

Impact of diabetic stress conditions on renal cell metabolome – Supplementary material

Simon Lagies^{1,2,3*}, Roman Pichler^{4*}, Tillmann Bork⁴, Michael M. Kaminski⁴, Kevin Troendle⁵, Stefan Zimmermann⁵, Tobias B. Huber⁶, Gerd Walz^{4,7}, Soeren S. Lienkamp^{4,7,8}, Bernd Kammerer^{1,7*}

¹Center for Biological Systems Analysis (ZBSA), Albert-Ludwigs-University Freiburg, Habsburgerstr. 49, 79104 Freiburg, Germany

²Spemann Graduate School of Biology and Medicine (SGBM), University of Freiburg, Albertstr. 19a, 79104 Freiburg, Germany

³Faculty of Biology, University of Freiburg, Schänzlestr. 1, 79104 Freiburg, Germany

⁴Department of Medicine, Renal Division, Medical Center - University of Freiburg, Faculty of Medicine, University of Freiburg, Hugstetter Str. 55, 79106 Freiburg, Germany

⁵Laboratory for MEMS Applications, IMTEK - Department of Microsystems Engineering, University of Freiburg, Georges-Koehler-Allee 103, 79110 Freiburg, Germany

⁶III. Department of Medicine, University Medical Center Hamburg-Eppendorf, 20246 Hamburg, Germany

⁷BIOSS Centre of Biological Signalling Studies, University of Freiburg, Schänzlestr. 18, 79104 Freiburg, Germany

⁸Institute of Anatomy, University of Zurich, Winterthurerstr.190, 8057 Zurich, Switzerland

* Equal contribution

* Corresponding author:

Prof. Dr. Bernd Kammerer

Email: bernd.kammerer@zbsa.uni-freiburg.de

Center for Biological Systems Analysis

Albert-Ludwigs-University Freiburg

Habsburgerstraße 49

79104 Freiburg i. Br.

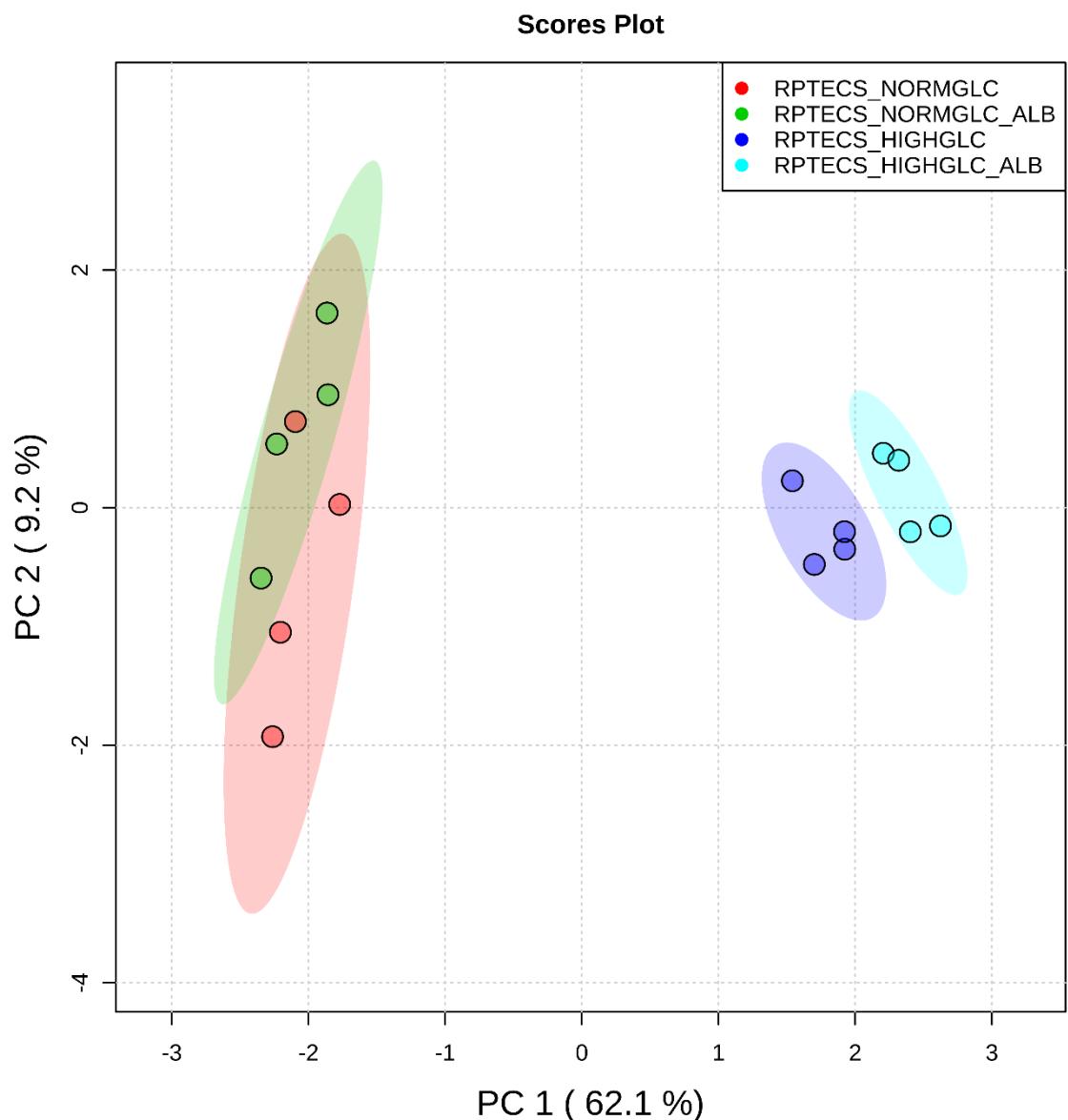
Germany

Tel: 0761-203-97137

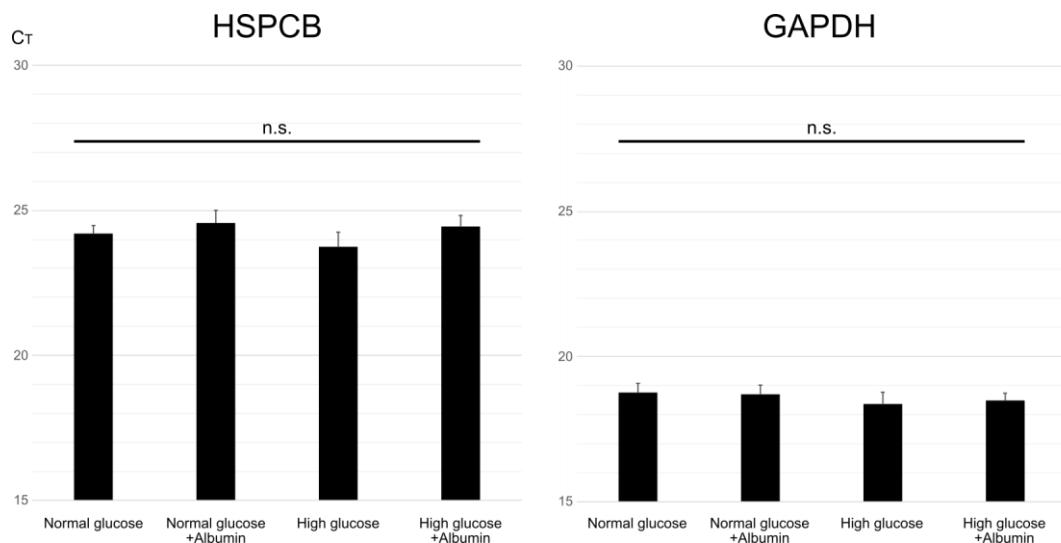
Fax: 0761-203-97177

Supplementary Table 1: Primers used in this study for qPCR analyses.

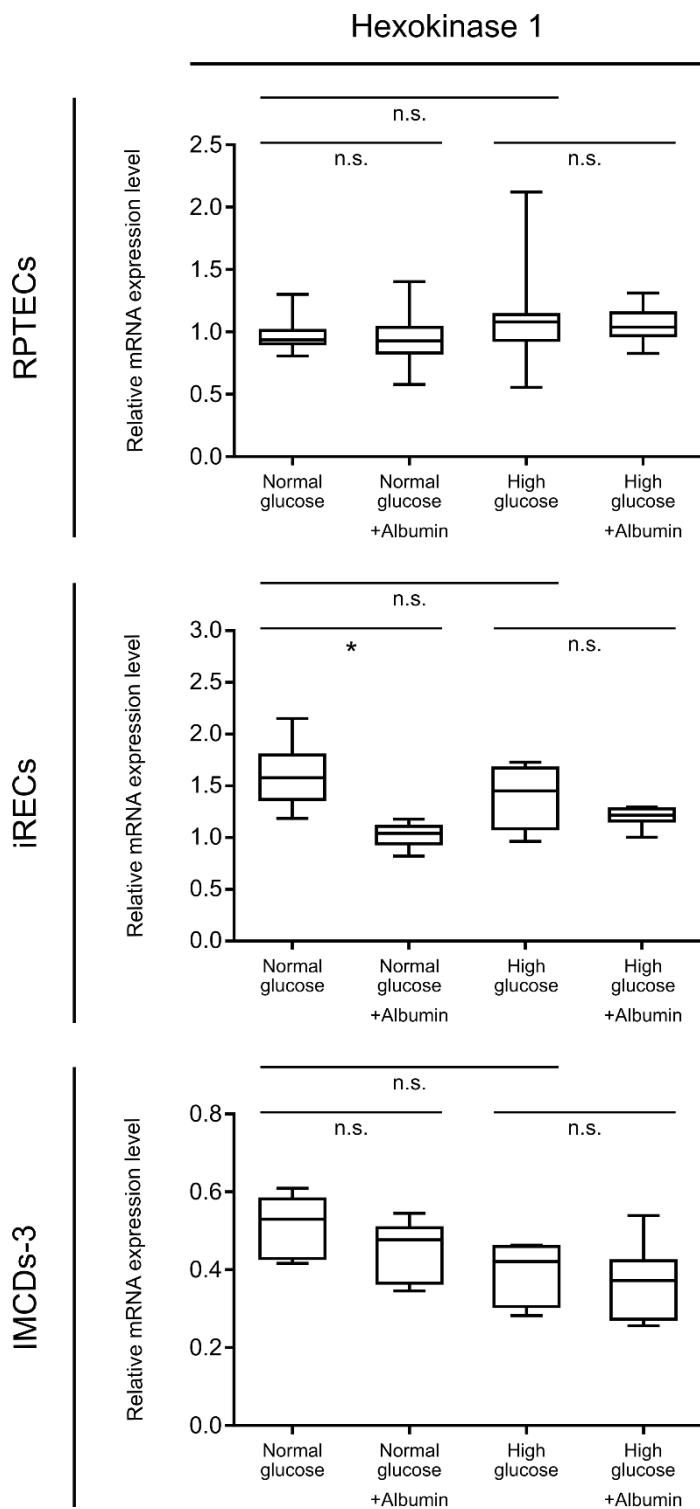
Gene	Accession number		Sequence	Source
mTBP	NM_013684	FP	5'-CCCCTTGACCCCTCACCAAT-3'	
		RP	5'-GAAGCTGCGGTACAATTCCAG-3'	[44]
mHk1	NM_001146100	FP	5'-GAAAGGAGACCAACAGCAGAGC-3'	
		RP	5'-TTCGTTCCCTCGAGATCCAAGG-3'	
mSord	NM_146126	FP	5'-GGATGGTCACTTGCTTGTGGC-3'	
		RP	5'-GGTCTCTTGCCAACCTGGATG-3'	
mAkr1b3	NM_009658	FP	5'-GGTAAAAGGAGCCTCCAGAAG-3'	
		RP	5'-ATCCAGTGGAAATAGTCGGGC-3'	
hHK1	NM_000188.2	FP	5'-CTGCTGGTAAAATCCGTAGTGG-3'	
		RP	5'-GTCCAAGAACGTAGAGATGCAGG-3'	[45]
hAKR1B1	NM_001628	FP	5'-CCAACCTCAACCACATCTCCAGGTG-3'	
		RP	5'-GTCACCACGATGCCTTGACT-3'	
hSORD	NM_003104	FP	5'-GCCGATACAATCTGTCACCTTCC-3'	
		RP	5'-CGCCTCCTCAAAGGTGACATTG-3'	
hGAPDH	NM_001256799.2	FP	5'-GTCTCCTCTGACTTCAACAGCG-3'	
		RP	5'-ACCACCCCTGTTGCTGTAGCCAA-3'	[46]
hHSPCB	NM_001271969	FP	5'-CTCTGTCAGAGTATGTTCTCGC-3'	
		RP	5'-GTTTCCGCACTCGCTCCACAAA-3'	



Supplementary Figure 1: Principal component analysis of RPTECs. PC2 reflected the biological variance and was not affected by glucose or albumin concentration.



Supplementary Figure 2: Ct values of housekeeping genes used for quantitative RT-PCR in RPTECs. Expression levels of both housekeeping genes used for this study are not dependent on glucose or albumin concentrations. Columns display mean Ct- values of three replicates with 95% confidence interval (error bars). Ct – cycle threshold; HSPCB – heat shock 90kDa protein 1, beta; GAPDH – glyceraldehyde-3-phosphate dehydrogenase; n.s. – not significant.



Supplementary Figure 3 : Relative mRNA expression level of hexokinase 1 as detected by qPCR analysis. Comparison between normal and high glucose, high glucose w/wo albumin overload and normal glucose w/wo albumin overload. Data are shown for RPTECs (upper line), iRECs (intermediate line) and IMCD-3 (lower line). Boxplots display mean values of three replicates with 95% confidence interval (whiskers). p-values: n.s. not significant, * <0.05.

References

44. Ferreira MJ, McKenna LB, Zhang J, Reichert M, Bakir B, Buza EL et al. Spontaneous Pancreatitis Caused by Tissue-Specific Gene Ablation of Hhex in Mice. *Cell Mol Gastroenterol Hepatol* 2015;1(5):550–69.
45. Hariton F, Xue M, Rabbani N, Fowler M, Thornalley PJ. Sulforaphane Delays Fibroblast Senescence by Curbing Cellular Glucose Uptake, Increased Glycolysis, and Oxidative Damage. *Oxid Med Cell Longev* 2018;2018:5642148.
46. Izumi-Nakaseko H, Kanda Y, Nakamura Y, Hagiwara-Nagasawa M, Wada T, Ando K et al. Development of correction formula for field potential duration of human induced pluripotent stem cell-derived cardiomyocytes sheets. *Journal of Pharmacological Sciences* 2017;135(1):44–50.