

Results CFD enabled measurement of WSS throughout the thoracic aorta. WSS was significantly elevated in aortic stenosis, highest in AS-BAV(RN) (mean WSS=37.1 ± 4.0 dyn/cm², compared to 19.9 ± 1.9 dyn/cm² for AS-BAV(RL), 25.7 ± 1.2 dyn/cm² in AS-TAV, 12.3 ± 3.4 dyn/cm² in AR-TAV, and 9.9 ± 5.4 dyn/cm² in healthy volunteers, *p* < 0.05). Aortic stenosis patients displayed asymmetrical WSS distributions, the greater curvature experiencing the highest WSS. OSI was lower in bicuspid right-non fusion (*p* < 0.05).

Eccentricity of flow was higher in bicuspid patients (Flow_{a-symmetry} = 84.1 ± 5.4%, compared to 28.1 ± 21.5 for tricuspid, *p* < 0.05). Helicity of flow was assessed by the Helical Flow Index (HFI), which was higher in bicuspid right-left fusion (HFI_{systole} = 0.39 ± 0.04, compared to 0.28 ± 0.03 for all others, *p* < 0.05).

Conclusions BAV displays eccentric flow with high helicity. Presence of AS, particularly in BAV-RN led to higher WSS and lower OSI in the greater curvature of the ascending aorta. Patient-specific CFD provides non-invasive functional assessment of the thoracic aorta, and enables development of a personalized approach to diagnosis and management of aortic disease beyond traditional guidelines.

184 CARDIOPROTECTIVE ROLE OF HEXARELIN IN A MOUSE MODEL OF MYOCARDIAL INFARCTION

¹Hayley McDonald*, ²Jason Peart, ¹Nyoman Kurniawan, ¹Graham Galloway, ³Chrisan Samuel, ¹Chen Chen. ¹University of Queensland; ²Griffith University; ³Monash University; *Presenting Author

10.1136/heartjnl-2016-309890.184

This study aimed to determine whether Hexarelin (HEX), a synthetic growth hormone secretagogue, preserves cardiac function and attenuates remodelling in mouse models of myocardial infarction.

Myocardial ischemia was induced by ligation of the left descending coronary artery in C57BL/6J mice followed by HEX (n = 16) or vehicle (VEH) (n = 16) administration at 0.3 mg/kg/day for 21 days. Treated and Sham mice were subjected to magnetic resonance imaging using a T₁-weighted late gadolinium enhancement sequence (LGE) at 9.4 Tesla (T) to measure left ventricular (LV) function, mass and infarct size at 24hrs and 21 days. HEX mice demonstrated a significant improvement (*P* < 0.05) in ejection fraction (EF) compared with VEH at 24 h (42% vs 34% respectively) and at 21 days (49% vs 36%).

A significant decrease in LV mass, interstitial collagen and collagen concentration was demonstrated after 21 days within the HEX group. This was accompanied by a decrease in TGF-β₁ and I ± -SMA and increase in MMP-13 in the HEX group. Furthermore, heart rate variability analysis demonstrated that HEX treatment shifted the balance of autonomic nervous activity towards a parasympathetic predominance, evidenced by a smaller low/high-frequency power ratio and increased normalised high frequency power. This was combined with a significant decrease in Troponin-I, IL1-β and TNF-α ± levels with HEX treatment compared with VEH treatment after 24 h.

These results demonstrate that GHS may preserve ventricular function and favourably remodel the process of fibrotic healing in mouse models of myocardial infarction; this may be through anti-inflammatory mechanisms.

185 GLYCOPROTEOMICS REVEALS DECORIN PEPTIDES WITH ANTI-MYOSTATIN ACTIVITY IN HUMAN ATRIAL FIBRILLATION

¹Javier Barallobre Barreiro*, ²Shashi Gupta, ¹Anna Zoccarato, ¹Rika Kitazume-Taneike, ¹Marika Fava, ¹Xiaoke Yin, ¹Anna Zampetaki, ³Alessandro Viviano, ¹Mei Chong, ⁴Marshall Bern, ³Antonios Kourliouros, ⁵Nieves Domenech, ¹Peter Willeit, ¹Ajay M Shah, ³Marjan Jahangiri, ⁶Liliana Schaefer, ⁷Jens W Fischer, ⁸Renato V Iozzo, ⁹Rosa Viner, ²Thomas Thum, ¹⁰Joerg Heineke, ¹¹Antoine Kichler, ¹Kinya Otsu, ¹Manuel Mayr. ¹King's College London; ²Institute for Molecular and Translational Therapeutic Strategies, MH-Hannover; ³St Georges Hospital, NHS Trust; ⁴Protein Metrics; ⁵Biobanco a Corua, INIBIC-Complejo Hospitalario Universitario de A Corua; ⁶Klinikum Der Goethe-Universität; ⁷Institute for Pharmacology and Clinical Pharmacology, Heinrich-Heine-University; ⁸Sidney Kimmel Medical College at Thomas Jefferson University; ⁹Thermo Fisher Scientific; ¹⁰Experimental Cardiology, Department of Cardiology and Angiology, MH-Hannover; ¹¹Laboratoire Vecteurs: Synthèse Et Applications ThA©Rapeutiques, A UMR 7199A CNRS Université de Strasbourg; *Presenting Author

10.1136/heartjnl-2016-309890.185

Background Myocardial fibrosis is a feature of many cardiac diseases. We used proteomics to profile glycoproteins in the human cardiac extracellular matrix (ECM).

Methods and Results Atrial specimens were analysed by mass spectrometry after extraction of ECM proteins and enrichment for glycoproteins or glycopeptides. Out of all ECM proteins identified, the small leucine-rich proteoglycan decorin was found to be the most fragmented. Within its protein core, eighteen different cleavage sites were identified. In contrast, no cleavage was observed for biglycan, the most closely related proteoglycan. Decorin processing differed between human ventricles and atria and was altered in disease. The C-terminus of decorin, important for the interaction with connective tissue growth factor, was predominantly detected in ventricles compared to atria. In contrast, atrial appendages from patients in persistent atrial fibrillation had higher levels of full-length decorin but also harboured a cleavage site that was not found in atrial appendages from patients in sinus rhythm. This cleavage site preceded the N-terminal domain of decorin that controls muscle growth by altering the binding capacity for myostatin. Myostatin expression was decreased in atrial appendages of patients with persistent atrial fibrillation and hearts of decorin null mice. A synthetic peptide corresponding to this decorin region dose-dependently inhibited the response to myostatin in cardiac myocytes and in perfused mouse hearts.

Conclusion This proteomics study is the first to analyse the human cardiac ECM. Novel processed forms of decorin protein core, uncovered in human atrial appendages can regulate the local bioavailability of anti-hypertrophic and pro-fibrotic growth factors.

186 PLATELETS AS KEY REGULATORS OF FIBRIN-CLOT ARCHITECTURE AS ASSESSED BY FRACTAL ANALYSIS OF VISCOELASTIC PROPERTIES; EFFECTS OF STANDARD ANTI-PLATELET THERAPIES

¹Rebecca Knowles*, ²Lindsay D'Silva, ¹Paul Armstrong, ¹Melissa Chan, ²Matthew Lawrence, ²Sophie Standford, ²Ahmed Sabre, ¹Arthur Tucker, ²Phylip Williams, ¹Timothy Warner, ²Phillip Evans. ¹WHRI; ²NISCHR Haemostasis Biomedical Research Unit, ABMU Health Board; *Presenting Author

10.1136/heartjnl-2016-309890.186

Introduction Platelets are critical drivers of thrombus formation with high on-treatment platelet reactivity despite dual

antiplatelet therapy (DAPT) representing a risk factor for thrombosis. Coagulation pathways also play a central role and configuration of the fibrin network is a vital determinant of clot stability and outcome, with dense clots composed of compact thin fibres associated with thrombotic events. However, despite the interwoven nature of platelet activation and the coagulation system, studies relating both aspects of thrombosis in the same population are limited.

Methods Four groups of healthy volunteers ($n = 8$) received single anti-platelet therapy aspirin (75 mg) or prasugrel (10 mg) or DAPT with aspirin (75 mg) plus prasugrel (10 mg) or aspirin (75mg) plus ticagrelor (90mg) for 7 days. In this study, we characterised the influences of these standard anti-platelet therapies on platelet function using standard tests of platelet activation and aggregation. We then associated platelet reactivity to fibrin clot microstructure which was determined using advanced rheological techniques to analyse the viscoelastic properties of incipient clots; fractal dimension (D_f) and mean relative mass (RM).

Results Aspirin alone caused inhibition of platelet responses to arachidonic acid (AA), but did not reduce ATP release, P-selectin or PAC-1 binding. Aspirin also had no effect upon the measures of clot structure D_f (1.71 ± 0.01 to 1.69 ± 0.01 , $p = 0.41$) or RM ($-8 \pm 19\%$, $p = 0.47$). Prasugrel significantly reduced D_f (1.72 ± 0.02 to 1.67 ± 0.01 , $p = 0.03$) and RM ($-40 \pm 11\%$, $p = 0.03$), as well as caused a significant decrease in ATP release, PAC-1 and p-selectin expression ($p < 0.05$). Both aspirin plus prasugrel and aspirin plus ticagrelor inhibited platelet responses to all agonists tested and decreased P-selectin expression. Significant platelet inhibition was associated with a reduction in D_f ; 1.73 ± 0.02 to 1.68 ± 0.02 ($p = 0.03$) and 1.72 ± 0.03 to 1.62 ± 0.02 ($p = 0.04$) by aspirin+prasugrel and aspirin+ticagrelor, respectively. This corresponded to reductions in RM both for aspirin plus prasugrel ($-35 \pm 16\%$ change, $p = 0.04$) and for aspirin plus ticagrelor ($-45 \pm 14\%$ change, $p = 0.04$).

Conclusion We demonstrate that platelets are important determinants of clot microstructure which is modifiable by anti-platelet therapies. These therapies may rely on their abilities to reduce thrombus density to exert their therapeutic effects with increased levels of platelet inhibition associated with the formation of more open, porous fibrin clots. We also suggest that it is activation of P2Y₁₂ receptors rather than TX (A₂) receptors that are key to clot formation. We conclude that there lies both diagnostic and therapeutic potential in determining the functional relationship between platelet reactivity, eventual clot quality and clinical outcome and D_f could be a useful novel biomarker in risk stratification and tailored anti-platelet therapies.

187

MITOCHONDRIAL DYSFUNCTION IN THE DIABETIC HEART A€ " IMBALANCE IN FUSION/FISSION AXIS?

Lucy Murfitt*, Mohammad M Monib, Hayley Bennett, Bernard Davenport, Christian Pinali, Garth Cooper, Elizabeth Cartwright, Ashraf Kitmitto. *University of Manchester*; *Presenting Author

10.1136/heartjnl-2016-309890.187

Background Cardiovascular disease is the leading cause of morbidity and mortality among diabetic patients. Diabetic cardiomyopathy is closely linked to mitochondrial dysfunction, however the pathophysiological mechanisms responsible are not known. Maintenance of mitochondrial function relies on

the balance between fusion and fission events. The fusion protein mitofusin-2 (Mfn2) has been implicated in the pathogenesis of diabetes. Alongside fusion, Mfn2 is widely believed to function as a molecular tether, binding mitochondria to the sarcoplasmic reticulum (SR) to form specialised Ca²⁺ microdomains. Nonetheless, the role of Mfn2 in the heart is poorly characterised. Therefore, the aim of this study was to investigate changes to cardiac mitochondrial protein expression and function in diabetes with a particular focus upon the fusion/fission axis.

Methods and results Protein expression levels were measured in control and streptozotocin-treated (STZ) Wistar rat heart using Western Blot. Mitochondrial OXPHOS function was assessed using enzyme activity assays. Lastly, changes to the mitochondrial proteome were investigated using Mass Spectrometry (MS). Western Blot showed a significant increase in Mfn1 and Mfn2 expression levels in STZ compared to controls with no change to the fission protein Drp1. Enzymatic assays revealed that mitochondrial function was altered in the STZ rat heart compared to control. Lastly, MS identified 1437 proteins, of which there was an upregulation of proteins involved in beta oxidation in the STZ compared to controls. In contrast, there was a downregulation of proteins associated with OXPHOS in the STZ suggesting mitochondrial dysfunction that corroborates the functional data.

Conclusion These data suggest that mitochondrial dysfunction may be linked to an imbalance of the mitochondrial fusion/fission axis in the diabetic heart. Future work will focus on the 3-D reconstruction of the mitochondrial networks using electron microscopy to determine whether changes to mitochondrial function are linked to structural alterations. These studies will enhance our understanding of the pathogenesis of cardiac mitochondrial dysfunction in diabetes, with the hope to elucidate potential targets for therapeutic intervention.

188

CHANGES IN PERIVASCULAR ADIPOSE TISSUE IN AGEING AND ATHEROSCLEROTIC MICE

¹Rachel Walker*, ²Clare Austin, ¹Cathy Holt. ¹University of Manchester; ²Edge Hill University; *Presenting Author

10.1136/heartjnl-2016-309890.188

Perivascular adipose tissue (PVAT) encases the majority of blood vessels and is an important component of the vasculature. PVAT is an active endocrine organ but until recently was not considered important in the pathogenesis of atherosclerosis. In healthy vessels PVAT exerts an anti-contractile effect which is attenuated in states of metabolic disturbance. The following experiments were conducted to determine the influence of PVAT, age and progression of atherosclerotic disease on isolated arterial reactivity.

Upon weaning, male ApoE^{-/-} or C57/BL6 (control) mice were fed a standard rodent chow diet for a period of eight or twenty-six weeks. Aortic atherosclerotic lesion area was analysed by en face quantification with Oil Red O. Aortic PVAT was collected and weighed. Contractile responses of the thoracic aortae to cumulative doses of phenylephrine (1×10^{-10} – 3×10^{-5} M) were measured in PVAT intact or PVAT denuded vessels using myography. Adipokine expression from PVAT was examined using an adipokine proteome profiler.