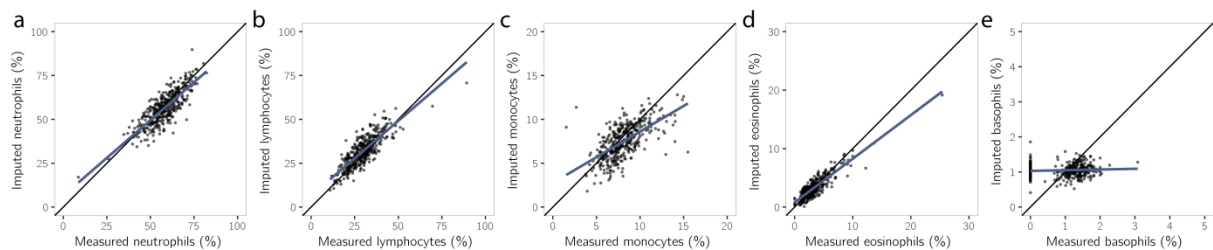


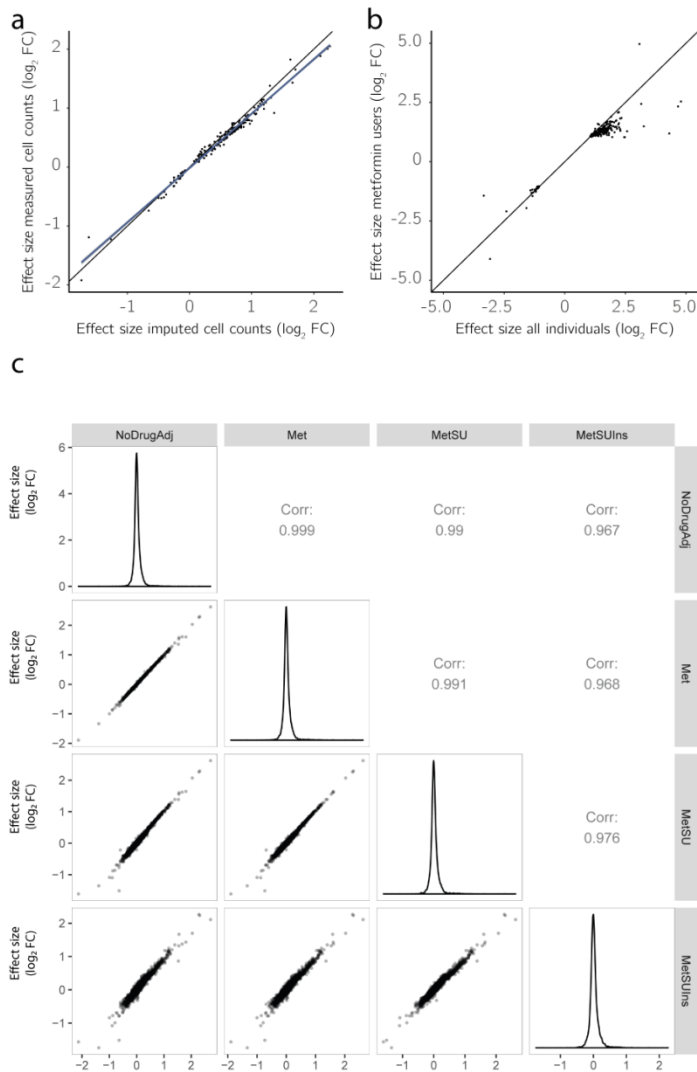
Electronic supplemental material

ESM Table 1, Characteristics of overall cohort, those within the biobank and the subsets for RNA and RNA-seq.

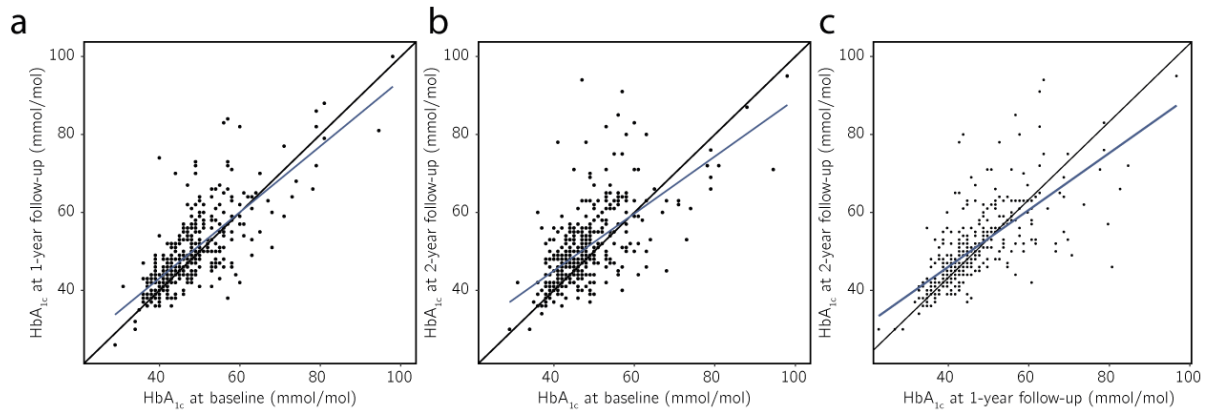
	Entire population (2013)		Population in biobank (2013)		RNA population		RNA-seq population	
	(n=8063)		(n=4921)		(n=1033)		(n=391)	
	Median	IQR	Median	IQR	Median ¹	IQR	Median ¹	IQR
Age (years)	67.5	[60.0-75.3]	67.1	[59.9-74.2]	65.0	[57.7-72.4]	64.0	[57.3-70.0]
Sex (% female)	47.3		44.5		43.4		41.2	
Diabetes duration (years)	6.9	[3.2-12.0]	7.6	[4.1-12.5]	4.5	[2.5-8.7]	3.7	[2.1-5.5]
Glucose (mmol/l)	7.8	[6.8-9.0]	7.8	[6.9-9.1]	7.8	[7.0-8.9]	7.9	[7.2-9.1]
HbA _{1c} (%)	6.5	[6.1-7.1]	6.5	[6.1-7.1]	6.4	[6.0-7.0]	6.4	[6.0-7.0]
HbA _{1c} (mmol/mol)	47	[43-54]	47	[43-54]	47	42-53]	47	[42-53]
BMI (kg/m ²)	29.2	[26.3-33.0]	29.2	[26.5-32.9]	29.5	[26.5-33.0]	29.5	[26.4-33.0]
LDL (mmol/l)	2.3	[1.8-3.0]	2.3	[1.8-2.9]	2.3	[1.8-3.0]	2.3	[1.8-2.9]
HDL (mmol/l)	1.2	[1.0-1.5]	1.2	[1.0-1.5]	1.2	[1.0-1.5]	1.2	[1.0-1.5]
Triacylglycerol (mmol/l)	1.5	[1.1-2.1]	1.5	[1.1-2.1]	1.5	[1.1-2.1]	1.6	[1.1-2.2]
Systolic BP (mmHg)	139	[126-155]	138	[126-154]	134	[124-152]	134	[124-152]
Diastolic BP (mmHg)	78	[73-84]	78	[73-84]	78	[74-84]	80	[75-85]
eGFR (ml/min)	81.0	[67.2-95.2]	81.3	[67.8-95.2]	83.0	[70.5-96.4]	85.8	[73.4-98.5]
Metformin use (%)	68.5		70.5		71.2		89.5	
SU use (%)	29.9		31.9		29.9		11.8	
Insulin use (%)	23.3		24.5		17.1		12.0	



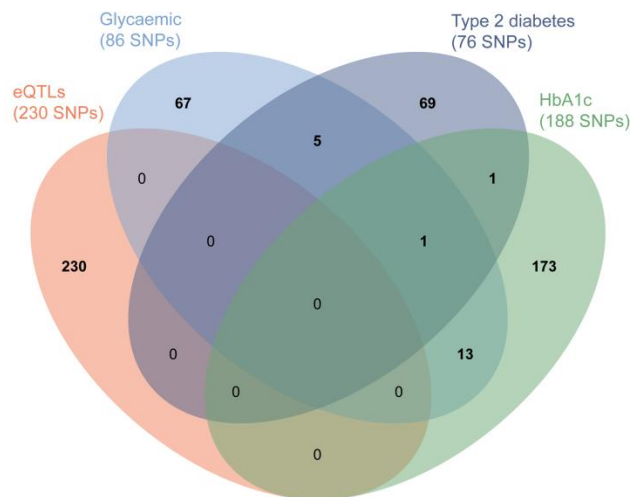
ESM Fig. 1. Measured blood cell percentages against imputed blood cell percentages. **a** Neutrophils, $r = 0.81$. **b** lymphocytes, $r = 0.83$. **c** monocytes, $r = 0.64$. **d** eosinophils, $r = 0.91$. **e** basophils, $r = 0.08$.



ESM Fig. 2. **a** Comparison of effect size (log₂ fold change) with imputed cell counts and measured cell counts. Pearson's correlation between two models: $r = 0.99$. **b** Comparison of effect size of the model on all individuals and a stratified analysis on metformin users only. Pearson's correlation between two models: $r = 0.94$. **c** Comparison of effect size of models 1) without adjustment for medication usage and with for 1) metformin, 2) metformin and SU and 3) metformin, SU and insulin. FC, fold change; Corr, correlation; NoDrugAdj, no adjustment for drug use; Met, adjustment for metformin only; MetSU, adjustment for metformin and sulfonylureas; MetSUIns, adjustment for metformin, sulfonylureas and insulin.



ESM Fig. 3. Correlation of HbA1c between successive years. **a** Baseline HbA_{1c} vs. 1-year follow-up HbA_{1c}, $n = 370$, $r = 0.76$, $p < 0.0001$. **b** Baseline HbA_{1c} vs. 2-year follow-up HbA_{1c}, $n = 360$, $r = 0.64$, $p < 0.0001$. **c** 1-year follow-up HbA_{1c} vs. 2-year follow-up HbA_{1c}, $n = 354$, $r = 0.74$, $p < 0.0001$.



ESM Fig. 4. Overlap between 239 eQTL-SNPs and known diabetes-related traits.