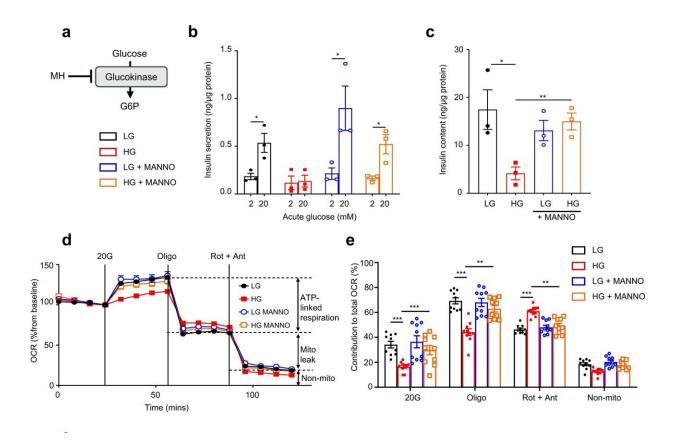


Study uncovers key cause of type 2 diabetes

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Inhibition of glucokinase prevents the effects of chronic hyperglycaemia. **a** Schematic showing how mannoheptulose (MH) inhibits glucose metabolism. **b**, **c** Insulin secretion (**b**) and insulin content (**c**) in LG-cells and HG-cells cultured for 48 h \pm 10 mM mannoheptulose (MANNO) and then stimulated with 2 mM or 20 mM glucose. Mannoheptulose was omitted during the assay (n=3 biologically independent experiments). **d** Oxygen-consumption rate (OCR) in LG-cells and HG-cells cultured for 48 h \pm 10 mM MANNO. OCR was recorded at 2 mM glucose and after sequential addition of 20 mM glucose (20 G), 1 μ M oligomycin (Oligo) and 0.5 μ M rotenone + 0.5 μ M antimycin A (Rot + Ant). Data are expressed as the percentage change from baseline (2 mM glucose); n=10



biologically independent experiments per group. **e** Percentage change in OCR when glucose was raised from 2 to 20 mM (20 G), ATP-linked OCR (Oligo), OCR required to maintain the mitochondrial leak (Rot + Ant) and non-mitochondrial OCR (non-mito); n = 10 biologically independent experiments per group. Same data as in (**d**). **f**, **g** mRNA levels of the indicated genes involved in glycolytic (**f**) and mitochondrial (**g**) metabolism as assessed by qPCR in LG-cells and HG-cells cultured for 48 h \pm 10 mM mannoheptulose (Pdk1, Idh2 and Ndufs8, n = 6 biologically independent experiments; Pfkl, Pfkfb3, Eno1, Sdha and Mdh2, n = 3 biologically independent experiments; Aldob, n = 6 biologically independent experiments for LG and HG but n = 5 for LG + MANNO and HG + MANNO; Ndufs8, n = 6 biologically independent experiments for LG but n = 3 for LG + MANNO, HG and HG + MANNO). All panels show individual data points and mean \pm s.e.m. *P

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