04 THE CARDIAC EXTRACELLULAR MATRIX IS REMODELLED DIVERGENTLY WITH AGE IN HEART FAILURE: A ROLE FOR ALTERED COLLAGEN DEGRADATION IN AN OVINE RAPID PACING MODEL

doi:10.1136/heartjnl-2012-302951.004

Presenter: Margaux Horn

MA Horn, HK Graham, KM Dibb & AW Trafford. Unit of Cardiac Physiology, The University of Manchester, Manchester, UK

Alterations to the amount, quality and/or distribution of the cardiac extracellular matrix (ECM) are defining features of the structural remodelling which occurs in heart failure (HF). However, whether the ECM remodelling which occurs in the aged failing heart occurs to the same extent as in the young remains to be determined.

HF was instigated in sheep aged either 18 months (young) or >8years (aged) by rapid ventricular pacing (210 bpm). HF increased LV diameter and reduced fractional shortening (measured by echocardiography) in both young and aged animals (all P<0.001), although these changes were more pronounced in the aged (P < 0.05). LV collagen content measured from picro-sirius red-stained LV sections was altered with HF in an age-dependent manner - with collagen accumulation in young HF (P<0.001) and depletion in aged HF (P<0.05). Matrix metalloproteinase-2 (MMP-2) activity determined from gelatin zymograms was enhanced with both ageing and in young HF (both P<0.05). Protein levels of tissue inhibitor of metalloproteinases (TIMPs) 3 & 4 quantified by immunoblotting were reduced in aged HF only (P<0.05). Levels of secreted protein acidic and rich in cysteine (SPARC) were increased in aged hearts compared to young controls (P<0.05) whilst serum procollagen type I C-peptide (PICP) was increased in both young failing (P < 0.05) and aged failing (P<0.01) animals, as measured using specific ELISA.

In conclusion, remodelling of the cardiac ECM in HF is agedependent. Diminished TIMP levels only in aged HF alongside enhanced collagen synthesis in HF provide a potential mechanism for this age-dependent response.

05 NOX2-DERIVED OXIDATIVE STRESS AND BONE MARROW-DERIVED HEMATOPOIETIC STEM CELL DYSFUNCTION IN MIDDLE-AGE OBESITY

doi:10.1136/heartjnl-2012-302951.005

Presenter: Daniel Meijles

DN Meijles, FHMS, J-M Li, FHMS. University of Surrey, Guildford GU2 7XH

Systematic oxidative stress is a characteristic of metabolic disorders and cardiovascular diseases associated with middle-age obesity. Bone marrow-derived multi-potent stem cells (BMSC) hold the hope in regenerating damaged tissues; however, the effect of oxidative stress on BMSC function remains unknown. In this study, we investigated the BMSC function in mouse models of middle-age obesity. Littermates of C57BL/J6 wild-type and Nox2 (an O2.-generating enzyme) knockout mice (7 m old, n=15) were fed with high fat diet (HFD, 45% kcal fat, 20% kcal protein and 35% kcal carbohydrate) or normal chow diet (NCD, 12% kcal fat, 28% kcal protein and 60% kcal carbohydrate) for 16 weeks. BMSCs were isolated from mice at 11 m of age. Compared to NCD controls, the numbers of CD133+/VEGFR2+ endothelial progenitor cells (EPC) were significantly decreased (2.2%±0.3 NCD vs. $0.8\% \pm 0.5$ HFD) in HFD mice. There were significant increases (82±9.2 %) in the levels of O2.- production by HFD BMSC, and this was accompanied with accelerated cell proliferation $(160\pm5.2\%)$, cell cycle progression from G1/G0 phase to S phase, and significant increases in cell apoptosis (6.9±2.5% NCD vs. 29.8±8.2% HFD) as examined by annexin V flow cytometry.

Moreover, the levels of PCNA and p53 expression were significantly increased in HFD BMSC. However, all these changes were absent in BMSC isolated from Nox2 knockout mice fed with HFD. In conclusion, an obesity environment activates Nox2 and oxidative stress damages BMSC function and reduces EPC population. Nox2 may present a therapeutic target for the prevention and treatment of obesity-related diseases.

06 DUAL ENERGY CT HAS THE POTENTIAL TO IMPROVE NON-INVASIVE IDENTIFICATION OF NECROTIC CORE

Presenter: Daniel Obaid

doi:10.1136/heartjnl-2012-302951.006

¹D Obaid, ¹P Calvert, ²M Goddard, ²D Gopalan, ^{1,3}J Rudd, ^{1,3}M Bennett. ¹Dept of Cardiovascular Medicine, University of Cambridge. UK; ²Papworth Hospital NHS Trust. UK; ³Addenbrookes Hospital NHS trust. UK

Background CT can classify plaque based on its x-ray attenuation. However, identifying vulnerable plaque is limited by overlap between the attenuation of necrotic core and fibrous plaque. Changes in attenuation of plaque components to x-rays of differing energies may allow better discrimination. We tested whether Dual Energy CT (DECT) (simultaneous image acquisition at two energies) improved identification of necrotic core, both in-vivo and ex-vivo.

Methods 20 patients underwent DECT and 3-vessel Virtual Histology-IVUS (VH-IVUS). Attenuation was sampled in 1088 plaque areas co-registered with VH-IVUS and used to define dual energy indices (changes in attenuation of plaque components at 100 kV and 140kV). 42 plaques were analysed by DECT to determine whether DECT increased sensitivity to detect VH-IVUS defined necrotic core. 10 post-mortem coronary arteries were also examined with DECT prior to histological analysis to determine whether DECT increased sensitivity to detect histologically proven necrotic core.

Results Dual energy indices of necrotic core and fibrous plaque were significantly different (mean: 0.0071 vs. 0.0283, p<0.05). Utilising these increased diagnostic accuracy for DECT to detect necrotic core in 87 segments of post-mortem arteries (sensitivity-64%, specificity-96%) compared with single energy CT (sensitivity-54%, specificity-92%). Sensitivity to detect necrotic core was lower in plaques analysed in-vivo due to the impact of temporal resolution on moving coronaries. However, DECT still provided marginal improvements in sensitivity (45%) compared with single energy CT (39%). **Conclusions** Dual Energy CT has the potential to improve the differentiation of necrotic core and fibrous plaque allowing more accurate non-invasive identification of vulnerable plaque.

SIMULTANEOUS POSITRON EMISSION TOMOGRAPHY AND MAGNETIC RESONANCE IMAGING OF RECEPTORS USING A NOVEL COMBINED PRE-CLINICAL MICROPET/MR SYSTEM

doi:10.1136/heartjnl-2012-302951.007

Presenter: Anthony Davenport

07

¹JLE Bird, ²RC Hawkes, ³R Manavaki, ²SJ Sawiak, ²DJ Williamson, ²FI Aigbirhio, ⁴JHF Rudd, ⁴M Bennett, ⁵JH Gillard, ²TA Carpenter, ⁶EA Warburton, ¹AP Davenport. ¹Clinical Pharmacology Unit; ²Wolfson Brain Imaging Centre; ³Department of Medicine; ⁴Division of Cardiovascular Medicine; ⁵Department of Radiology; ⁶Department of Clinical Neurosciences, University of Cambridge, Addenbrooke's Hospital Cambridge CB2 200, UK

Vascular and cardiac diseases are complex pathologies and preclinical models are required to fully investigate the multifactorial interactions. In vivo imaging techniques are important research tools in quantifying pathogenic mechanisms and positron emission tomography (PET) is an imaging modality which has the chemical specificity and the sensitivity to quantify biological processes in vivo. However, PET tracer uptake does not usually provide sufficiently detailed anatomical structure to accurately allocate receptor activity to precise tissue regions. This may be overcome by simultaneous imaging with magnetic resonance imaging (MRI). The aim of the study was to evaluate the combined PET/MRI scanner for the investigation of biological processes in rodents. N-(5-fluoro-2-phenoxyphenyl)-N-(2-[18F]-fluoroethyloxy-5-methoxybenzyl)acetamide ([18F]FEDAA1106) binds the translocator protein (TSPO) which is up-regulated in activated macrophages and may quantify vascular inflammatory pathologies. Dynamic in vivo imaging was carried out using a modified Focus F120 microPET incorporated into a bespoke 1 Tesla MR magnet (1). The pharmacokinetic profile of [18F]-FEDAA was characterised in mice. Simultaneously acquired PET and MRI reconstructed images were aligned together and the combined images revealed rapid uptake of [18F]-FEDAA into the heart, liver, lungs, kidneys and brain. Time activity curves were constructed using regions of interest delineated by the MR images and showed the expected pharmacokinetic profile. Thus, the results show that the fused anatomical-functional image not only provides anatomical context to the PET data, but can also allow improved quantification by more accurately defining the region of radioactive emission

(1) Hawkes RC, et al. Technol Cancer Res Treat. 2010; 9(1):53-60.

08 RAS-ASSOCIATION DOMAIN FAMILY 1 ISOFORM A (RASSF1A) IS A NOVEL REGULATOR OF TNF-ALPHA SIGNALLING IN CARDIOMYOCYTES

doi:10.1136/heartjnl-2012-302951.008

Presenter: Tamer Mohamed

TMA Mohamed, M Zi, A Maqsood, S Prehar, L Neyses, D Oceandy. Cardiovascular Medicine Research Group, School of Biomedicine, University of Manchester

Tumour necrosis factor-alpha; (TNF-alpha) plays key roles in the pathogenesis of heart failure. Cardiomyocytes express the TNFalpha; receptor (TNFR), however, the mechanism of TNF-alpha; signal transmission in cardiomyocytes is not completely understood. Recent studies showed that in cancer cells TNFR is regulated by Ras-association domain family 1 isoformA (RASSF1A). Therefore, we investigated whether RASSF1A modulates TNFalpha; signalling in cardiomyocytes. We used RASSF1A knockout (KO) mice and wild type (WT) controls and stimulated them with TNF-alpha; ($10\mu g/kg i.v.$). In WT mice acute treatment with low dose of TNF-alpha increased cardiac contractility and intracellular calcium transient amplitude, which is consistent with previously published data (Circulation 2004; 109:406-411). However, KO mice showed a blunted contractile response following acute TNFalpha treatment as indicated by the change in end systolic elastance (in vivo) and intracellular calcium transient amplitude (isolated adult cardiomyocytes). We also found that RASSF1A formed a molecular complex with TNF-alpha; receptor in cardiomyocytes and this interaction was essential in the recruitment of TRADD and TRAF2, the major downstream effectors of TNFalpha; signalling. By mapping the interaction domain we found that the C-terminal region of RASSF1A was responsible for the formation of TNF-alpha; receptor complex. Furthermore, using an adenoviral-mediate d shRNA construct we found that cardiomyocytes lacking RASSF1A exhibited reduced activation of NFkB, a downstream target of TNF-alpha. Overall, our data indicate an essential role of RASSF1A in regulating TNF-alpha; signalling in cardiomyocytes, with RASSF1A being key in the formation of TNF receptor complex and in the signal transmission to the downstream targets.

09 BIOMARKERS FOR DETECTION OF ACTIVATED MACROPHAGES IN ATHEROSCLEROSIS

Presenter: Joseph Bird

doi:10.1136/heartjnl-2012-302951.009

¹JLE Bird,¹L Burzynski, ²R Manavaki, ³JHF Rudd, ⁴EA Warburton, ¹AP Davenport. ¹Clinical Pharmacology Unit; ²Department of Medicine; ³Division of Cardiovascular Medicine; ⁴Department of Clinical Neurosciences, University of Cambridge, Addenbrooke's Hospital Cambridge CB2 200, UK

Atherosclerosis is the pathogenic mechanism underlying the majority of strokes, heart attacks and peripheral vascular disease. These are the principal causes of morbidity and mortality in the developed nations. The atherosclerotic lesion (or plaque) is characterised by a build up of lipid and inflammatory cells in the artery wall. Arterial stenosis is the current clinical predictor for endarterectomy, although macrophage burden in the lesion correlates strongly with likelihood of rupture and could be a better proxy of plaque vulnerability. Activated macrophages have up-regulated expression of translocator protein (TSPO), which is readily quantified by the positron emission tomography (PET)-based ligands (1) such as [11C](R)-1-(2-chlorophenyl)-N-methyl-N-(1-methylpropyl)-3isoquinoline-carboxamide ([11C](R)-PK11195). Studies examining TSPO plaque biodistribution frequently use CD68, a pan-phenotype marker, for identification of macrophage presence. However, this biomarker demonstrates some inconsistencies in these analyses (1). We hypothesise that more specific macrophage phenotype markers may perform better as correlates for TSPO PET-ligand binding. In this study, we examined the expression of phenotyperestricted macrophage markers in endarterectomy tissue, and correlated these with the distribution of [3H](R)-PK11195 using autoradiographic, immunohistochemical and immunofluorescence techniques. Autoradiographic and immunohistochemical images were digitised and co-registered to provide more accurate correlation of expression. We identified several markers of activated macrophage phenotype which have better correlation with [3H] (R)-PK11195 binding patterns compared with CD68. These data indicate that TSPO PET ligands bind principally to activated macrophage phenotypes, and the phenotype markers have the potential to more accurately identify PET-ligand binding macrophages in ex vivo analysis.

1) Bird JL et al. Atherosclerosis, (2010), 210(2): 388-91.

10 REGULATION OF MONOCYTE-ENDOTHELIAL CELL INTERACTIONS BY NEUTROPHIL-DERIVED MICROPARTICLES

doi:10.1136/heartjnl-2012-302951.010

Presenter: Amanda Burnett

A Burnett, P Hellewell, V Ridger. Department of Cardiovascular Science, University of Sheffield, Sheffield, S10 2RX

Rationale Monocytes play a major role in atherosclerosis progression through migration into the arterial wall. Neutrophil involvement in atherosclerosis was previously only thought to be via enzymatic weakening of the fibrous cap. However, neutrophil depletion can delay atherogenesis and conversely increasing circulating neutrophil enhance plaque progression in mice. Lack of evidence for the presence of neutrophils in atherosclerotic plaques makes their role in disease progression less clear. We have found neutrophil-derived microparticles increase migration of neutrophils. Our hypothesis: neutrophil-derived microparticles increase endothelial cell-monocyte interactions and facilitate monocyte transendothelial migration.

Methodology Neutrophils were incubated with various agents to stimulate microparticle formation. Microparticles were characterised and quantified using a novel, standardised flow cytometry