208

PREDICTION OF CONTRAST INDUCED NEPHROPATHY USING NOVEL BIOMARKERS FOLLOWING ELECTIVE CONTRAST CORONARY ANGIOGRAPHY

¹Michael Connolly*, Michelle Kinnin, David Mc Eneaney, Ian Menown, Neal Morgan, ²Mark Harbinson. ¹Craigavon Area Hospital; ²Queens University Belfast; *Presenting Author

10.1136/heartjnl-2016-309890.208

Introduction Chronic Kidney Disease (CKD) is a risk factor for contrast induced nephropathy (CIN), defined as an increase in serum creatinine of >25% from baseline or a delta rise of $>26.5~\mu \text{mol/L}$ within 48 h. Early diagnosis of CIN requires validated novel biomarkers.

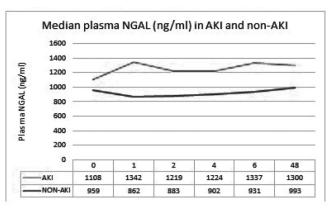
Methods A prospective observation study of 301 consecutive CKD patients undergoing elective invasive coronary angiography was performed. Low-osmolar contrast was standard. Demographics and Mehran risk score were recorded. Samples for plasma neutrophil gelatinase-associated lipocalin (NGAL), serum liver fatty acid-binding protein (L-FABP), serum kidney injury marker 1 (KIM-1), serum interleukin 18 (IL-18) and serum creatinine were taken at 0, 1, 2, 4, 6 and 48 h post contrast. Urinary NGAL and urinary cystatin C (CysC) were collected at 0, 6 and 48 h. Incidence of major adverse clinical events (MACE); acute myocardial infarction, heart failure hospitalisation, stroke and death were recorded at 1 year.

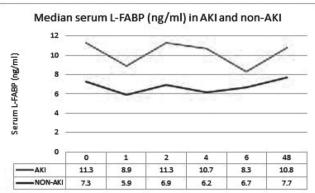
Results CIN occurred in 28 (9.3%) patients and were independently associated with older age, diabetes, higher Mehran score, larger contrast volume and anaemia (p < 0.05). Logistic regression analysis showed diabetes, CKD stage and GFR to be most predictive of CIN. The predictive power of plasma NGAL was greatest at 6 h with median levels of 1,337 ng/ml in CIN patients compared with 931 ng/ml in non-CIN patients (p = 0.002, AUC 0.71, sensitivity 75.0%, specificity 96.1%, OR 2.86), see figure 1 and table 1. L-FABP performed best at 4 h with median levels of 10.7 ng/ml in CIN patients compared with 6.2 ng/ml in non-CIN patients, p = 0.001, AUC 0.69, sensitivity 42.3%, specificity 90.2%, OR 6.75, Figure 1 and Table 1. Median urinary NGAL was higher only after 48 h, 487 ng/ml in CIN patients versus 155 ng/ml in non-CIN patients, p = 0.008, AUC 0.63. CysC, IL-18 and KIM-1 were not predictive at any time-point (p > 0.05). A Mehran score â%%¥10 performed prior to procedure achieved an AUC of 0.65, p = 0.006. MACE occurred in 7 (25.0%)

Abstract 208 Table 1 Summary of NGAL (ng/ml) and L-FABP (ng/ml) in AKI and non-AKI patients

Biomarker	Time (hr)	Median CIN	Median non-CIN	AUC	P value
	0	1108	959	0.62	0.046
0.000	2	1219	883	0.68	0.004
NGAL	4	1224	902	0.65	0.014
	6	1337	931	0.71	0.002
L-FABP	0	11.3	7.3	0.65	0.011
	2	11.3	6.9	0.67	0.007
	4	10.7	6.2	0.69	0.001
	6	8.3	6.7	0.63	0.045

Sensit: sensitivity; specif: specificity; PPV: positive predictive value; NPV: negative predictive value; RR: relative risk; OR: odds ratio





Abstract 208 Figure 1 Median plasma NGAL (ng/ml) and serum L-FABP (ng/ml) in AKI and non-AKI

CIN patients but only 17 (6.2%) non-CIN patients (p < 0.001). CIN cases also had considerably higher mortality (10.7% compared to 3.3%, p = 0.037). Exploratory analysis showed that the combination of Mehran score >10, 6 hr NGAL and 4 hr L-FABP improved specificity to 96.7%. Figure 2 highlights a proposed pathway of how biomarkers could be used to identify CIN early and facilitate timely therapeutic intervention to reduce morbidity and mortality.

Conclusions/implications Mehran risk score, 6 h plasma NGAL and 4 h serum L-FABP performed best at early CIN prediction. CIN patients were four times more likely to develop MACE and had a trebling of mortality risk at 1 year. The implications of our results, translated to the design of safer elective coronary intervention services able to more efficiently manage the increasing volume of contrast studies, should be a key health priority for cardiac and renal services.

209 WHOLE EXOME SEQUENCING IDENTIFIES GENETIC CAUSE OF HISTIOCYTOID CARDIOMYOPATHY

¹Gillian Rea*, ²Tessa Homfray, ²Jan Till, ²Ferran Roses-Noguer, ¹Rachel J Buchan, ¹Sam Wilkinson, ¹Roddy Walsh, ³Shane McKee, ³Fiona J Stewart, ⁴Victoria Murday, ⁵Robert W Taylor, ²A John Baksi, ²Sanjay K Prasad, ¹Paul JR Barton, ¹James S Ware, ⁷Stuart A Cook. ¹Imperial College London & Royal Brompton and Harefield NHS Foundation Trust; ²Royal Brompton & Harefield NHS Foundation Trust; ³Northern Ireland Regional Genetics Service; ⁴Department of Clinical Genetics, Laboratory Medicine, The Queen Elizabeth University Hospital; ⁵Wellcome Trust Centre for Mitochondrial Research; ⁶National Heart Centre; *Presenting Author

10.1136/heartjnl-2016-309890.209