

SUPPLEMENTARY INFORMATION

A distribution-centered approach for analyzing human adipocyte size estimates and their association with obesity-related traits and mitochondrial function.

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Table S1: Study participants' characteristics according to BMI categories.

Variable	All Samples n = 188	Normal weight n = 23	Overweight n = 18	Obesity class I n = 8	Obesity class II n = 13	Obesity class III n = 122
Sex						
Female	129 (69%)	8 (35%)	6 (33%)	6 (75%)	10 (77%)	96 (79%)
Male	59 (31%)	15 (65%)	12 (67%)	2 (25%)	3 (23%)	26 (21%)
Age [years]	48 ± 13 18 - 78	51 ± 17 22 - 78	61 ± 11 32 - 77	53 ± 8 42 - 64	47 ± 13 28 - 74	46 ± 11. 18 - 73
BMI [kg/m²]	43.6 ± 13.3 18.2 - 83.3	22.6 ± 2.1 18.2 - 24.9	27.1 ± 1.2 25.0 - 29.9	32.6 ± 1.5 30.5 - 34.7	37.4 ± 1.6 35.2 - 39.3	51.4 ± 8.3 40.2 - 83.3
Unknown	4	0	0	0	0	0
T2D						
N	140 (76%)	22 (96%)	18 (100%)	8 (100%)	9 (69%)	81 (67%)
Y	45 (24%)	1 (4.3%)	0 (0%)	0 (0%)	4 (31%)	40 (33%)
Unknown	3	0	0	0	0	1
Glucose [mmol/l]	6.0 ± 3.0 2.2 - 22.1	5.0 ± 0.6 3.9 - 6.7	5.1 ± 1.4 2.2 - 7.7	4.8 ± 0.8 3.7 - 5.8	5.5 ± 3.1 2.9 - 13.0	6.4 ± 3.4 2.9 - 22.1
Unknown	49	7	5	3	4	30
HbA1c [%]	6.0 ± 1.2 4.6 - 11.5	NA	5.4 5.4 - 5.4	5.2 ± 0.6 4.8 - 6.2	5.8 ± 1.5 4.6 - 9.3	6.1 ± 1.2 4.8 - 11.5
Unknown	89	23	17	4	5	40
Cholesterol [mmol/l]	5.1 ± 1.0 1.4 - 8.0	5.6 5.6 - 5.6	4.6 ± 1.0 3.9 - 5.4	5.6 ± 0.6 4.9 - 6.2	5.4 ± 1.4 3.8 - 8.0	5.1 ± 1.0 1.4 - 7.4
Unknown	91	22	16	4	5	44
LDL [mmol/l]	3.1 ± 0.8 1.3 - 5.3	3.5 3.5 3.5	2.6 ± 1.1 1.8 - 3.4	3.2 ± 0.6 2.6 - 3.9	2.8 ± 0.8 1.8 - 3.9	3.1 ± 0.9 1.3 - 5.3
Unknown	96	22	16	4	6	48
HDL [mmol/l]	1.3 ± 0.4 0.6 - 2.4	1.5 1.5 1.5	1.4 ± 0.4 1.1 - 1.6	1.9 ± 0.6 1.2 - 2.4	1.3 ± 0.3 0.9 - 1.9	1.3 ± 0.3 0.6 - 2.3
Unknown	94	22	16	4	6	46
Triglycerides [mmol/l]	1.9 ± 1.0 0.6 - 6.8	1.6 1.6 - 1.6	1.8 ± 1.5 0.7 - 2.9	1.5 ± 1.0 0.8 - 3.1	2.7 ± 1.8 1.1 - 5.7	1.8 ± 0.9 0.6 - 6.8
Unknown	92	22	16	4	5	45

Continuous variables are specified as mean ± SD with the second row specifying minimal and maximal values. Categorical variables are given as count with percentages specified in brackets

Table S2: Pearson correlations between mean adipocyte size and anthropometry/laboratory chemistry. Associations with a significance level of $p < 0.05$ are written in bold. Associations that remained significant after Bonferroni correction for multiple testing are written in bold and cursive.

Dep.		BMI [kg/m ²] $n_{sc} = 158$ $n_{vc} = 184$		Age [years] $n_{sc} = 161$ $n_{vc} = 188$		Glucose [mmol/l] $n_{sc} = 120$ $n_{vc} = 139$		HbA _{1c} [%] $n_{sc} = 90$ $n_{vc} = 99$		Cholesterol [mmol/l] $n_{sc} = 89$ $n_{vc} = 97$		LDL [mmol/l] $n_{sc} = 83$ $n_{vc} = 92$		HDL [mmol/l] $n_{sc} = 86$ $n_{vc} = 94$		Triglycerides [mmol/l] $n_{sc} = 88$ $n_{vc} = 96$	
		$r_{pearson}$	p	$r_{pearson}$	p	$r_{pearson}$	p	$r_{pearson}$	p	$r_{pearson}$	p	$r_{pearson}$	p	$r_{pearson}$	p	$r_{pearson}$	p
Mean area [μm ²]	sc	0.49	5E-11	-0.19	0.015	0.12	0.193	0.07	0.529	-0.20	0.055	-0.11	0.313	-0.01	9.2E-01	0.11	0.328
	vc	0.47	2E-11	-0.02	0.762	0.26	0.002	0.32	0.001	-0.10	0.345	0.04	0.706	-0.34	7.4E-04	0.20	0.050
Mean diameter [μm]	sc	0.49	6E-11	-0.18	0.024	0.11	0.232	0.06	0.575	-0.22	0.041	-0.12	0.281	0.00	1.000	0.09	0.420
	vc	0.48	9E-12	-0.01	0.921	0.26	0.002	0.31	0.002	-0.06	0.542	0.03	0.783	-0.32	1.6E-03	0.19	0.063
Mean volume [pL]	sc	0.48	1E-10	-0.20	0.013	0.12	0.181	0.07	0.517	-0.19	0.072	-0.10	0.357	-0.02	0.872	0.12	0.280
	vc	0.46	6E-11	-0.03	0.677	0.27	0.002	0.31	0.002	-0.13	0.221	0.03	0.783	-0.35	5.4E-04	0.20	0.051
Mean surf. area [μm ²]	sc	0.49	5E-11	-0.19	0.015	0.12	0.193	0.07	0.529	-0.20	0.055	-0.11	0.313	-0.01	0.917	0.11	0.328
	vc	0.47	2E-11	-0.02	0.762	0.26	0.002	0.32	0.001	-0.10	0.345	0.02	0.831	-0.34	7.4E-04	0.20	0.050

Table S3: Multiple linear regression analysis on mean adipocyte diameter and anthropometry/laboratory values. Mean adipocyte diameter from histology was set as the dependent variable. To exclude an influence of the BMI on the correlation between mean adipocyte size and the independent variable BMI was used as a second independent variable to adjust for in multiple linear regression. p-values < 0.05 are written in bold.

Variable	Depot (n)	β adj. for BMI	p adj. for BMI	Adj. R ² Model
Age	sc (n = 158)	-0.02	0.644	0.23
	vc (n = 184)	0.01	0.028	0.24
Sex (male)	sc (n = 158)	-0.98	0.426	0.23
	vc (n = 184)	3.34	0.006	0.25
T2D	sc (n = 158)	0.56	0.666	0.23
	vc (n = 183)	3.34	0.010	0.25
Glucose [mmol/l]	sc (n = 117)	0.16	0.443	0.22
	vc (n = 135)	0.56	0.004	0.28
HbA1C [%]	sc (n = 87)	0.18	0.775	0.06
	vc (n = 95)	1.59	0.002	0.15
Cholesterol [mmol/l]	sc (n = 86)	- 1.22	0.084	0.11
	vc (n = 93)	- 0.19	0.769	0.10
LDL [mmol/l]	sc (n = 80)	- 0.82	0.353	0.09
	vc (n = 88)	0.17	0.834	0.12
HDL [mmol/l]	sc (n = 83)	1.60	0.446	0.08
	vc (n = 90)	-4.55	0.015	0.14
Triglycerides [mmol/l]	sc (n = 85)	0.60	0.392	0.09
	vc (n = 92)	1.58	0.017	0.15

Table S4: Correlations between mean adipocyte diameter and mitochondrial respiratory capacity of mature adipocytes. Dependent on the distribution of the data either pearson or spearman correlations were used to assess the relationship between mean adipocyte diameter and respiratory chain function. Significant correlations are written in bold.

Respiratory state	Depot	Mean Diameter		
		Pearson/Spearman Correlation coefficient	p-value	Method
Free Oxphos capacity (pmol/(s*ng DNA))	sc (24)	- 0.41	0.045	Pearson
	vc (35)	- 0.21	0.209	Pearson
Oxphos capacity (pmol/(s*ng DNA))	sc (24)	- 0.53	0.008	Pearson
	vc (35)	- 0.36	0.035	Spearman
Leak respiration oligomycin (pmol/(s*ng DNA))	sc (24)	- 0.40	0.050	Pearson
	vc (35)	- 0.37	0.026	Pearson
Electron transfer system capacity (pmol/(s*ng DNA))	sc (24)	- 0.59	0.003	Pearson
	vc (35)	- 0.33	0.058	Pearson

Table S5: Multiple linear regression analysis on mean adipocyte diameter and adipocyte respiratory chain function. Mean adipocyte diameter from histology was set as the dependent variable. To exclude an influence of the BMI on the correlation between mean adipocyte size and mitochondrial respiratory capacity BMI was used as a second independent variable to adjust for in multiple linear regression. p-values < 0.05 are written in bold.

Dependent variable	Depot (n)	β adj. for BMI	p adj. for BMI	Adj. R² Model
Free Oxphos capacity (pmol/(s*ng DNA))	sc (24)	-0.05	0.630	0.31
	vc (35)	-0.07	0.273	0.11
Oxphos capacity (pmol/(s*ng DNA))	sc (24)	-0.09	0.254	0.34
	vc (35)	0.17	0.126	0.14
Leak respiration oligomycin (pmol/(s*ng DNA))	sc (24)	-0.16	0.227	0.35
	vc (35)	-0.28	0.053	0.18
Electron transfer system capacity (pmol/(s*ng DNA))	sc (24)	-0.14	0.131	0.37
	vc (35)	-0.11	0.083	0.14

Table S6: Correlations between mean adipocyte diameter and mitochondrial respiratory capacity of mature adipocytes with 200 sampled cells to determine mean adipocyte diameter. Dependent on the distribution of the data either pearson or spearman correlations were used to assess the relationship between mean adipocyte diameter and respiratory chain function. Significant correlations are written in bold.

Respiratory state	Depot	Mean Diameter		
		Pearson/Spearman Correlation coefficient	p-value	Method
Free Oxphos capacity (pmol/(s*ng DNA))	sc (32)	-0.45	0.010	Pearson
	vc (41)	-0.21	0.182	Pearson
Oxphos capacity (pmol/(s*ng DNA))	sc (32)	-0.59	3.83E-04	Pearson
	vc (41)	-0.33	0.033	Spearman
Leak respiration oligomycin (pmol/(s*ng DNA))	sc (32)	-0.50	0.003	Pearson
	vc (41)	-0.32	0.037	Pearson
Electron transfer system capacity (pmol/(s*ng DNA))	sc (32)	-0.67	3.18E-05	Pearson
	vc (41)	-0.34	0.031	Pearson

Table S7: Multiple linear regression analysis on mean adipocyte diameter and adipocyte respiratory chain function with 200 sampled cells to determine mean adipocyte diameter. Mean adipocyte diameter from histology was set as the dependent variable. To exclude an influence of the BMI on the correlation between mean adipocyte size and mitochondrial respiratory capacity BMI was used as a second independent variable to adjust for in multiple linear regression. p-values < 0.05 are written in bold.

Variable	Depot (n)	β adj. for BMI	p adj. for BMI	Adj. R ² Model
Free Oxphos capacity (pmol/(s*ng DNA))	sc (32)	-0.13	0.101	0.25
	vc (41)	-0.08	0.255	0.08
Oxphos capacity (pmol/(s*ng DNA))	sc (32)	-0.16	0.007	0.36
	vc (41)	-0.09	0.104	0.10
Leak respiration oligomycin (pmol/(s*ng DNA))	sc (32)	-0.29	0.011	0.34
	vc (41)	-0.29	0.057	0.13
Electron transfer system capacity (pmol/(s*ng DNA))	sc (32)	-0.221	0.001	0.44
	vc (41)	-0.14	0.045	0.12

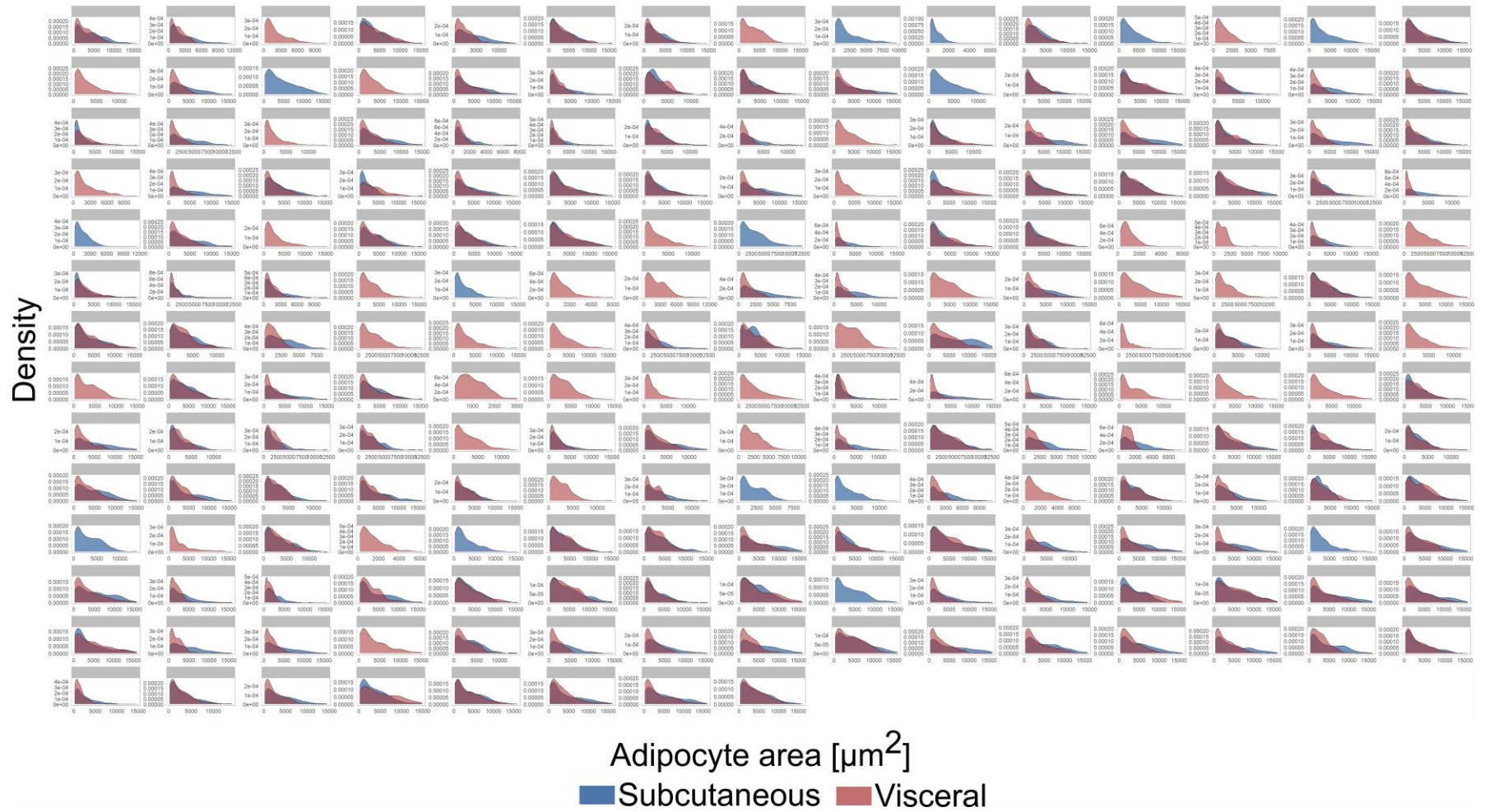


Figure S1: Individual density plots for sc and vc adipocyte area based on 500 sampled cells

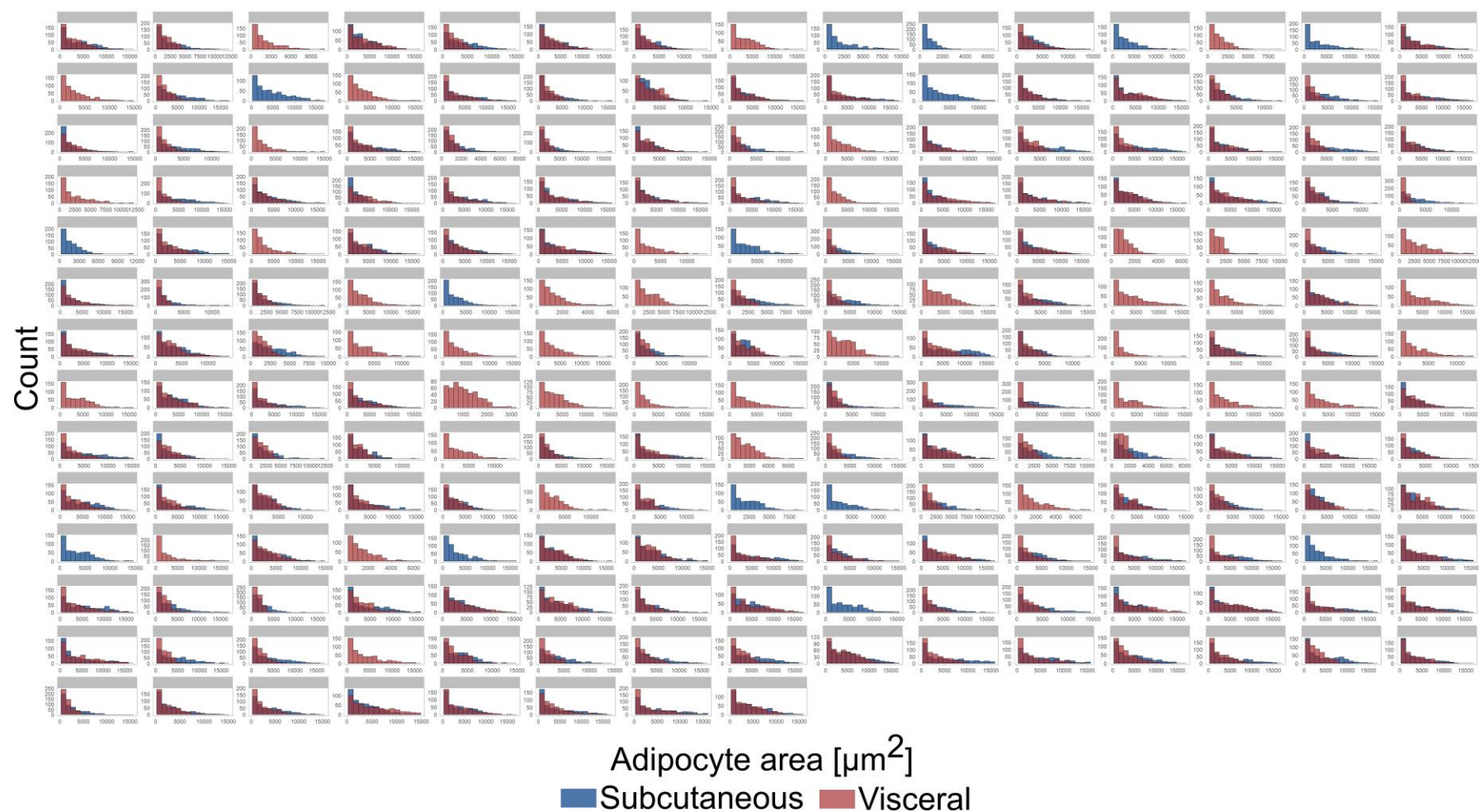


Figure S2: Individual histograms for sc and vc adipocyte area based on 500 sampled cells

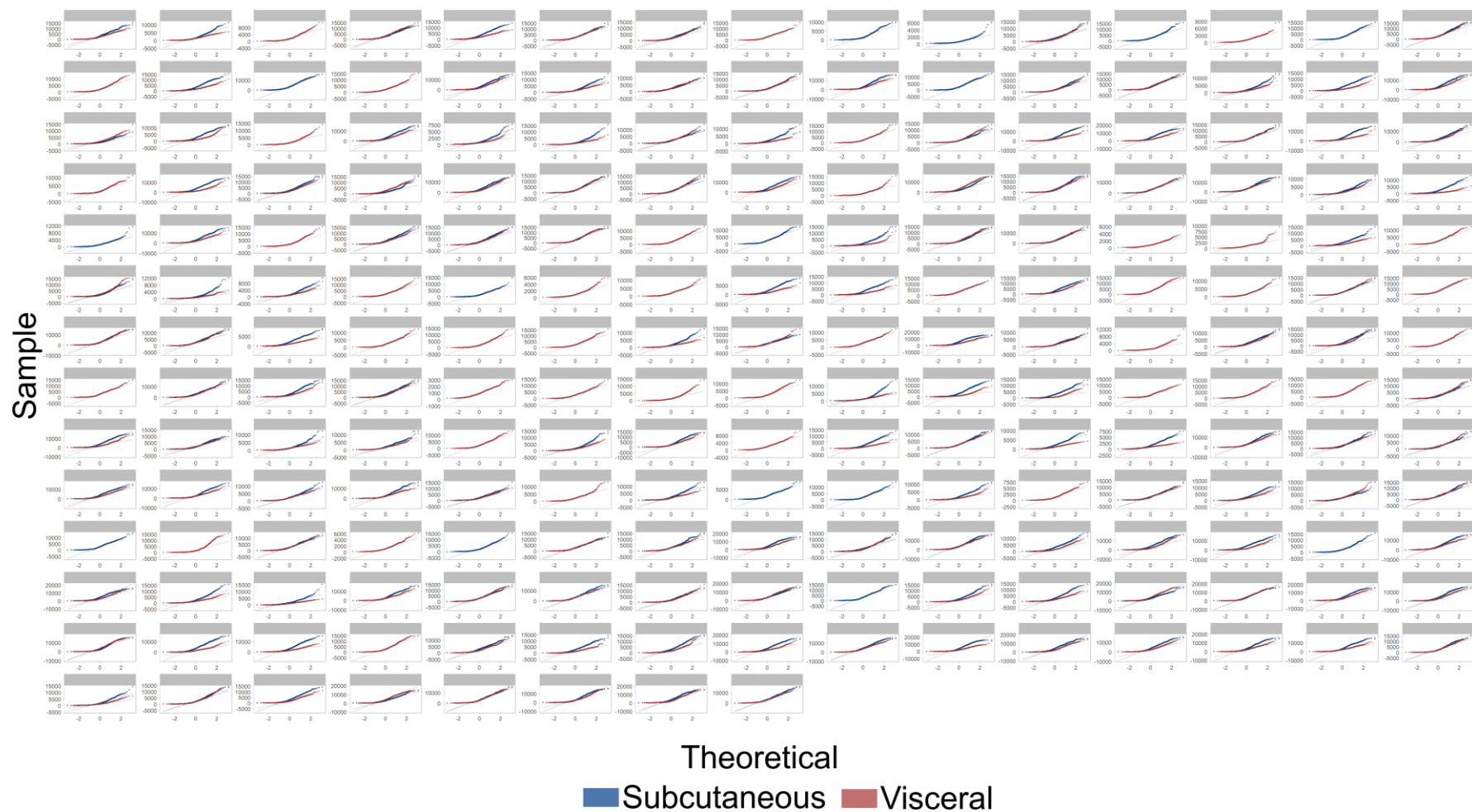


Figure S3: Individual QQ plots for sc and vc adipocyte area based on 500 sampled cells

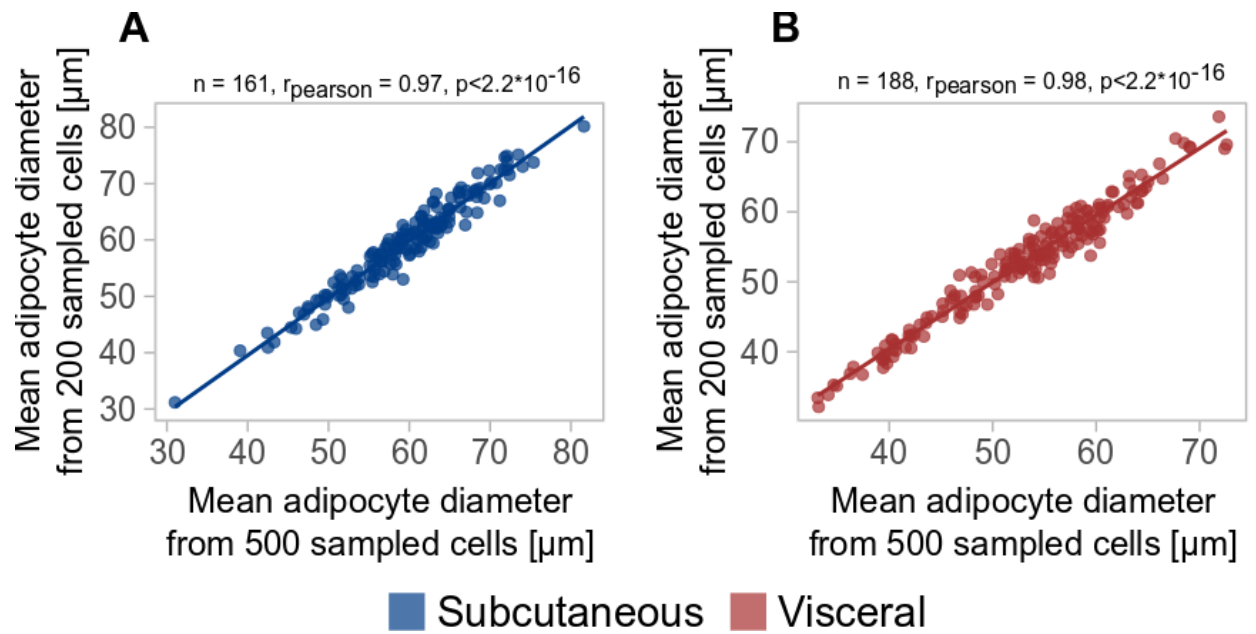
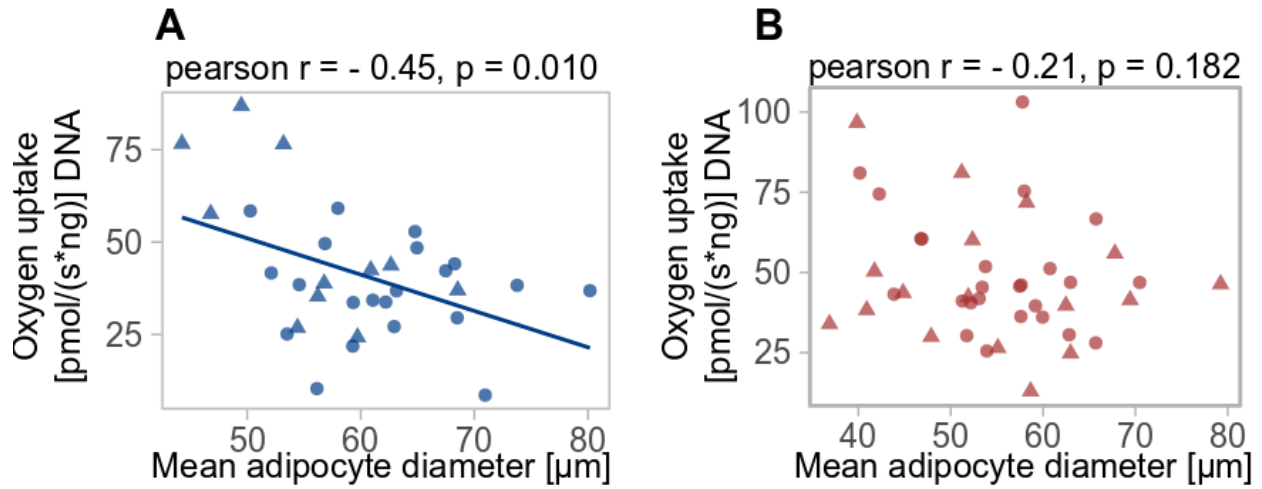


Figure S4: Correlations between sc (A) and vc (B) mean adipocyte diameter from 500 and 200 sampled cells.

Free OXPHOS Capacity



Electron transfer system capacity

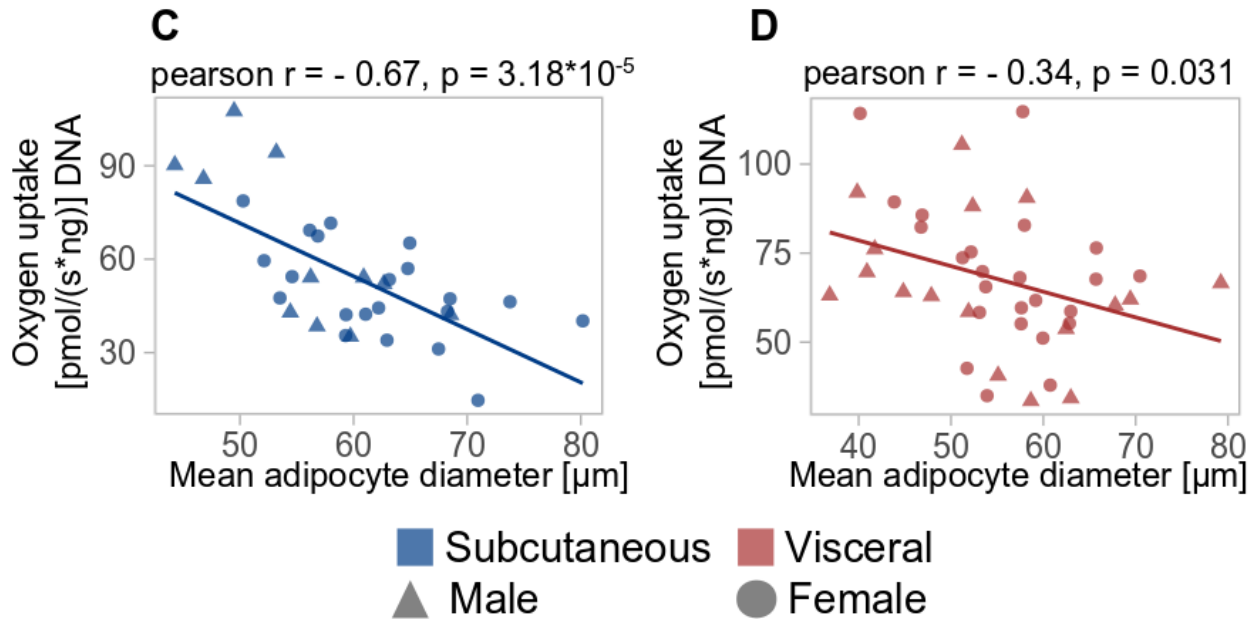


Figure S5: Correlations between mean adipocyte diameter and mitochondrial respiratory capacity of mature adipocytes with 200 sampled cells to determine mean adipocyte diameter.

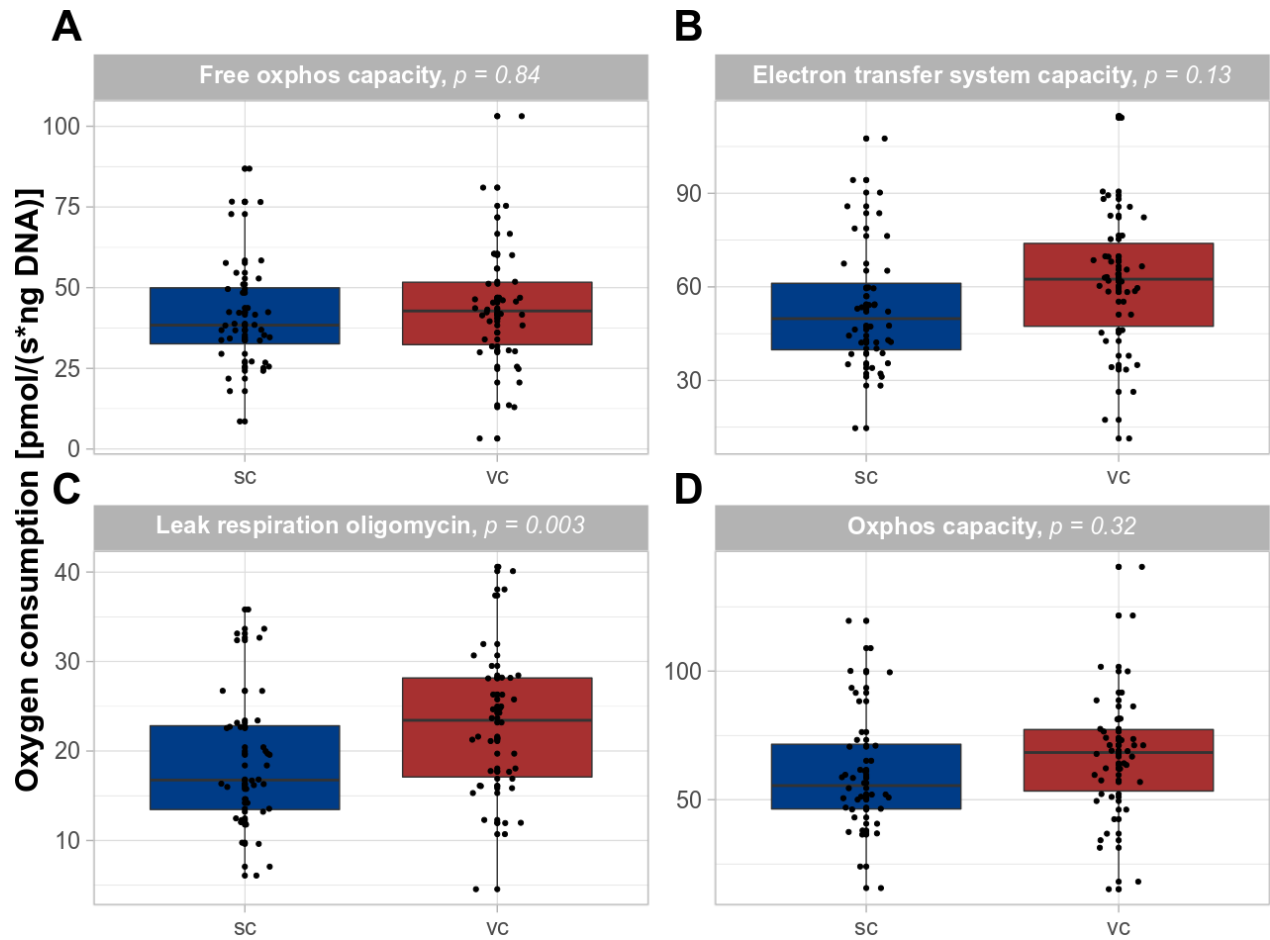


Figure S6: Differences in respiratory states according to adipose tissue depot. Paired sample t-tests were used to test for differences in respiration in 51 samples. No significant differences were found except for leak respiration in the presence of oligomycin ($p = 0.003$). P-values originate from two-sided paired t-tests.