CA1区，对缺氧、缺糖极其敏感[1]。有文献证实构成灰质(GM)和白质的细胞组成不同与二者的脑血流(CBF)水平、代谢率不同有关，这为二者对低灌注不同的易损性提供了依据[1]。

尽管先前对选择性的局部缺血易损性已经有了相关的研究，但是神经影像学相关的空间多相性尚未被确立。确定区域敏感性能够帮助预测缺血半暗带区低灌注脑组织的预后。另外，相对不易受损的脑组织在卒中发生后对局部缺血能够耐受更长时间，这些组织对延迟的治疗有效。

在我们的研究中，运用CT灌注成像(CTP)和MRI的弥散加权成像(DWI)成像评估了急性脑卒中患者脑组织对低灌注的区域缺血易损性。CTP非常适用于CBF的定量分析，能够得出静脉内的对比浓度和CT像素强度的线性关系。首先，我们进行区域分析，分析局部每单位血流量的减少与梗塞灶百分比增加的关系，以确定不同部位缺血敏感性的不同并确认位置所在。另外，我们构建的以体素为基础的“易损图”使这种空间多相性更形象，并确定最适宜评估梗塞灶的以体素为基础的CTP血流量阈值。
材料和方法

患者

研究中我们回顾了医疗中心2008年5月到2009年6月就诊的行CTP和MRI检查的脑卒中患者。入选标准：单侧首发的缺血性脑卒中；症状出现12小时内行头部CTP及MR-DWI检查；既往史及头部MRI检查否认存在任何脑部疾病史。我们的研究得到了国家学术审查委员会的批准，并符合健康保险与责任法案的规定。

影像采集

所有的CT扫描均由多层螺旋扫描仪获取。行头部CT平扫及CT血管造影（CTA）后，行CTP检查，静脉注入35mL非离子型等渗碘对比剂后，每3秒钟完成一帧图像的扫描，由90秒模式获得的图像是CTP检查。采集参数分别是管电压80kVp、管电流200mAs，沿8cm的中轴分为16个相邻的5mm厚的层面。总辐射剂量<450 mSv，低于食品和药物管理局规定的上限500 mSv。然后将CTP影像移至GE工作站用重叠法商务软件（CT Perfusion 3; General Healthcare）进行后处理，而不是运用除外血管的方法计算。

MRI运用1.5 T的Signa扫描仪（GE Medical Systems）完成。我们标准的MR卒中协议包括带有2个180度脉冲的DWI序列以减少涡流的扭曲效应。重复时间为5000 ms；反射时间最小。轴向图像采用5 mm的层厚和1 mm的间隔厚度扫描。

图像分析

对于MRI，我们手动在DWI相上对梗塞灶进行分割并形成一组双相的数据集。这个数据集中梗塞灶的所有体素均赋值为“1”，梗塞灶外的体素赋值为“0”。使用FLIRT 5.5软件（FMRIB线性图登记工具）自动地把这些二元的DWI损伤图，和CTP参数图标注在MNI-152脑空间图上。其次进行如下两个序列的分析：基于区域和基于体素的分析。前者用于监测随每单位血流量的减少，区域梗塞灶百分比增加的不同，以确定脑内不同区域缺血敏感性的差别。对于以区域为基础的分析，在用传统书写软件(custom-written)程序构建的 Talairach图集的基础上，自动地把CTP和二元的DWI相损伤图分成125对对称的、皮质和皮质下区域。此外，计算有症状半球每个转化区域的梗塞百分比和相对脑血流量（rCBF）。将二元的DWI体素值（0 或者 1）线性转化到MNI-152脑空间图中，以致于把部分体素值分配于0-1之间，此值反映了相应体素发生梗塞的概率。因为不同的脑图整合到MNI-152上，体素会有差异，因此单纯的用是与否判断梗塞灶是否存在是不可行的。对于每个Talairach定义的部位，区域梗塞百分比被认为是梗塞概率体素值的均数。每个脑内部位rCBF值被确定为有症状侧半球该部位与对侧相应部位的平均绝对CBF的比率。

本研究为每个患者在MNI-152空间图上构建了以体素为基础的易损图。用类似于计算区域值的方法计算以体素为基础的rCBF值。对于每个体素来说，易损指标被定义为以体素为基础的rCBF值和该体素梗塞概率的乘积。因此，梗塞概率高的体素尽管rCBF高却有最高的易损指数(VI)值，而梗塞概率低的体素尽管低血流量却有最低的VI值。每个患者易损图的体素值被标在MNI-152脑空间图上，且计算出了所有患者每个体素的平均VI值。

为展示研究中患者的脑梗塞空间分布，我们开发了一个以MNI-152为基础的脑图，图上每个体素值代表梗塞的平均概率。

统计分析

在综合性的回归性分析中，我们首先在所有区域和患者中进行分析，以评价局部rCBF和梗塞百分比之间是否存在线性关系。然后为了验证随每单位rCBF的减少，梗塞百分比的增加在被分出的125对脑组织中是不同的，研究者对以区域为基础的数据集进行了多变量回归分析。研究者构建了一个回归模型，这一模型在每个脑内区域将梗塞容积百分比与如下输入变量相联系：(1)区域的rCBF；(2)代表每一大脑区域的任意类别变量；(3)前两个变量的交感项。该模型中交感项(rCBF×区域)的一个显著概率值会支持脑内各区域中，随每单位rCBF的减少梗塞容积百分比增加不同的这一假设。

我们用简单线性回归确定脑内缺血易损性最高的区域。每一区域，我们都用线性回归方程评估缺血百分比与局部脑血流量之间的关联。回归线的斜率[|B|]反映了血流量每减少一单位区域，梗塞百分比的增加(|B|越高说明缺血易损性越高)。

对于体素为基础的分析，我们制定了最佳的基于体素的rCBF阈值，使用接收-操作特性曲线分析时，这一阈值在DWI相分段的扫描中能区分出梗塞区和非梗塞区。首先，我们在接收-操作特性曲线上作用点确定最佳阈值，在平均易损图上将易损
表 1 多元回归分析以验证在 125 对脑内区域随 rCBF 的减少，梗塞百分比增加的情况

<table>
<thead>
<tr>
<th></th>
<th>B</th>
<th>SE</th>
<th>β</th>
<th>P</th>
</tr>
</thead>
<tbody>
<tr>
<td>rCBF</td>
<td>0.535</td>
<td>0.487</td>
<td>0.019</td>
<td>0.273</td>
</tr>
<tr>
<td>部位</td>
<td>0.063</td>
<td>0.003</td>
<td>0.279</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>rCBF × 部位</td>
<td>-0.057</td>
<td>0.004</td>
<td>-0.344</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>恒量</td>
<td>2.96</td>
<td>0.426</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

用上表中的 3 个变量构建与每个脑内区域相关联的梗塞百分比回归模型。变项 (rCBF × 部位) 支持不同部位的梗塞易损性不同的假设。统计，从分出的 125 对不同区域中，随每单位 rCBF 的减少，梗塞体积百分比增加不同。

B：回归系数; β，标准回归系数; P，输入变量值; rCBF，相对脑血流量; SE，B 系数的标准误。

### 结果

我们的研究纳入了 90 名首次发作单侧急性脑卒中的患者。这些患者中，有 51 名 (57%) 是左侧半球脑卒中，54(60%) 名为男性。头部 CTA 报告显示：前循环动脉闭塞者 72 名 (80%)，7 个大脑前动脉；后循环动脉闭塞者 6 名 (7%)；12 名 (13%) 未见明显的血管闭塞。研究中大部分患者存在大脑中动脉供血区的梗塞 (见图 1)。入院头部 CTP 检查在脑卒中后 0.5-8.5 小时内完成 (平均时间 3.7±2.0)，头部 MRI 在行 CTP 检查后 0.2-2.7 小时 (平均时间，0.4±0.3) 内完成。所有结果均显示区域梗塞百分比和其 rCBF 之间存在线性关系 ($R^2=0.35$; $P<0.001$)。

多元回归模型显示 rCBF 和局部变量之间存在显著相关性，统计证实不同区域间随每单位 rCBF 指标体素值定在第 20 百分位以上。然后我们确定剩余脑组织的合并体素的阈值。

所有的测试值均以百分率或均数 ± 标准差来表达。统计分析用 STATA 10 软件完成。

### 图 1
入组患者脑梗塞部位分布图。所有患者均有相应的体素值。这些体素值反映了梗塞发生的平均可能性。比色刻度尺是基于五分位分组的分层。
减少，区域梗塞百分比增加不同 \( (P<0.001, \text{表 } 1) \)。因此不同部位缺血易损变量亦不同。

这种区域变异性在被分出 125 对区域的每部分脑组织中，可通过简单的线性分析显示出来。总的来说，区域梗塞百分比随着 \( rCBF \) 的减少而增加。表 2 列举了回归线斜率最高的区域，与缺血易损性最敏感的部位相一致 \( (\text{斜率} = |B| \text{系数，随每单位血液流量的减少，区域梗塞百分比增加}) \)。胼胝体的斜率值最低 \( (\text{左侧} |B|=0.10 ; R^2=0.23 ; \text{右侧} |B|=0.03 ; R^2=0.30) \)。而脑岛、中央前回和基底神经节斜率值最高。

基于体素的分析结果为以区域为基础的结果 \( \text{(图 } 3 \) 提供了直观的证据。在平均易损率脑图上，基底神经节区、脑岛、中央前回、旁中央小叶、额中回、额下回均呈现出较高的易损性。基于对这些 \( VI \) 值在第 20 百分位以上的 \( (\text{例如：高易损区；图 } 3 \) 体素的工作点受试者操作特征性曲线的分析，基于体素的 \( rCBF \) 阈值为 0.42 最适于识别缺血灶。其他部位脑组织，基于体素的 \( rCBF \) 阈值为 0.16 对于鉴别梗塞灶有较高的敏感性和特异性。
讨论

我们的研究显示了不同脑组织的缺血易损性不同，且尾状核、壳核、岛叶、中央前回、额下回及额中回是缺血最敏感的区域。借助特有的CTP获取协议和后处理软件，我们对数据进行了量化统计，在高度易损部位，rCBF减少约60%即能鉴别出梗塞灶，而脑内其他部位rCBF减少约85%才能被鉴别。


有趣的是，在我们的研究对象中，对低灌注敏感脑组织的空间布局（图2、3）与报道的缺血缺氧脑病早期DWI显示的白质高信号分布类似[7,8]。这提示选择性的区域神经元缺失的病理生理机制可能成为栓塞-闭塞和缺血-缺氧性卒中患者脑损伤的基础。

不少文献证实白质和灰质对缺血的神经化学反应不同，这可能归因于GM的代谢需求更高[9]。我们的研究结果显示：在灰质中，某些区域脑皮层较其他部位更易受损，包括岛叶、中央前回及额下回。我们的脑平均易损图（图2）提示，皮层特定区域低灌注的缺血敏感性最高，在某种程度上区域回归分析结果（表2）可归因于在这些部位复杂的灰质结构容积较大所致。我们的结果亦反映了这些区域选择性的神经生理学易损性。

Woo等人[10]的研究发现，长期心功能衰竭患者的额部、岛叶皮质有选择性的脑灰质缺失，推测原因可能与灌注障碍导致脑缺血有关。这一研究支持我们的假设。CTP数据的获得方式不同、后处理模式的供应商也可能不同，这些均未经标准化处理，故我们报道的rCBF阈值由于不同的影像资料来源也会有轻微的变化[13]。尽管图3所示的20%界限的选择是随机的，但这使我们直观的看到了最高易损区的阈值。

结论

综上，我们表明了低灌注时脑组织缺血敏感性的区域差异。在我们研究群体的梗塞灶所在区域中，我们发现尾状核和壳核极易受损，一些特殊皮质区，包括岛叶、中央前回、额中回及额下回也易受损。
此项研究结果支持以下假设：在用 CTP 评估梗塞发生的可能性时，区域特异性阈值可能比全脑阈值更为准确。因此，区域特异性阈值在临床中具有很好的应用前景。

参考文献