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### **PET Imaging of Regional vs Global Myocardial Sympathetic Activity to Improve Risk Stratification in Patients with Ischemic Cardiomyopathy**

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#### **Abstract**

**Background:** Current risk assessment approaches fail to identify the majority of patients at risk of sudden cardiac arrest (SCA). Non-invasive imaging of the cardiac sympathetic nervous system (SNS) using SPECT and PET offers the potential for refining SCA risk assessment. While various [<sup>11</sup>C]meta-hydroxyephedrine (HED) quantification parameters have been proposed, it is currently unknown whether regional denervation or global innervation yields greater SCA risk discrimination. The aim of the study was to determine whether the global innervation parameters yield any independent and additive prognostic value over the regional denervation alone.

**Methods:** In a post hoc competing risks analysis of the PAREPET trial, we compared global innervation and regional denervation parameters using the norephinephrine analog HED for SCA risk discrimination. Patients with ischemic cardiomyopathy (n=174) eligible for an implantable cardiac defibrillator (ICD) for the primary prevention of SCA were recruited into the trial. HED uptake and clearance rates were measured to assess global (LV mean) retention index (RI) and volume of distribution (DV). Regional defects were quantified as the percentage of the LV having values <75% of the maximum.

**Results:** During a median follow-up of 4.2 years, there were 56 cardiac related deaths, of which 26 were SCAs. For any given regional denervation volume there was substantial heterogeneity in global innervation scores. Global RI and DV did not decrease until regional defects exceeded 40%

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LV. Global scale parameters, RI and DV (AUC=0.61, p=0.034, p=0.046, respectively), yielded inferior SCA risk discrimination compared to regional heterogeneity (AUC=0.74).

**Conclusion:** Regional denervation volume has superior cause-specific mortality prediction for SCA versus global parameters of sympathetic innervation. These results have widespread implications for future cardiac sympathetic imaging, which will greatly simply innervation analysis.

#### **Introduction**

Sudden cardiac arrest (SCA) remains a lethal and unpredictable event, accounting for half of all heart related deaths.<sup>1</sup> Survivors of myocardial infarction (MI) have a four-fold elevated risk of arrhythmic death compared to persons without a history of MI.<sup>2</sup> Clinically, ejection fraction (EF) is the main parameter used to delineate risk and guide the need for primary prevention ICD therapy (EF  $35\%$ ).<sup>3</sup> Yet, the long-term appropriate discharge rate among these patients is only 11-35% over a 3-4 year period.<sup>4-6</sup> Furthermore, despite lower rates of SCA in patients with LVEF>35%, this population actually accounts for the majority of SCA cases.<sup>7</sup> Non-invasive imaging of the cardiac sympathetic nervous system (SNS) has been proposed to compliment SCA risk assessment. This may not only enable more cost-effective patient selection and improved outcomes, but also provide a way to identify the large number of patients that develop SCA with an EF > 35%.

Alterations in the sympathetic nervous system have been implicated in the development of lethal ventricular arrhythmias. $8-14$  These can arise from global alterations in myocardial sympathetic tone without structural alterations in sympathetic nerve density. In addition, since sympathetic nerves are exquisitely sensitive to ischemia and infarction, regional defects can develop with the zone of denervation often extending beyond the ischemic penumbra.<sup>15</sup>

Both positron emission tomography (PET) and single photon emission computed tomography (SPECT) tracers have been developed to characterize the cardiac sympathetic nervous system. Of them, 123I-meta-iodobenzylguanidine (MIBG) for SPECT and planar imaging and  $[11C]$ *meta*-hydroxyephedrine (HED) for PET have been most widely studied in large prospective trials. The ADMIRE-HF trial used planar MIBG imaging for predicting cardiac events in patients with LVEF<35%.12 The heart-to-mediastinum (H/M) ratio was assessed, serving as a global index of SNS dysfunction. In this trial, a H/M<1.6 portents a worsening prognosis. In contrast, the PAREPET trial (Prediction of ARrhythmic Events with Positron Emission Tomography) demonstrated that regional sympathetic denervation volume, assessed using HED PET, predicts SCA independently of LVEF and infarct volume. <sup>15</sup> The superior spatiotemporal resolution and well validated attenuation correction for PET enables the separate quantification of global scale and regional heterogeneity.

Various methods of HED quantification for positron emissions tomography (PET) have been proposed to assess the burden of sympathetic denervation and dysfunction. While these methods may require varying degrees of quantitative sophistication and imaging time, there remains a paucity of research evaluating which method yields the most robust SCA risk discrimination. Quantification of both global (i.e. total tracer uptake and retention) and

regional (i.e. defect volume) sympathetic innervation have been reported in the literature, but lack direct comparisons with prognosis data. The PAREPET trial calculated a regional uptake defect score, expressed as percentage of the total left ventricle (LV).15 In contrast, other studies report global parameters with HED quantification of tracer retention index (RI) and/or volume of distribution (DV). While there is some debate as to which better reflects global cardiac sympathetic innervation<sup>16-20</sup>, initial reports suggest DV may be less flowdependent and thus may be a more relevant parameter, particularly for patients with ischemic cardiomyopathy.20 Conceptually, patients with a given defect size may have varying degrees of global LV uptake. Mechanistically, this may be due to differing defect severities, heterogenous uptake in remote regions or a global reduction in tracer uptake. To date, no study has determined the prognostic significance of global tracer uptake parameters, independent of defect size and whether they offer any additive value in SCA risk prediction.

The objectives of the present analysis were two-fold: 1) in a competing-risk analysis compare the robustness of regional and global  $[{}^{11}$ C]HED quantification parameters for predicting cause-specific mortality from SCA and 2) determine whether the global RI and DV assessments yield any independent and additive prognostic value over the regional defects alone.

#### **Methods**

The data in the present analysis will not be made available to other researches as this was not included in the original patient consent form in the PAREPET trial.

#### **Study Population**

The PAREPET trial ([NCT01400334\)](https://clinicaltrials.gov/ct2/show/NCT01400334) enrolled 204 subjects with ischemic cardiomyopathy, eligible for an ICD for primary prevention of SCA. <sup>15</sup> The University of Buffalo Institutional Review Board approved the study protocol and all subjects signed an informed consent form. Of the 204 participants, 190 patients underwent HED PET scans. N=16 were excluded from the present study due to reasons listed in Figure 1, leaving n=174 included for analysis.

#### **PET Imaging**

PET imaging was performed as described in prior methods.<sup>15,21</sup> An intravenous bolus of 740 MBq (20 mCi) of HED was administered, followed by dynamic cardiac PET imaging for 60 minutes (6×30 sec, 2×60 sec, 2×150 sec, 2×300 sec, 2×600 sec, 1×1200 sec frames). The images were reconstructed with all corrections enabled using filtered back-projection and a Hann window of the Ramp filter, resulting in  $\sim$ 10 mm isotropic resolution. Dynamic Images were analyzed using FlowQuant® v2.4 (University of Ottawa Heart Institute, Ottawa, Canada).

#### **HED Parameters**

**Retention Index—**Retention index is a measure utilized to estimate the net uptake of HED at scan end-time, calculated using the following equation which relates the activity in myocardial tissue  $(C_m)$  to the unchanged tracer activity in plasma  $(C_p)$  and the function evaluation time  $(T_R)$ :

$$
RI_{(RC)} = \frac{C_m(T_R)}{\int_0^T R_{C_p(t)dt}} \times 100\,\%
$$
 (1)

No partial-volume recovery correction was applied  $(RC=I)$  as fractional blood volume is not estimated using the retention model.

**Distribution Volume—**The distribution volume is defined as the ratio of tracer concentration in the myocardium to arterial plasma, at equilibrium. For receptor ligands, the DV physiologically reflects tracer binding affinity (i.e.  $B_{max}/K_d$ ), where  $B_{max}$  is Uptake1 density and  $1/K_d$  is the affinity of HED for the Uptake1 transporter. <sup>22</sup> Volumes of distribution were calculated using a one-tissue compartment model as previously described. <sup>23</sup> This method can be expressed by the following equation, where  $C_m$  is myocardial tracer concentration,  $C_p$  is arterial plasma tracer concentration, and  $T_E$  is the time to equilibrium:

$$
DV = \frac{K_1}{k_2} = \frac{C_m(T_E)}{C_p(T_E)}
$$
(2)

Quantitative analysis requires subtraction of radiolabeled blood metabolites accumulating in the blood stream to ensure accurate computation. The arterial whole-blood time-activity curve was obtained from a 20 mm<sup>3</sup> region in the LV cavity. The following equation was used to correct for blood metabolites:  $C_p(t) = C_{wb}(t) \times pfp(t)$ , where  $C_p(t)$  is the unchanged parent tracer concentration in plasma,  $C_{wb}(t)$  is the arterial whole-blood tracer concentration, and  $pfp(t)$  is the parent fraction in plasma function.<sup>18,24</sup>

No partial volume recovery correction ( $RC = I$ ) was applied. The fraction of blood volume  $(f_{wb})$  within myocardial tissue was estimated together with the one-tissue compartment model parameters  $(K_1, k_2)$  as follows:

$$
C_m(t) = f_{wb} \times C_{wb}(t) + RC \times C_p(t) * K_1 e^{-k_2 t}
$$
\n<sup>(3)</sup>

where  $*$  is the mathematical convolution operator.

**Regional Heterogeneity—**Regional defect scores based on tracer normalized uptake (NUDS) and distribution volume (DVDS) reflect the extent  $\times$  severity of relative defects, expressed as a percent of the left ventricle.<sup>25</sup> The defect scores were calculated by quantifying tracer uptake or DV in 496 myocardial sectors; those with 75% relative to the peak tracer activity were defined as normal and set to equal 100%. The defect score was then calculated as 100% minus the average polar-map value.

#### **Clinical Endpoint**

The primary PAREPET endpoint was SCA, including ICD discharge for fast ventricular tachycardia (>240 bpm) or ventricular fibrillation as an equivalent. The secondary endpoint in the competing risks analysis was total cardiac mortality, including both SCA and nonsudden cardiac arrest. Events were arbitrated blindly by 3 board-certified cardiologists.<sup>15</sup>

#### **Statistical Analysis**

Continuous variables were represented as mean±SD. Correlations between regional heterogeneity and global scale parameters were performed using a Pearson's Correlation coefficient. A D'Agostino-Pearson omnibus normality test was first performed on the data. For the PET parameters, both regional heterogeneity parameters (NUD score and DV defect score) were normally distributed. For the global scale parameters, DV was normally distributed but RI was not. In figure 2, we used a one-way analysis of variance (ANOVA), or a Kruskal-Wallis test to assess statistical differences in global parameters at increasing severity of regional uptake defect values, where appropriate. The degree of agreement/ disagreement between the PET parameter tertiles was assessed using the weighted kappa. Time-to-endpoint was analyzed using the Kaplan-Meier method and differences between tertiles were assessed by the Log-Rank test. A Cox-proportional hazards regression was also used to assess the association between the HED PET parameters and SCA. A competing risks analysis was performed to simultaneously assess the HED PET parameters associated with SCA and NSCD. In the aforementioned analysis, a multivariate analysis was also performed with global scale values (RI or DV) and the Normalized Uptake Defect Score entered as predictors to identify any independent value of global quantification. Area under (AUC) the receiver operator characteristic (ROC) curve was used to assess the ability of the individual HED PET parameters to discriminate between patients who developed SCA versus those who did not. For this analysis, discrimination was compared at the median follow-up time of 4.2 years. We also evaluate the incremental value of global scale parameters when added to regional defect parameters using the methods of Delong, Delong, and Clarke-Pearson 26. In the original PAREPET trial, a 4 variable clinical model was created consisting of i) the HED uptake defect score, ii) creatinine, iii) indexed LVEDV, and iv) no ACE/ARB therapy. 15 Stepwise selection was used to generate this multi-variate model to predict time to SCA from PET using the Cox proportional hazard model. From this model, optimized cut points were derived (NU defect score >37% of LV, creatinine >1.49 mg/dL, LVEDVI >99 mL/m<sup>2</sup>, and no ACE/ARB therapy). Patients with none of these risk factors had a SCA risk of  $\langle 1\% \rangle$ year, whereas patients with 2 risk factors had a SCA event rate of 11.7%/year. In this clinical model, we scrutinized the ability of the global scale HED parameters to re-classify patients across these event thresholds. The incremental additive value of the global parameters were assessed using the net reclassification index (NRI) and the integrated discrimination improvement  $(ID).^{27}$  The IDI evaluates reclassification as a continuous outcome across the spectrum of risk, whereas NRI is defined by prespecified thresholds, chosen *a priori*. P<0.05 was considered statistically significant. Statistical analysis and graphical representation were performed using SAS 9.4 (SAS Institute, NC), and GraphPad Prism 6.0 (GraphPad, La Jolla, CA, USA).

#### **Results:**

Patient demographics and characteristics appear in Table 1. Overall patients were optimally treated with 97% on β-blocker therapy, 90% on an ACE or ARB, and 99% receiving either warfarin or antiplatelet therapy. After a median follow-up of 4.2 years (IQR: 2.7-5.3 years) there were 56 cardiac related deaths (n=26 SCA, n=30 non-sudden cardiac death (NSCD)), and 25 non-cardiac related deaths (NCD).

#### **Agreement Between HED Quantification**

Four HED quantification methods were performed, two regional heterogeneity defect parameters (NUDS and DVDS) and two global scale parameters (RI and DV). To investigate the agreement between the quantification methods, patients were stratified into tertiles based on their respective regional heterogeneity defect and global scale parameters. There was good to excellent agreement within the global (weighted kappa=0.53 [95%CI 0.43,0.64]) and regional heterogeneity defect (weighted kappa=0.71 [95%CI 0.63,0.79]) quantification values (Table 2). In contrast, there was poor agreement between regional heterogeneity defect and global scale parameters tertiles (weighted kappa 0.16-0.34). Similarly, there were weak linear correlations between the global scale and regional heterogeneity defect parameters (RI:  $r^2$ =0.049; DV:  $r^2$ =0.082; Figure 2 C,D). Interestingly, for a given regional defect size, there was substantial heterogeneity in global HED DV (Figure 2A) and RI (Figure 2B) values, with no consistent reductions in these parameters until the regional defect size increased above 40% LV ( $p<0.05$ ). Representative polar maps (Figure 3) illustrate this concept. We aimed to investigate whether the global parameters offer any independent, or additive prognostic value.

#### **HED Parameters and Survival**

The HED regional defect scores and global scale parameters association with time to the cardiac end points are summarized in Table 3. Only the regional defect parameters were associated with total cardiac mortality  $(p<0.05)$ , whereas both regional defect and global scale parameters were associated with SCA ( $p<0.05$ ). None of the HED parameters were associated with non-sudden cardiac death. Kaplan-Meier curves summarize this data (Figure 4). After adjusting for competing risks, SCA was associated with increased regional defect size, NU ( $p=0.0025$ ) and DV ( $p=0.0005$ ), and decreased global scale parameters, DV (p=0.04), RI (p=0.02) as shown in Table 4. However, after statistically controlling for regional defect scores (NUDS), these global scale parameters were no longer significantly associated with SCA in the competing-risk model (Table 4). In the unadjusted competing risk analysis, the effect of each of these HED quantification parameters on probability of SCA, NSCD and non-cardiac death is illustrated in Figure 5.

#### **HED Parameters and SCA Risk prediction**

Table 5 and Figure 6 summarize the AUCs for the individual HED quantification methods for predicting SCA. Of these parameters, the regional normalized uptake defect score (NUDS) had the best SCA risk discrimination (AUC 0.74) and had a significantly higher AUC than both global RI ( $\Delta U = 0.13$ , p=0.03) and global DV ( $\Delta U = 0.12$ , p=0.04). The addition of either global parameter to the regional normalized uptake defect score did not significantly improve the AUC (Table 5). In the original PAREPET trial, a 4-variable clinical model was created consisting of i) the normalized uptake defect score, ii) creatinine, iii) indexed LVEDV, and iv) no ACE/ARB therapy (Table 6).  $^{15}$  From this model, optimized cut points were derived (NUDS > 37% of LV, creatinine > 1.49 mg/dL, LVEDVI > 99 mL/m<sup>2</sup>, and no ACE/ARB therapy). Patients with none of these risk factors had a SCA risk of <1%/year, whereas patients with 2 risk factors had a SCA event rate of 11.7%/year. In this clinical model, we scrutinized the ability of the global scale HED parameters to re-classify patients

across these event thresholds. Global Scale RI and DV did not significantly improve the model and appropriately reclassify patients in accordance with their SCA risk (NRI and IDI not significant, Table 7).

#### **Discussion**

This study represents the first assessment of both regional heterogeneity and global scale (or intensity) of sympathetic activity measurements for predicting cause-specific mortality from SCA. In this *post hoc* analysis of the PAREPET trial, two regional parameters (NUDS and DVDS) and two global parameters (RI and DV) were investigated. Four major findings were derived in this study. 1) For a given regional defect size, there is substantial variability in global scale quantification values. We further assessed whether there was any independent and additive prognostic value in these global scale parameters. 2) In a competing-risk analysis, both global scale and regional heterogeneity parameters were associated with SCA risk, but not non-sudden cardiac death and non-cardiac death. 3) Global scale parameters have inferior SCA risk discrimination as compared to regional parameters and yield no additive or independent prognostic value. 4) Overall, there were no differences between uptake versus DV defect scores with respect to SCA risk prediction. These results have widespread implications for future cardiac sympathetic HED analysis and imaging protocols. Taken together we demonstrate that SCA risk stratification is sufficiently predicted with conventional HED uptake defect scores, whereas other more sophisticated quantification measures of regional heterogeneity (DV defect score) and global scale (RI and DV) may not be required for risk stratification.

#### **Relationship Between Global and Regional Sympathetic Activity**

Our results corroborate previous findings of a strong correlation between global RI and DV parameters.23 The poor correlation between global and regional parameters is novel and arises from substantial heterogeneity in global tracer uptake and retention. As a result, for a given regional defect size, there can be either global preservation or downregulation of tracer uptake. Alterations in HED uptake can arise from both anatomical and functional properties of the cardiac SNS.28 Indeed, it is not possible to differentiate the relative contribution of dysinnervation or neuronal stunning from denervation (true loss of sympathetic nerves) using a single tracer approach.<sup>28</sup> Preclinical studies indicate that sympathetic nerve function is particularly susceptible to ischemia, whereas denervation and a loss of sympathetic nerves arise in the infarct and peri-infarct risk region. In addition, sympathetic nerve sprouting occurs at the border between infarcted and normal tissue and this hyperinnervation carries significant arrhythmogenic risk<sup>29</sup>. All of these contribute to inhomogeneity in sympathetic innervation and the resulting substrate for lethal ventricular arrhythmias.<sup>30,31</sup> In contrast, global reductions in tracer uptake are affected by these regional alterations as well as elevated SNS tone, high systemic norephinephrine (NE) or a global reduction in Uptake1 (NE transporter).28,32,33 Mechanistically, the distinction between neuronal stunning and denervation may be clinically important, as the former may be reversible and less likely to induce nerve sprouting as compared to denervated myocardium.

#### **Competing Risks Analysis**

A recent competing risk analysis of the PAREPET trial identified PET and clinical variables associated with cause-specific cardiac mortality. In this analysis, SCA was correlated with greater regional HED defect score, lack of ACE/ARC therapy, elevated BNP and larger LVEDV index.13 The denervation volume was not associated with non-sudden cardiac death. <sup>13</sup> The present analysis extends these findings by comparing regional defect and global scale HED parameters in the competing risk analysis. This analysis enables the assessment of the independent components of cardiac mortality, namely SCA and NSCD. In our analysis both regional and global HED parameters were associated with cause-specific mortality from SCA, but not NSCD or NCD. However, after adjusting for regional defect size, the global RI and DV were not associated with SCA, indicating that global scale parameters offer no independent prognostic information.

#### **SCA Risk Discrimination**

In the original PAREPET trial, global RI values were not statistically different in patients that had an arrhythmic event or ICD equivalent.<sup>15</sup> These results were corroborated in a separate HED imaging study of ischemic cardiomyopathy patients  $(EF < 35\%)$ .<sup>6</sup> In the latter study, patients underwent an electrophysiological study (EPS) after ICD implantation. Global RI values were not different in patients with inducible ventricular arrhythmias (RI EPS positive 2.41 versus EPS negative 2.81, p=0.07). However, in both of these analysis regional and global HED quantification SCA risk prediction were not formally compared. We demonstrated that the global scale parameters have inferior SCA risk discrimination as compared to the regional uptake defect score used in the PAREPET trial. Previous studies have demonstrated that inhomogeneity in myocardial sympathetic innervation results in a substrate at particular risk of arrhythmogenesis.<sup>34</sup> In myocardial infarction, this arises from regional sympathetic denervation hypersensitivity and spatial alterations in action potential duration that may promote re-entrant arrhythmias.<sup>35</sup> Inhomogeneity can also arise in viable dysfunctional myocardium as a result of alterations in sympathetic nerve density, nerve sprouting and alterations in myocardial beta-receptor density and function.<sup>30,36-38</sup> There are several potential explanations why imaging this heterogeneity may be superior to assessing it indirectly with global indices of 11-CHED uptake in our analysis. First, heterogeneity in sympathetic activation was assessed by comparing reductions in 11-CHED relative to normal regions in each individual patient. In contrast, global alterations in 11-CHED uptake reflect reductions in both dysinnervated regions as well as altered uptake in normal myocardium. Uptake in normal myocardium could vary with myocardial sympathetic tone, circulating catecholamine levels and potentially pharmacological interventions that would modulate neurohormonal activation.<sup>39</sup> As a result, the same regional defect could have variable levels of global uptake in normal myocardium. Furthermore, while our analysis compared global indices to regional indices in individual patients, application of a global index clinically would require identification of a specific cut point and/or comparison to a normal data set. The variability in uptake among individuals may require a greater difference in global uptake and result in a further deterioration in predictive value. Based on these considerations, we believe that quantifying regional defects offers advantages over global indices using PET 11-CHED. Whether this extends to other norepinephrine analogs and imaging platforms will require further study.

The global scale parameters also offered no incremental discrimination as compared to the regional defect parameters alone. In a recent test-retest analysis, we demonstrated that HED regional defect parameters also have better reproducibility than global scale parameters.<sup>23</sup> Together, these results suggest regional defect parameters may be the optimal approach for routine clinical SCA risk prediction in patients with ischemic cardiomyopathy.

#### **Comparison to Previous Studies**

The present study is the first to directly compare the SCA discrimination of various HED PET imaging parameters. Numerous tracers and modalities have been studied in the context of cardiac denervation, notably MIBG with SPECT imaging. A number of studies have examined global cardiac sympathetic innervation using the tracer MIBG to predict cardiovascular outcomes in heart failure patients, but there also remains equipoise regarding the optimal parameters to predict cardiovascular risk. While global indices of tracer quantification including the heart-to-mediastinum ratio and MIBG washout rate have been studied, quantitative analysis of regional defect size with SPECT has been challenging.<sup>12</sup> The lower detection sensitivity and resolution of SPECT imaging compared to PET renders it difficult to separate regional defects from global levels of innervation.40 The ADMIRE-HF trial enrolled 961 participants with an LVEF 35% and NYHA Class II/III symptoms and examined a composite outcome of heart failure progression, potentially life-threatening arrhythmias and cardiac death.<sup>12</sup> This large study included ischemic and nonischemic cardiomyopathy patients and found the H/M ratio (similar to the global scale RI in our study) to be predictive of the composite primary endpoint. Another study (n=116), in patients with an ICD indication for both primary and secondary prevention, examined the MIBG H/M ratio, washout rate, and defect score, and found only the defect score by SPECT was predictive of ventricular arrhythmias.<sup>10</sup> This is consistent with our analysis of quantifying regional defect score and the primary endpoint of SCA. While quantifying defect volume with SPECT is challenging, technical advances such as CZT cameras may allow this in the future.

In a post-hoc analysis of the ADMIRE-HF data, Travin et al. proposed a 'bell-shaped' relationship between regional MIBG defect scores (regional parameter) and arrhythmic events, such that patients with 'intermediate-range' defects had the highest proportion of events.<sup>41</sup> This hypothesis has been subsequently supported by two separate research groups. In these studies, Verschure et al. and Nakajima et al. similarly revealed a bell-shaped between MIBG H/M ratio (global scale parameter) and combined arrhythmic death and appropriate ICD discharge.<sup>42,43</sup> To our knowledge, no such relationship has been observed for HED. In the present manuscript, Figure 4B visually hinted at such a relationship with the relationship between RI (global scale parameter) and SCA. However, there was no statistical difference between the Low and Intermediate Tertiles. In Supplemental Figure 1, we further explored the relationship between regional heterogeneity or global scale parameters and SCA on a continuous scale. Most importantly, the regional defect scores based on HED Uptake (A) or Distribution Volume (B) showed no hint of a bimodal distribution or 'bellshaped' risk profile, contrary to the MIBG SPECT data. For the global scale parameters (C, D), specifically exploring the relationship between SCA-free survival and the retention index, a few patients with relatively low values of the RI (C) appear to show a slight

improvement in survival (see red arrow), but this subsequently decreases again as the RI approaches zero. Given the very small sample size in this range (only 2 patients with RI values < 2%/min), it seems most likely that this effect is due to normal heterogeneity in the data and not a real physiological effect.

#### **Study Limitations**

While appropriate ICD therapy for VF or VT at rates of 240 BPM and faster were used as a surrogate for aborted arrhythmic death, ventricular arrhythmias at slower rates were not considered as a primary end-point. Currently there is no consensus regarding which classification of ICD discharge represents the most appropriate definition. Available evidence suggests that using the definition of the present study, the rate of ICD discharge is approximately similar to the difference in mortality rate in clinical trials comparing patients receiving a primary prevention ICD with coronary artery disease versus a medically treated population. 44 A broader analysis of ventricular arrhythmias in the PAREPET trial demonstrated that the HED defect size did not predict slower ventricular arrhythmias and the prognostic impact of HED for VF and fast VT remained.45 It is important to note that while these data have important imaging implications for patients with ischemic cardiomyopathy, the global data findings cannot be applied to patients with non-ischemic cardiomyopathy as they were not part of the present cohort.

#### **Clinical Implications**

The findings of this study may have implications for broader implementation of imaging myocardial sympathetic innervation. Current HED quantification methods have varying degrees of sophistication. Regional HED activity uptake quantification capabilities are likely available with most commercial PET software. In contrast, parametric imaging of distribution volume, which also requires kinetic modeling and long image acquisition times, may only be available to academic institutions that have access to compartmental analysis. A key finding of our study was that absolute global scale values such as RI and DV yield no independent or additive benefit clinical use in distinguishing SCA risk. This greatly simplifies future PET innervation imaging for predicting clinically meaningful outcomes and should facilitate translation into clinical care as fluorinated norepinephrine analogs become available to assess regional inhomogeneity in sympathetic innervation with PET.

#### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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#### **Abbreviations**



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#### **Figure 2.**

Relationship between regional heterogeneity and global scale [11C]HED PET parameters. For a given regional heterogeneity defect, patients display a range of global DV (A) and global RI (B) values. Correlations between the regional normalized uptake defect score (used in the PARREPET trial) and global scale parameters displayed (C,D). Excellent correlations were observed within global scale (E) and regional defect (F) [11C]HED PET parameters. DV, distribution volume; RI, retention index; LV, left ventricle.



#### **Figure 3.**

Polar maps of regional normalized uptake defect score (A), and global scale retention (B) and distribution volume (C) for two patients with similar regional defect scores but differing global RI and DV scores.



#### **Figure 4.**

Kaplan-Meier curves show the incidence of cardiac endpoints for tertiles of regional heterogeneity (A) and global scale (B) [11C]HED quantification methods. Note that the coloring of the tertiles for the global scale parameters (B) is reversed to correspond with the theoretical risk. For the regional defect parameters, the low (green) and high (red) tertiles for Regional Normalized Uptake Defect Score (low: <22.87%, high 32.6%) and DV Defect Score (low:  $\langle 35\%, \text{ high} \rangle$  43%) are shown. Similarly, for the global scale parameters, the low (red) and high (green) tertiles for DV (Low: <8.7mL/g, High:  $12.2$  mL/g) and RI (Low: <3.84 %/min, High: 4.85%/min). Corresponding P values for the univariate analysis using continuous variables are in Table 3. SCA, sudden cardiac arrest; C/NS, non-sudden cardiac death.



#### **Figure 5.**

Estimated event rates from the competing risk analysis for HED PET regional defect and global scale quantification. P values are derived from the competing-risk analysis in Table 4. Figure illustrates the anticipated effect of each HED quantification parameter on SCA (red line), non-sudden cardiac arrest (blue line) and non-cardiac death (black line) within the follow-up period. All HED quantification parameters were associated with SCA after adjusting for competing risks. The effect of each PET parameter was assessed individually. LV, left ventricle; SCA, sudden cardiac arrest; NSCD, non-sudden cardiac death; NCD, noncardiac death.



#### **Figure 6.**

Comparison of Regional Heterogeneity and Global Scale [11C[HED PET parameters for SCA risk discrimination. Time-dependent AUC illustrate discrimination over time (A). PET parameters were compared up to the median follow-up (4.2years). ROC curve comparison for regional defect and global scale parameters up to median follow-up (B). Corresponding P values are displayed in Table 5. \* AUC superior than both global scale parameters, p<0.05.

**Table 1.**

Patient Characteristics Patient Characteristics



 Author Manuscript**Author Manuscript**  BMI, Body mass index; ACE, Angiotensin-converting enzyme; ARB, Angiotensin II Receptor Blocker; ASA, Acetylsalicylic acid; LV, Left ventricle; LA, Left Atrial; BMI, Body mass index; ACE, Angiotensin-converting enzyme; ARB, Angiotensin II Receptor Blocker; ASA, Acetylsalicylic acid; LV, Left ventricle; LA, Left Atrial; **Author Manuscript** 



DV, Distribution volume; DVDS, DV Defect Score; NUDS, Normalized Uptake Defect Score; RI, Retention index; CI, Confidence interval. DV, Distribution volume; DVDS, DV Defect Score; NUDS, Normalized Uptake Defect Score; RI, Retention index; CI, Confidence interval.

Univariate Associates with Cardiac Endpoints Univariate Associates with Cardiac Endpoints



DVDS, DV Defect Score. NUDS, Normalized Uptake Defect Score; DVDS, DV Defect Score.  $\frac{1}{2}$ NUDS, Normalized Uptake Defect **Author Manuscript** 

Competing Risk Model for SCA vs Non-Sudden Cardiac Death and Non-Cardiac Death. Competing Risk Model for SCA vs Non-Sudden Cardiac Death and Non-Cardiac Death.



Comparison of PET Parameter AUC for Predicting SCA up to the median follow-up (4.2 yrs) Comparison of PET Parameter AUC for Predicting SCA up to the median follow-up (4.2 yrs)



NUDS, Normalized Uptake Defect Score; DVDS, DV Defect Score NUDS, Normalized Uptake Defect Score; DVDS, DV Defect Score Author Manuscript

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LVEDV, LV End Diastolic Volume Index; ACE, Angiotensin-converting enzyme; ARB, Angiotensin II Receptor Blocker; CI, Confidence interval. LVEDV, LV End Diastolic Volume Index; ACE, Angiotensin-converting enzyme; ARB, Angiotensin II Receptor Blocker; CI, Confidence interval.

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# **Table 7.**

Net Reclassification Improvement and Integrated Discrimination Improvement for Global PET Parameters to the PAREPET Multivariate Model. Net Reclassification Improvement and Integrated Discrimination Improvement for Global PET Parameters to the PAREPET Multivariate Model.



\* (Kisk Thresholds for NRI set at 0.9%, 3.9% 11.7% for low, intermediate and high risk, respectively. These are the event rates that correspond to patients with 0,1 or >2 high risk parameters (in Table 6). Risk Thresholds for NRI set at 0.9%, 3.9% 11.7% for low, intermediate and high risk, respectively. These are the event rates that correspond to patients with 0 ,1 or >2 high risk parameters (in Table 6).