Chronic systemic immune dysfunction in African Americans with small vessel-type ischemic stroke

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Abstract

Background—The incidence of small vessel-type (lacunar) ischemic strokes is greater in African-Americans compared to whites. The chronic inflammatory changes that result from lacunar stroke are poorly understood. To elucidate these changes, we measured serum inflammatory and thrombotic biomarkers in African-Americans at least 6 weeks post-stroke compared to control individuals.

Methods—Cases were African-Americans with lacunar stroke (n=30), and controls were age-matched African-Americans with no history of stroke or other major neurologic disease (n=37). Blood was obtained >6 weeks post-stroke and analyzed for inflammatory biomarkers. Freshly isolated peripheral blood mononuclear cells were stimulated with lipopolysaccharide (LPS) to assess immune responsiveness in a subset of cases (n=5) and controls (n=4).

Results—After adjustment for covariates, the pro-inflammatory biomarkers, soluble vascular cadherin adhesion molecule-1 (sVCAM-1) and thrombin anti-thrombin (TAT), were independently associated with lacunar stroke. Immune responsiveness to LPS challenge was abnormal in cases compared to controls.

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Conclusions—African-Americans with lacunar stroke had elevated blood levels of VCAM-1 and TAT and an abnormal response to acute immune challenge >6 weeks post-stroke, suggesting a chronically compromised systemic inflammatory response.

Keywords

small vessel disease; lacunar stroke; African-Americans; chronic inflammation; race-ethnic disparities; biomarkers

Background

About 795,000 people in the U.S. have a stroke each year with African-Americans having twice the risk of stroke as whites, especially at younger ages [1, 2]. African-Americans are also more likely to have silent lacunar infarcts on MRI compared to whites and are three times more likely to have lacunar strokes [3]. The reasons for this disparity are not fully explained by differences in common risk factors, nor are the underlying mechanisms known. Peripheral inflammatory processes are implicated as a causative factor in small vessel-type cerebrovascular disease, of which lacunar stroke is one phenotype [4–6].

There are at least two major gaps in current data related to the relationship between inflammation and lacunar stroke. Several studies report inflammatory biomarker data in the acute phase after stroke, thus restricting analysis to the immediate damage responses after stroke. Few studies have reported long term, chronic post-stroke changes in inflammatory markers. Second, the majority of studies have been done in whites, yet African-Americans are more likely to have clinical and subclinical small vessel-type cerebrovascular disease. Importantly, there are race-ethnic differences in inflammation at baseline after stroke with African-Americans having an increased systemic inflammatory state, as reflected by increased levels of C reactive protein (CRP) compared to whites. In addition to traditional small vessel risk factors, this increase in CRP identifies patients as being at higher stroke risk [7].

The goal of this study was to determine whether African-American patients have chronic immune dysfunction at >6 weeks post-lacunar stroke. We measured serum inflammatory, endothelial, and thrombotic biomarkers in African-Americans after lacunar stroke compared with age-matched African-American controls. Peripheral immune cells taken from blood samples of patients and control individuals were further evaluated for responsiveness to a pro-inflammatory stimulus (treatment with lipopolysaccharide (LPS). We hypothesized that there would be prolonged immune activity and higher pro-thrombotic marker levels in African-Americans after lacunar stroke, and that subjects with stroke would have immune dysregulation as assessed by LPS stimulation compared with controls.

Materials and Methods

Subjects

Cases—Patients were enrolled at Wake Forest Baptist Medical Center (WFBMC), Duke University Medical Center (DUMC), Carolinas Medical Center—Northeast, and University of California, Davis. All subjects were self-reported to be Black or African-American in

order to be eligible. Potential subjects were identified during a hospital admission for acute stroke and their medical records screened for clinical symptoms/signs, imaging results, and history. Patients were approached regarding study participation either during the index hospitalization or in the outpatient setting. Individuals with ischemic stroke (focal neurological deficit of vascular origin) with symptoms lasting greater than 24 hours or those with symptoms lasting less than 24 hours but having neuroimaging (computerized tomography (CT) or magnetic resonance (MR) imaging) evidence of a small vessel distribution (i.e. lacunar) infarction were selected for study. Lacunar ischemic stroke subtype was defined by the presence of a clinical syndrome (pure motor hemiparesis, pure sensory, mixed sensorimotor, ataxic hemiparesis, or dysarthria-clumsy hand) or CT or MR imaging evidence of an infarct in a small vessel distribution. Patients with cardioembolic, large vessel carotid or vertebral artery causes (greater than 50% stenosis by ultrasound, CT, or MR angiogram in the relevant artery) or non-atherothrombotic cause were excluded.

Controls—Control subjects self-reported as Black or African-American were recruited from the community via advertising and direct mailings at 2 of the sites (WFBMC and DUMC) to match the age frequency of cases. To be eligible, controls had no prior history of stroke (screened using the stroke-free questionnaire) [8], or other major neurologic diseases (Parkinson's disease, vascular or Alzheimer's dementia, multiple sclerosis, normal pressure hydrocephalus, traumatic brain injury with loss of consciousness greater than 30 minutes, epilepsy, or mental retardation).

Blood Collection

Subjects had a fasting phlebotomy at least 6-weeks after the stroke using standard procedures. The blood was immediately centrifuged at $1200g$ for 15 minutes, and plasma (from citrated tubes) and serum (from non-citrated tubes) aliquots were stored at $-20°C$ until analysis.

Biomarker quantitation

All cytokine or other biomarker analyses were carried out by custom Quantibody array multiplex analysis performed by RayBiotech (Norcross, GA). Briefly, 50 µl of each sample was diluted in sample diluent and incubated with target cytokine antibodies arrayed on a solid surface, followed by incubation with a biotin-labeled detection antibody. The cytokine-antibody-biotin complex was then visualized through the addition of a streptavidin-labeled Cy3 equivalent dye and detected using a laser scanner. Cytokine standards were run in parallel with all samples. The limits of detection (pg/ml) for cytokines measured using the custom Quantibody array were: interferon-γ (IFNγ: 16.8–30,000); S-100b (464.5–1,500,000); plasminogen activator inhibitor-1 (PAI-1: 112.8–150,000); tumor necrosis factor-α (TNFα: 7.0–3,000); vascular adhesion molecule-1 (VCAM-1: 1639–1,500,000); interleukin-1 receptor antagonist (IL-1ra: 23.0 – 6,000); interleukin-4 (IL-4: 1.6 – 3,000); interleukin-6 (IL-6: 5.9–3,000); interleukin-8 (IL-8: 1.3 – 3,000); and D-Dimer (0.08- 60pg/ml). Thrombin-Anti Thrombin complex (TAT: 1.48 – 120 ng/ml) was measured by ELISA (AbCam, Cambridge MA).

To measure immune cell response to a pro-inflammatory stimulus, fresh whole blood aliquots containing mixed peripheral blood mononuclear cells (PMBCs) was collected from a random subset of the stroke patients (n=5) and age-race matched controls (n=4). These subjects were randomly recruited from the case and control participants enrolled at WFBMC. Participants signed an additional consent form for this portion of the study. PBMCs were stimulated with either LPS (*E. coli* O111:B4; 100 ng/ml) or saline vehicle for 3 hours at 37°C. Following treatment, blood samples were centrifuged (for 10 minutes, 1100*g*) and the separated plasma was stored in aliquots at −20°C. Non-stimulated and LPS-stimulated samples were then assayed for cytokine levels as described above.

Statistical analysis

The distributions of the biomarkers were assessed for skew and log-transformed as appropriate, except if the majority of values were below the level of detection (i.e., zero). Odds ratios were estimated to determine the association of each biomarker with stroke status before and after adjustment for baseline characteristics that differed between groups in the descriptive analysis using logistic regression. Wilcoxon rank sum tests were used to compare medians of markers between stroke cases and controls, and chi square tests were used to compare proportions in these two groups. Spearman rank correlation tests were performed to determine correlations amongst the biomarkers. In the LPS study, data were analyzed using a paired Student's t-test (two-tailed) with $p < 0.05$ considered significant.

Results

Study population characteristics are given in Table 1. A total of 30 stroke cases and 37 controls were enrolled and included in the analysis. Controls were more likely to have high school education, whereas higher proportions of stroke subjects had hypertension, hyperlipidemia, and a history of substance abuse. There was no difference in the history of diabetes or cigarette smoking in stroke cases compared to controls. The median NIH Stroke Scale score among stroke cases at the time of enrollment was 1, with a range of 0 to 4. Table 2 shows differences in median log-transformed levels of biomarkers between cases and controls. Stroke cases had significantly elevated levels of sVCAM-1, IL-1ra, TAT, and IL-6 compared to age-matched controls. In contrast, D-dimer levels were significantly higher in controls than in stroke cases.

The associations between biomarkers and small vessel-type stroke, both unadjusted and adjusted for the variables that were different between cases and controls (age, sex, hyperlipidemia, and hypertension) are given in Table 3. Although IL-1ra, IL-6, IL-10, VCAM-1, and TAT were all higher in the unadjusted models, only VCAM-1 and TAT were independently associated with stroke after covariate adjustment. Table 4 shows correlations between individual biomarkers, with the most significant positive correlations between IL-1ra and IL-10 and IL-6 and IFNγ. Other highly significant correlations were between IL-1ra and IL-6, IL-1ra and VCAM-1, IL-1ra and TAT, IL-6 and IL-8, IL-10 and VCAM-1, and IL-10 and TAT.

The final part of this study addressed if acute immune function >6 weeks post-stroke was compromised in cases compared to controls. *Ex vivo* stimulation of PBMCs with LPS in a

randomly selected subset of patients and controls resulted in significantly elevated cytokine production compared with vehicle stimulation for IL-1ra, TNFα, IL-6, IL-8, and IL-10 in samples from controls (Fig 1a). In contrast, both vehicle- and LPS-stimulated blood from stroke cases produced equivalent, elevated levels of the same cytokines (Fig 1b).

Discussion

We measured changes in inflammatory and thrombotic biomarkers in African-Americans during the chronic phase (>6wks) after small vessel-type ischemic (lacunar) strokes compared with age-matched, stroke-free African-Americans. Circulating levels of two of these markers, sVCAM-1 and TAT, were independently associated with the chronic phase of lacunar stroke. Soluble vCAM-1 (CD106) is an established marker of inflammation and endothelial dysfunction, whereas TAT is an established marker of thrombin/coagulation activation. To our knowledge, there are no other studies that have evaluated either inflammatory or endothelial biomarkers and none measured plasma levels of sVCAM-1 in the chronic phase after lacunar stroke in African-Americans [9]. These later time points are likely to be important for repair after stroke and to the potential for re-occurrence of stroke.

The increased blood sVCAM-1 levels we observed in this study suggest prolonged changes in the vascular endothelium. VCAM is an important component of the vascular endothelium where it plays critical roles in immune surveillance of tissues, immune signaling and transmigration of neutrophils, monocytes, dendritic cells (DCs) and other immune or tumor cells [10, 11]. Activation of VCAM–mediated signaling pathways varies depending on the type of vascular injury and has been noted as an early pathological characteristic of endothelial dysfunction found in atherosclerotic lesions [12] or in focal cerebral ischemia in rodent models [13]. Unlike the relationship of intracellular cell adhesion molecule (ICAM), however, the relationship of sVCAM-1 to ischemic stroke is more tenuous [9, 14, 15]. Kozuka *et al.* [16] detected changes in sVCAM-1 and other endothelial factors as part of an acute phase in lacunar infarctions. However, a systemic review concluded that data on sVCAM-1 was insufficient to establish a firm relationship between sVCAM-1 and acute or chronic lacunar stroke. [9]

Our study of a group of African-Americans found a significant relationship between sVCAM-1 and lacunar stroke after adjustment for co-variables that included age, sex, hyperlipidemia, and hypertension. One explanation for the association between sVCAM-1 and stroke has been that sVCAM-1 is also associated with vascular risk factors such as hypertension, older age and hyperlipidemia. For example, patients with essential and acute hypertension have elevated plasma levels of soluble adhesion molecules, such as sVCAM-1 and sICAM-1 [17]. VCAM-1 could reflect the cumulative impact of these risk factors. But, after controlling for differences in baseline risk factors between cases and controls, we found that sVCAM-1 remained associated with chronic stroke suggesting the relationship may not be entirely related to these established stroke risk factors. Thus, although our study was not designed to measure plasma levels of either inflammatory or vascular endothelial factors prior to stroke, we speculate that, after controlling for baseline risk factors, the elevation of sVCAM-1 levels was more likely to be increased as a result of the stroke rather than a pre-existing condition.

We also found an association between higher TAT levels, a measure of the amount of thrombin inhibition (and hence indirectly of coagulation and fibrinolysis) and lacunar stroke after adjustment for potential confounders. The increased post-stroke TAT levels implies long-lasting changes in thrombin production after the index lacunar stroke. Vascular endothelial activation is proposed to change the blood vessel wall leading to platelet activation and TAT formation, providing a possible mechanism by which VCAM-1-mediated endothelial activation and immune signaling may contribute to coagulation within small vessels [18]. Thus, VCAM-1 and TAT may interact to contribute to the apparent prolonged endothelial damage found in African-Americans with lacunar stroke. A meta-analysis of measurements of plasma levels of tissue plasminogen activator (tPA), plasminogen activator inhibitor (PAI) and fibrinogen, however, showed no association between these coagulation and fibrinolysis markers and lacunar infarction [9]. Unfortunately, TAT was not included in this meta-analysis. Other studies show that TAT is increased in Binswanger's disease, which is characterized by dementia in the setting of severe leukoaraiosis and lacunar strokes [19]. The results of our study, therefore, are consistent with an association between TAT and small vessel disease.

The *ex vivo* assay of immune activation in a random subset of cases and controls in our study provides important insights into the immune phenotype after lacunar stroke. Cytokine measurement following *ex vivo* stimulation of whole blood with LPS is a validated, cost-effective surrogate assessment of monocytic, systemic cytokine production and immune responsiveness [20, 21]. PBMCs from non-stroke controls responded with a typical increase in cytokine biomarker production when challenged with LPS. In contrast, stroke cases had cytokine levels in vehicle- and LPS-stimulated samples that were not significantly different from those in LPS-stimulated control samples. Significantly elevated markers included canonical pro-inflammatory cytokines such as TNF-α and IL-6 as well as anti-inflammatory cytokines such as IL-1Ra and IL-10. The observation that baseline levels of cytokines were much higher in cases compared to controls and the failure of LPS to significantly increase cytokine production is indicative of a "ceiling effect," which suggests that these PBMCs have maximized their immune capacity in the chronic stroke phase. Taken together, these results suggest a lack of immune resolution and may reflect a maintained high inflammatory state after stroke.

Several biomarkers were highly correlated (Table 4), which could suggest additional mechanistic targets for further study in the chronic phase of lacunar stroke. Cytokines, particularly IL-1 and TNF-α, have pivotal roles in regulating inflammatory and innate immune responses to acute infection, but also regulate cellular pathways that are important in acute ischemic stroke [22, 23]. In general, TNF-α and IL-6 blood levels are elevated in acute lacunar stroke and associated with larger infarct volumes and poorer outcomes [24, 9]. Acute pro-inflammatory processes are followed at varying times by anti-inflammatory immune activation that leads to release of cytokines associated with repair and resolution of the immune response to injury. IL-1ra is an anti-inflammatory protein that reversibly blocks the pro-inflammatory actions of IL-1, and increased blood levels of IL-1Ra reduced the extent of acute ischemic injury in experimental models [25]. High blood IL-1ra levels immediately after stroke increase the risk of infection and may thereby lead to poorer

outcomes [26]. Interestingly, abnormally elevated levels of IL-1Ra were found up to 6 months after ischemic stroke [27]. We also found highly significant correlations between IL-1Ra and many proinflammatory cytokines. Overall, the results of these studies are consistent with our *ex vivo* data suggesting the presence of a dysregulated inflammatory response after lacunar stroke.

Our study was limited to a relatively small cohort (30 individuals with stroke, 37 control individuals) and was homogenous in terms of race/ethnicity, age and specific stroke subtype. All subjects with stroke had careful stroke subtype characterization and non-stroke controls were carefully age-matched. Thus, our intriguing findings in an African-American population may not be generalizable to other race-ethnic groups or other populations. We also specifically designed the study to evaluate candidate biomarkers that represented endothelial, inflammatory and coagulation/thrombin-based factors thought to be associated with lacunar stroke. Other biomarkers not assessed in our study may also be important. Importantly, we have examined an understudied but critical phase of stroke recovery, that is, the chronic phase, which may reveal unresolved stroke-based sequelae. Chronic elevation of VCAM-1 and TAT, markers of downstream inflammation, endothelial dysfunction, and thrombin generation, were associated with small vessel stroke in African-Americans. Further research comparing race/ethnic groups, and other stroke types is required to fully substantiate the pathophysiology of chronic changes in VCAM-1 and TAT levels and the changed immune response in our study population. In addition, further research using these biomarkers and immune challenges could have clinical implications for personalized medicine, such as identifying patients at risk of recurrent stroke, identifying at-risk immune phenotypes prior to the onset of first stroke, or identifying novel drug targets for prevention.

Conclusions

Chronic elevations of VCAM-1 and TAT are associated with small vessel stroke in African-Americans. Our data suggest that elevation of immune factors represents a failure to repair and resolve immune responses in the chronic phase after stroke when compared to stroke-free controls in this population.

Acknowledgments

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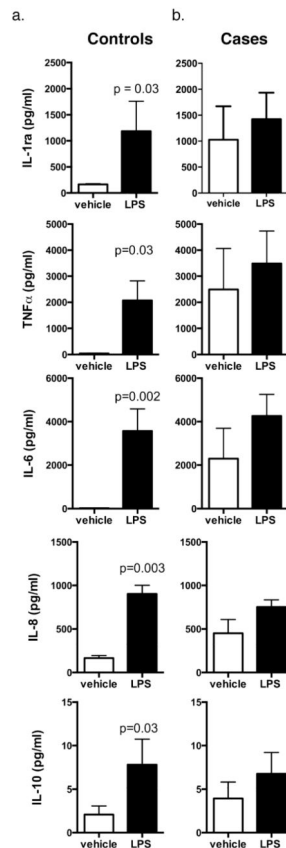
Fig. 1. *Ex vivo* **LPS stimulation of PBMCs from stroke cases and age-matched controls**
PBMCs were stimulated with saline vehicle or LPS (100 ng/ml) followed by cytokine analysis. **A.** LPS stimulation significantly increased cytokine levels compared to vehicle in controls. **B.** The same cytokines were similarly elevated with no significant difference between vehicle and LPS-stimulated samples from cases. Data=mean±SEM; n = 4–5 individuals, with p <0.05 as significant

Table 1

Baseline characteristics of cases and controls.[a]

Variable	Total (n=67)	Case (n=30)	Control (n=37)	P value
Age, yrs, median (IQR)	59 (53–66)	57.5 (50–62)	60 (55–67)	0.093
Female, n (%)	43 (64)	16 (53)	27 (73)	0.096
High school education, n (%)	59 (89.4)	23 (79.3)	36 (97.3)	0.018
Hypertension, n (%)	48 (71.6)	25 (83.3)	23 (62.2)	0.056
Diabetes, n (%)	19 (28.4)	10 (33.3)	9 (24.3)	0.416
Hyperlipidemia, n (%)	29 (43.3)	18 (60)	11 (29.7)	0.013
Tobacco smoker, n (%)	34 (50.8)	17 (56.7)	17 (46.0)	0.383
Substance abuse, n (%)	6 (9)	5 (16.7)	1 (2.7)	0.046
BMI, kg/m^2 (IQR)	31.0 (28.3–35.3)	29.6 (27.7–32.8)	32.0 (28.9–37.4)	0.065

[a]Numbers in parentheses refer to either interquartile range (IQR) or the percentage of individuals in each group (%).

Table 2

Comparison of biomarkers between stroke patients and controls using Wilcoxon rank sum test.[a]

Biomarker (log-transformed)	Stroke pg/ml, median (IQR)	Control pg/ml, median (IQR)	P value
IL-1ra	2.074 (1.793–2.274)	1.427 (−0.301–1.961)	0.001
IL-6	1.286 (1.017–1.529)	1.022 (−0.301–1.254)	0.009
IL-8	1.285 (0.922–1.477)	0.888 (.719–1.160)	0.017
IL-10	0.347 (−0.301–0.684)	−0.301 (−0.301–0.114)	0.027
VCAM-1	6.018 (5.902–6.084)	5.797 (5.686–5.958)	<0.001
IFNγ	0.478 (−0.301–2.211)	0.756 (−0.301–2.010)	0.507
TNF-α*	0.000 (0.000–6.766)	0.000 (0.000–8.102)	0.914
TAT	1.191 (0.736–1.619)	0.638 (0.534–0.722)	<0.001
D-Dimer	2.606 (2.348–2.784)	2.971 (2.576–3.406)	0.038

[a]TNF-α was not log transformed due to prevalence of zero values, and not included in the correlation analyses below.

Table 3

Association between biomarkers and prior small vessel stroke in African-Americans.

Biomarker	Unadjusted		Adjusted[a]	
	Odds Ratio	P value	Odds Ratio	P value
IL-1ra	3.024 (1.419–6.442)	0.004	1.932 (0.868–4.301)	0.107
IL-6	3.655 (1.378–9.698)	0.009	2.741 (0.810–9.269)	0.105
IL-8	1.987 (0.883–4.470)	0.097	1.841 (0.788–4.301)	0.158
IL-10	2.597 (1.018–6.629)	0.046	1.720 (0.571–5.179)	0.335
IFNγ	1.040 (0.685–1.579)	0.853	0.741 (0.435–1.262)	0.270
VCAM-1	274.710 (10.559–>999,999)	>0.001	722.715 (14.457–>999.999)	0.001
TAT	6.399 (1.904–21.506)	0.003	3.573 (1.010–12.638)	0.048

[a]Adjusted for age (continuous), sex, history of hyperlipidemia, and hypertension

Table 4

Correlation (Spearman r coefficients) amongst serum cytokine and thrombosis markers in the entire cohort (cases and controls, n=67)[a].

	IL-6	IL-8	IL-10	IFNγ	VCAM-1	TAT	D-Dimer
IL-1ra	0.419[†]	0.121	0.464[*]	0.322[‡]	0.409[†]	0.466[†]	−0.202
IL-6		0.465[‡]	0.229	0.508[*]	0.346[†]	0.262[‡]	−0.069
IL-8			0.178	0.114	0.314[‡]	0.347[†]	0.062
IL-10				−0.002	0.445[†]	0.390[†]	−0.289[‡]
IFNγ					0.023	0.036	0.089
VCAM-1						0.386[†]	−0.276[‡]
TAT							−0.221[‡]

[a]Statistical significance of correlations is indicated as:

[*]p<0.0005,

[†]p<0.005,

[‡]p<0.05