

Association of Elevated Urinary miR-126, miR-155 and miR-29b with Diabetic Kidney Disease



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Introduction

Diabetes Mellitus affects 3.8 million people in the UK

Approx. 40% diabetic patients go on to develop diabetic kidney disease (DKD) the incidence of which is increasing globally

Stage	Description	GFR (ml/min/1.73m ²)
1	Kidney damage with normal or ↑GFR	≥ 90
2	Kidney damage with mild ↓GFR	60 - 89
3	Moderate ↓GFR	30 - 59
4	Severe ↓GFR	15 – 29
5	Kidney failure	< 15 (or dialysis)



predictive The value OŤ current non-invasive prognostic indicators for









chronic kidney disease, such urine protein as quantification and declining eGFR is limited

UK Renal Registry 2008

Methods



Results

Biomarker candidates



MicroRNA mechanisms





Volcano plot: analysis of 4 pooled urine samples from 5 DKD (n = 20) and 4 pooled urine samples from 5 controls (n = 20)



Individual RT-qPCR analyses of miRs in DKD vs controls



Localization miRNA expression by Of capture laser microdissection (LCM) and in vitro models







Validation: Detection of selected microRNAs in healthy, diabetic DKD independent cohorts (n=151).



MiRNA expression in glomerular endothelial cells (CiGenC) and CiGenC growth media under diabetic conditions+/-cytokines, showing the release of miR-126 and 29b into the conditioned medium in response to cytokines

Conclusions

MicroRNAs show great promise as potential predictive biomarkers in CKD

Our targets show protective properties being released from the glomerulus under disease conditions in vitro



Acknowledgements

