File Name: Supplementary Information Description: Supplementary Figures



Supplementary Figure 1. Obese mice bearing AsPC1-CFP tumours display features of the metabolic syndrome. (a) Table describing the composition of the control chow and high fat diet (HFD) fed to mice in the current study. (b) Glucose tolerance test (GTT) performed on mice (n = 15 lean and n = 14 obese) from Figure 1b, 7 weeks post-HFD feeding (left) and quantification of the area under the curve (AUC, right). (c-e) Plasma glucose (c) insulin (d) and total IGF-1 (e) levels in mice from Figure 1b, at time of euthanasia (n = 15). Data represent the mean \pm s.e.m. in b, and mean \pm s.d. in c-e. $*P \le 0.05$, $***P \le 0.001$ by *t*-test.



Supplementary Figure 2. Human PDA tumours display mild stromal infiltration in both lean and obese mice. Haematoxylin and Eosin stain (H&E, left) and immunohistochemistry of DBA-Lectin, α -Smooth Muscle Actin (α -SMA) and α -Amylase, which stain respectively, ductal cells (normal and neoplastic), stromal cells, and acinar cells, in representative human AsCP1-CFP PDA tumours described in **Figure 1c**. Dashed lines demarcate areas of ductal adenocarcinoma; T, tumour; N, normal acinar tissue; S, stroma; BV, blood vessel. Framed top corner insets represent a 5-fold magnification of the area indicated by a square in the tumour (left), or a 5-fold magnification of stain-positivity for either α -SMA or α -Amylase (right). Scale bars: 50 µm.



Supplementary Figure 3. ARG2 knockdown does not suppress *in vitro* PDA cell growth. (a) Immunoblots of ARG2 in HPAC, PANC-1 and PA-TU-8988T PDA cells showing decreased levels upon ARG2 knockdown (hairpins A06, A07, A10, B09 and B10), compared to control knockdown (shScr or shGFP). (b) Proliferation curves of cells in a demonstrating that ARG2 silencing does not significantly decrease the growth of cultured PDA cells (n = 6). Data are representative of 2 independent experiments and represent the mean \pm s.e.m. $*P \le 0.05$, $**P \le 0.01$, $***P \le 0.001$, $***P \le 0.001$ for differences between control cells and cells with ARG2 knockdown at the indicated timepoints as assessed by two-way ANOVA followed by Tukey test.



Supplementary Figure 4. Nitrogen metabolism pathways are enriched upon *ARG2* knockdown in human PDA tumours. (a) Growth curves for 4-6 week old male *Rag1*^{-/-} mice described in Figure 3a that were fed either chow or HFD (n = 7 for shScr and n = 8 for shARG2) for 14 weeks. Arrow indicates orthotopic injection of control (AsPC1-shScr) or *ARG2* knockdown (AsPC1-shARG2-A10) cells into the pancreata of lean or obese mice. (b) Glucose tolerance test (GTT) performed on mice (n = 7-8) in **a**, 7 weeks post-HFD feeding (left) and quantification of the area under the curve in (AUC, right). (c) List of top metabolic pathways that are differentially altered upon *ARG2* knockdown in AsPC-1 PDA tumours described in Figure 3a. Metabolite data were processed by Metaboanalyst 3.0 and the pathways were ranked by $-\log(2)$ of the *P*-value. In color are pathways involved in nitrogen and arginine metabolism that are enriched in urea cycle metabolites. In **a**,**b**, data represent the mean \pm s.e.m. * $P \le 0.05$, *** $P \le 0.001$, **** $P \le 0.0001$, by two-way ANOVA in **a**, and one-way ANOVA in **b**, followed by Tukey test.



Supplementary Figure 5. *Arg2* loss suppresses the growth of KPC tumours in obese mice. (a) Immunoblots for Arg2 and Arg1 from tumours described in Figure 5c, showing that aberrant, inactive Arg2 is detected by western blotting in mouse KPC; $Arg2^{-/-}$ tumours. (b) Volumes of orthotopic PDA tumours derived from mouse KPC cells with or without *Arg2* loss, that were injected into the pancreata of female lean or obese 16-18 week old C57BL/6J-129 svJae mice (500,000 cells per mouse) and grown for 3 weeks; n = 6 from 2 independent experiments. Data represent mean \pm s.d. ***P* \leq 0.01 by one-way ANOVA followed by Tukey test.



Supplementary Figure 6. ARG2 expression is not induced *in vitro* by insulin, IGF-1, glucose or lipids. (a) Immunoblots of ARG2, pT308-AKT and total AKT in AsPC1, HPAC and SUIT-2 human PDA cells grown in 0.1% serum and treated with the indicated levels of insulin, IGF-1 or lipid mixture for 24 hours. (b) Immunoblots of ARG2 in cells described in **a** that were grown in 0.1% (top) or 10% (bottom) dialyzed serum and treated with the indicated levels of glucose for 24 hours. Data in **a**,**b** are representative of 2 independent experiments.



Supplementary Figure 7. Differential AKT activation and sensitivity to growth factors in human PDA cells. (a) Immunoblots for pS473-AKT, total AKT and PTEN in human PDA cells grown in 10% or 0.1% serum, indicating constitutive AKT activation in "serum-resistant" compared to "serum-sensitive" cells. (b) Proliferation curves of PDA cells described in a showing differential sensitivity to insulin or IGF-1, of "serum-sensitive" cells, compared to growth factor-independent "serum-resistant" cells. All cells were cultured in the presence of 0.1% serum and increasing concentrations of insulin or IGF-1 (n = 6). Data are representative of 2 independent experiments. (c) Proliferation curves of AsPC1-AKT and AsPC1-CFP cells cultured as in b, showing differential sensitivity of AsPC1-CFP, but not AsPC1-AKT cells to insulin or IGF-1 (n = 6). For all graphs, data represent the mean \pm s.e.m. * $P \le 0.01$, **** $P \le 0.0001$ for differences between 0 and 1000 ng/ml⁻¹ insulin, or between 0 and 100 ng ml⁻¹ IGF-1 at the indicated timepoints. Two-way ANOVA was conducted followed by Tukey test.



Supplementary Figure 8. Activated AKT leads to transcriptional induction of nitrogen and arginine metabolic pathways in PDA tumours independent of obesity. (a) Growth curves of 4-6 week old male Rag1^{-/-} mice described in Figure 6b that were fed either chow (Lean-AKT, n =14) or HFD (Obese-AKT, n =15). Arrow indicates orthotopic injection of AsPC1-CFP or AsPC1-AKT cells (10⁵ per mouse) into the pancreata of lean or obese mice. (b) Glucose tolerance test (GTT) performed on mice (n = 14 for Lean-AKT and n = 15 for Obese-AKT) described in **a**, 7 weeks post-HFD feeding (left) and quantification of the area under the curve (AUC, right). c-e, Plasma glucose (c), insulin (d) and total IGF-1 (e) levels in Lean-AKT and Obese-AKT mice, at time of euthanasia (n = 14 for lean and n = 15 for obese). (f) Heatmaps (left) of the induced genes in indicated metabolic pathways with corresponding enrichment plots (right), illustrating differential gene upregulation in AsPC1-CFP tumours grown in obese mice (Obese-CFP, 10-18) and all AsPC1-AKT tumours (Lean-AKT, 19-27; Obese-AKT: 28-36), compared to AsPC1-CFP tumours grown in lean mice (Lean-CFP, 1-9). Red signal indicates higher expression, and blue indicates lower expression relative to the mean expression level within each group. NES, normalized enrichment scores; FDR false discovery rate; Nom., nominal. In \mathbf{a}, \mathbf{b} , data represent the mean \pm s.e.m. In $\mathbf{c}-\mathbf{e}$, data represent the mean \pm s.d. ** $P \leq 0.001$, *** $P \leq 0.001$, **** $P \leq 0.0001$, by two-way ANOVA followed by Tukey test in **a**, and by *t*-test in **b**-**e**.



Supplementary Figure 9. AKT phosphorylation is not increased in PDA tumours of the obese. (a-d) Immunoblots for pT308-AKT, pS473-AKT and total AKT in AsPC1-CFP and AsPC1-AKT PDA tumours grown in lean or obese mice described in **Figures 1c** and **6b**.



Supplementary Figure 10. ARG2 is induced in obesity-associated and AKT-driven PDA *in vivo* but not *in vitro*. (a-c) Immunoblots of ARG2 and ARG1 in orthotopic AsPC1-CFP and AsPC1-AKT tumours described in Figures 1c and 6b, showing increased ARG2 levels in all AKT-driven tumours, independent of obesity. (d) Immunoblots of ARG2 and ARG1 in cultured AsPC1-CFP and AsPC1-AKT cells, showing lack of induction of ARG2 expression by activated AKT *in vitro*.

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Supplementary Figure 11. Scanned films of uncropped Western blots related to the indicated panels from Main Figures 1, 2, 3 and 6. Blots for different antibodies that were developed on the same film (i.e. Figure 2a) or blots including unrelated samples (i.e. Figure 6a) are separated by a black line.