

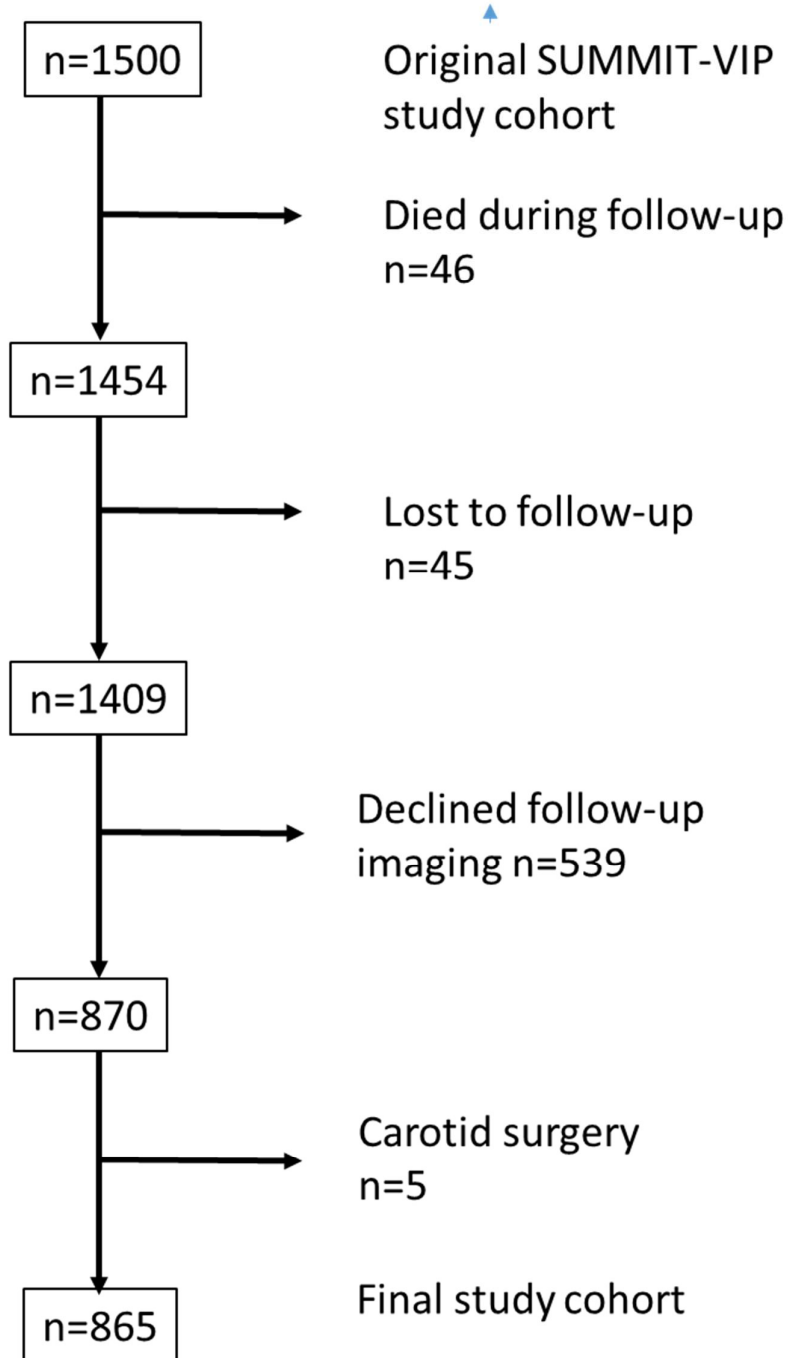
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**Supplemental information**

**Plaque characteristics and biomarkers  
predicting regression and progression  
of carotid atherosclerosis**

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**On-line supplemental data**



**Supplemental figure 1.** Consort flow diagram of study cohort. Related to table 1

**Supplemental table 1.** Baseline clinical characteristics of study subjects with and without T2D. Related to table 1.

	<b>T2D (n=566)</b>	<b>Non-T2D (n=299)</b>	<b>P</b>
Age (years)	68.5±8.0	67.4±9.1	0.07
Gender (% males)	70.0	59.4	0.09
Current smokers (%)	8.7	6.4	0.30
Prevalent CVD (%)	39.2	41.3	0.56
Duration of T2D (years)	10.7±7.8	-	
BMI (kg/m <sup>2</sup> )	30.3±5.0	26.9±4.0	<0.001
<b>Medications</b>			
Insulin (%)	25.4	0	
Statin (%)	74.3	49.8	<0.001
ACE inhibitors (%)	43.6	24.7	<0.001
Metformin (%)	70.0	0	
Betablockers (%)	34.5	26.6	0.02
<b>Metabolic factors</b>			
HbA1c (mmol/mmol)	57.1±13.2	38.9±3.9	<0.001
LDL (mmol/L)	2.22±0.88	2.81±0.90	<0.001
HDL (mmol/L)	1.28±0.39	1.52±0.42	<0.001
Triglycerides (mmol/L)	1.40 (1.00-2.09)	1.11 (0.83-1.54)	<0.001
<b>Blood pressure</b>			
Systolic (mmHg)	136±18	132±17	0.006
Diastolic (mmHg)	76±10	76±9	0.94
<b>Renal function</b>			
eGFR (mL/min <sup>-1</sup> per 1.73m <sup>2</sup> )	79.8±20.3	80.8±17.0	0.47

Variables with normal distribution are shown as mean±standard deviation and skewed variables as median and interquartile range. eGFR; estimated glomerular filtration rate. Differences between means of normally distributed continuous variables were assessed with independent sample *t* tests and between skewed variables with the Mann-Whitney U-test.  $\chi^2$  test was used for categorical variables.

**Supplemental table 2.** Relationships between medications and change in carotid IMT. Related to table 2.

	<b>Mean IMT</b>			
	Regression	No change	Progression	P for trend
<b>CCA</b>	(n=181)	(n=379)	(n=299)	
Statin (%)	69.1	66.4	63.5	ns
ACE inhibitors (%)	36.1	39.0	35.1	ns
Beta-blockers (%)	37.8	30.0	30.9	ns
Antiplatelet (%)	44.8	47.1	45.3	ns
<b>Bulb</b>	(n=192)	(n=206)	(n=412)	
Statin (%)	67.4	60.3	65.0	ns
ACE inhibitors (%)	34.2	33.3	38.1	ns
Beta-blockers (%)	29.6	26.1	33.2	ns
Antiplatelet (%)	45.5	39.2	43.3	ns

P for trend was calculated by Chi-square linear-by-linear association.

**Supplemental table 3.** Relationships between baseline clinical characteristics with change in carotid IMT using  $\pm 0.1$  mm as cut-off for no change in IMT. Related to table 2.

	<b>Mean bulb IMT</b>		
	Regression n=137	No change n=367	Progression n=302
Age (years)	68.2 $\pm$ 8.4	67.0 $\pm$ 8.6	69.5 $\pm$ 8.1***
Male gender (%)	54.0	61.9	69.9*
Current smokers (%)	6.9	7.4	7.5
BMI (kg/m <sup>2</sup> )	29.1 $\pm$ 4.9*	28.1 $\pm$ 4.8	28.8 $\pm$ 4.5
<b>Metabolic factors</b>			
HbA1c (mmol/mmol)	48.8 (40.8-60.0)	46.0 (40.0-57.0)	48.0 (39.0-59.0)
LDL (mmol/L)	2.23 $\pm$ 0.94**	2.48 $\pm$ 0.92	2.53 $\pm$ 0.92
HDL (mmol/L)	1.34 $\pm$ 0.40	1.40 $\pm$ 0.43	1.36 $\pm$ 0.42
TG (mmol/L)	1.48 (1.04-2.04)	1.20 (0.90-1.67)	1.30 (0.91-1.89)
<b>Blood pressure</b>			
Systolic (mmHg)	138 $\pm$ 17*	132 $\pm$ 17	137 $\pm$ 17**
Diastolic (mmHg)	77 $\pm$ 9	76 $\pm$ 9	76 $\pm$ 10
<b>Renal function</b>			
eGFR	78.5 $\pm$ 20.9	82.2 $\pm$ 17.5	79.0 $\pm$ 20.4
	<b>Mean CCA IMT</b>		
	Regression n=88	No change N=633	Progression n=142
Age (years)	68.2 $\pm$ 8.4	68.1 $\pm$ 8.5	68.1 $\pm$ 8.2
Male gender (%)	59.1	63.8	63.4
Current smokers (%)	9.1	7.8	7.7
BMI (kg/m <sup>2</sup> )	29.9 $\pm$ 5.2	29.1 $\pm$ 5.7	28.7 $\pm$ 4.9
<b>Metabolic factors</b>			
HbA1c (mmol/mmol)	49.9 (39.0-60.2)	48.0 (40.0-58.1)	47.0 (41.0-57.8)
LDL (mmol/L)	2.28 $\pm$ 0.97	2.44 $\pm$ 0.94	2.45 $\pm$ 0.87
HDL (mmol/L)	1.24 $\pm$ 0.44**	1.38 $\pm$ 0.42	1.28 $\pm$ 0.40
TG (mmol/L)	1.50 (1.00-2.10)	1.30 (0.93-1.85)	1.30 (0.90-1.86)
<b>Blood pressure</b>			
Systolic (mmHg)	137 $\pm$ 18	134 $\pm$ 17	137 $\pm$ 16
Diastolic (mmHg)	77 $\pm$ 9*	75 $\pm$ 9	77 $\pm$ 10
<b>Renal function</b>			
eGFR	77.9 $\pm$ 20.7	80.1 $\pm$ 19.0	82.2 $\pm$ 19.3

Variables with normal distribution are shown as mean $\pm$ standard deviation and skewed variables as median and interquartile range. eGFR (mL/min<sup>-1</sup> per 1.73m<sup>2</sup>). Differences between IMT change categories (regression and progression versus no change) were calculated with one-way ANOVA and significance of differences versus no change with Scheffé's post hoc test. Chi-square test was used to analyze difference between categorical variables. \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001.

**Supplemental table 4.** Relationships between biomarkers and change in carotid bulb IMT using  $\pm 0.1$  mm as cut-off for no change in IMT. Related to table 3.

	Regression (n=137)	No change (n=367)	Progression (n=302)
<b>Inflammatory</b>			
hsCRP (mg/L)	1.48 (0.69-3.15)	1.12 (0.60-2.32)	1.46 (0.75-2.90)
IL-6	42.5 (31.3-66.3)**	31.8 (22.2-48.2)	34.5 (23.7-56.2)
MCP-1	14.2 (10.9-18.0)***	11.4 (8.6-14.7)	11.5 (9.0-14.9)
MIP-1 $\alpha$	5.13 (4.22-6.50)**	4.71 (3.97-5.39)	4.59 (3.86-5.66)
<b>Matrix proteases</b>			
MMP-3	2.51 (1.91-3.40)	2.38 (1.90-2.97)	2.38 (1.91-3.03)
MMP-7	468 (282-676)	387 (283-559)	446 (311-690)*
MMP-12	154 (99-232)**	122 (91-169)	136 (98-199)*
<b>Apoptosis</b>			
TNFR-1	6608 (5185-8192)*	6039 (5008-7473)	6295 (5185-7643)
TRAILR-2	3.68 (2.51-4.63)*	3.24 (2.35-4.21)	3.18 (2.31-4.33)
Fas	226 (177-280)*	197 (168-242)	198 (167-244)
<b>SMC Growth factors</b>			
PDGF	70.0 (29.2-166.8)***	132.5 (61.2-246.0)	174.8 (90.4-263.2)*
EGF	35.6 (11.0-73.5)***	51.1 (21.1-116.2)	67.6 (32.4-121.1)*
HBEGF	22.8 (18.9-29.0)**	25.9 (21.0-34.8)	26.7 (21.6-34.5)
<b>EC growth factors</b>			
HGF	121 (93-150)**	101 (83-130)	103 (87-127)
PIGF	191 (144-246)**	163 (139-202)	168 (139-207)
VEGF	1510 (1193-1965)	1370 (1097-1783)	1370 (1097-1739)

All biomarkers except hsCRP are expressed as arbitrary units and values are shown as median and IQR. Differences between IMT change categories (regression and progression versus no change) were calculated with one-way ANOVA and Scheffe's post hoc test. \* $p < 0.05$ , \*\* $p < 0.01$  and \*\*\* $p < 0.001$ . If correcting for multiple comparisons in the present table, the threshold for significance is  $p < 0.004$ . SMC; smooth muscle cell, EC; endothelial cell.

**Supplemental table 5.** Relationships between biomarkers and change in common carotid IMT using  $\pm 0.1$  mm as cut-off for no change in IMT. Related to table 4.

	Regression (n=181)	No change (n=379)	Progression (n=299)
<b>Inflammatory</b>			
hsCRP (mg/L)	1.12 (0.62-1.90)	1.33 (0.66-2.87)	1.20 (0.70-2.53)
IL-6	40.8 (24.1-61.0)	34.9 (23.2-53.1)	36.9 (25.4-61.0)
MCP-1	13.4 (10.5-16.1)	11.5 (8.8-15.5)	12.1 (9.7-15.6)
MIP-1 $\alpha$	4.77 (3.79-6.00)	4.72 (3.95-5.74)	4.94 (4.00-5.99)
<b>Matrix proteases</b>			
MMP-3	2.60 (1.95-3.13)	2.36 (1.89-3.07)	2.62 (1.92-3.34)
MMP-7	440 (220-709)	422 (296-633)	471 (335-704)
MMP-12	143 (107-205)	133 (97-201)	144 (100-207)
<b>Apoptosis</b>			
TNFR-1	6427 (5293-8079)	6339 (5149-7804)	6295 (5104-7968)
TRAILR-2	3.89 (3.05-4.87)**	3.32 (2.37-4.32)	3.11 (2.42-4.19)
Fas	219 (173-267)	204 (169-247)	211 (170-272)
<b>SMC Growth factors</b>			
PDGF	84.4 (43.9-200.2)	144.0 (61.0-243.9)	133.9 (62.6-233.5)
EGF	41.1 (17.1-92.6)	53.8 (20.6-111.4)	51.2 (23.5-104.1)
HBEGF	24.1 (18.4-33.2)	25.3 (20.7-32.7)	26.4 (21.6-33.8)
<b>EC growth factors</b>			
HGF	110 (88-133)	106 (87-133)	114 (87-137)
PIGF	183 (144-218)	171 (141-214)	173 (144-228)
VEGF	1443 (1193-1786)	1339 (1121-1795)	1448 (1105-1761)

All biomarkers except hsCRP are expressed as arbitrary units and values are shown as median and IQR. Differences between IMT change categories (regression and progression versus no change) were calculated with one-way ANOVA and Scheffe's post hoc test. \* $p < 0.05$ , \*\* $p < 0.01$  and \*\*\* $p < 0.001$ . If correcting for multiple comparisons in the present table, the threshold for significance is  $p < 0.004$ . SMC; smooth muscle cell, EC; endothelial cell.



**Supplemental table 6.** Relationships between baseline vascular measurements and change in carotid IMT using  $\pm 0.1$  mm as cut-off for no change in IMT. Related to table 5.

	<b>Carotid bulb</b>		
	Regression	No change	Progression
<b>CCA</b>	n=191	n=200	n=412
Mean IMT (mm)	0.88 (0.81-1.05)	0.86 (0.75-0.99)	0.89 (0.78-1.03)
<b>Bulb</b>			
Mean IMT (mm)	1.31 (1.10-1.69)***	1.02 (0.87-1.19)	1.10 (0.90-1.40)
GSM baseline	40 (28-63)***	66 (48-83)	69 (50-90)
GSM follow-up	50 (35-70)***	67 (49-92)	73 (50-96)
<b>Arterial stiffness</b>			
PWV (m/s)	10.9 (9.3-13.4)**	10.0 (8.6-11.9)	10.5 (8.9-12.5)
<b>Endothelial function</b>			
RHI	2.03 (1.75-2.47)*	2.19 (1.79-2.66)	2.13 (1.79-2.62)
	<b>CCA</b>		
<b>CCA</b>	n=181	n=379	n=299
Mean IMT (mm)	1.03 (0.88-1.24)***	0.87 (0.77-0.99)	0.85 (0.74-0.99)
<b>Bulb</b>			
Mean IMT (mm)	1.10 (0.94-1.51)	1.06 (0.91-1.31)	1.10 (0.93-1.39)
GSM baseline	58 (36-79)	64 (43-83)	67 (46-88)
GSM follow-up	54 (43-71)	66 (46-87)	62 (47-87)
<b>Arterial stiffness</b>			
PWV (m/s)	10.8 (9.6-13.6)	10.2 (8.7-12.4)	10.6 (9.0-12.6)
<b>Endothelial function</b>			
RHI	1.98 (1.74-2.39)	2.14 (1.76-2.64)	2.16 (1.88-2.69)

Grey scale median (GSM) value is the median of all grey levels of the pixels in the plaque. PWV; pulse wave velocity, RHI; reactive hyperemia index. Differences between IMT change categories (regression and progression versus no change) were calculated with one-way ANOVA and Scheffe's post hoc test. Values are shown as median and IQR. \* $p < 0.05$ , \*\* $p < 0.01$  and \*\*\* $p < 0.001$ .

**Supplemental table 7.** Relationships between biomarkers and change in carotid bulb IMT in subjects with T2D. Related to table 3.

	Correlation	P	Regression (n=126)	No change (n=125)	Progression (n=269)
<b>Inflammatory</b>					
hsCRP (mg/L)	0.06	ns	1.42 (0.66-3.30)	1.65 (0.65-2.96)	1.56 (0.73-3.00)
IL-6	0.01	ns	41.4 (29.4-68.1)	33.4 (24.9-52.7)	36.3 (26.0-57.3)
MCP-1	-0.05	ns	13.7 (9.8-17.3)	11.4 (8.6-15.3)	12.0 (9.4-15.6)
MIP-1 $\alpha$	-0.07	ns	4.89 (3.97-6.11)	4.68 (3.92-5.36)	4.66 (3.94-5.74)
<b>Matrix proteases</b>					
MMP-3	0.00	ns	2.51 (1.91-3.40)	2.38 (1.90-2.97)	2.38 (1.91-3.03)
MMP-7	0.12	0.005	452 (233-779)	405(304-564)	495 (341-719)*
MMP-12	0.05	ns	154 (100-223)	128 (95-183)	151 (108-211)
<b>Apoptosis</b>					
TNFR-1	0.04	ns	6517 (5185-8422)	6320 (5113-7486)	6700 (5367-8192)
TRAILR-2	-0.08	ns	3.88 (2.59-5.07)	3.36 (2.57-4.32)	3.36 (2.48-4.53)
Fas	-0.08	ns	231 (176-286)	205 (168-247)	208 (175-251)
<b>SMC Growth factors</b>					
PDGF	0.28	8.8E-11	75.1 (33.3-164.3)**	137.7 (65.3-231.5)	177.3 (90.4-263.7)*
EGF	0.24	3.9E-8	33.8 (12.6-74.5)*	54.0 (25.1-106.0)	68.6 (32.4-123.6)
HBEGF	0.13	0.003	22.5 (18.5-29.2)	26.1 (20.7-33.6)	26.9 (21.6-34.3)
<b>EC growth factors</b>					
HGF	-0.04	ns	123 (92-153)	104 (87-130)	110 (92-134)
PIGF	-0.01	ns	187 (143-246)	161 (142-198)	174 (146-217)
VEGF	-0.03	ns	1531 (1184-2091)	1389 (1142-1827)	1448 (1144-1808)

All biomarkers except hsCRP are expressed as arbitrary units and values are shown as median and IQR. Correlations are shown as Spearman rank correlation coefficients. Differences between IMT change categories (regression and progression versus no change) were calculated with one-way ANOVA and Scheffe's post hoc test. \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001. If correcting for multiple comparisons in the present table, the threshold for significance is p<0.004. SMC; smooth muscle cell, EC; endothelial cell.

**Supplemental table 8.** Relationships between biomarkers and change in carotid bulb IMT in subjects without T2D. Related to table 3.

	Correlation	P	Regression (n=65)	No change (n=75)	Progression (n=143)
<b>Inflammatory</b>					
hsCRP (mg/L)	-0.04	ns	1.38 (0.70-2.55)	0.84 (0.60-2.06)	0.93 (0.60-2.01)
IL-6	-0.10	ns	35.3 (25.1-54.4)	26.7 (18.6-44.1)	25.5 (18.3-46.7)
MCP-1	-0.19	0.002	13.0 (10.1-16.1)	11.7 (8.5-16.3)	10.5 (8.4-13.2)
MIP-1 $\alpha$	-0.22	3.4E-4	5.24 (4.32-6.64)*	4.77 (3.97-5.48)	4.42 (3.74-5.39)
<b>Matrix proteases</b>					
MMP-3	-0.13	0.03	2.69 (1.90-3.42)	2.41 (2.03-3.07)	2.23 (1.77-2.87)
MMP-7	-0.08	ns	503 (322-628)	353(272-530)	384 (293-547)
MMP-12	-0.12	0.04	153 (96-238)**	116 (84-145)	117 (85-178)
<b>Apoptosis</b>					
TNFR-1	-0.15	0.01	6700 (5230-7954)*	5673 (4656-7145)	5576 (4713-6865)
TRAILR-2	-0.17	0.004	3.52 (2.36-4.37)	2.87 (2.03-4.00)	2.76 (1.99-3.51)
Fas	-0.22	2.6E-4	224 (181-271)	191 (166-241)	184 (159-222)
<b>SMC Growth factors</b>					
PDGF	0.27	6.0E-6	67.6 (26.9-177.3)**	124.5 (51.3-286.5)	168.3 (89.6-257.8)
EGF	0.25	2.5E-5	31.8 (7.09-72.8)**	46.9 (17.6-128.7)	65.1 (31.1-113.4)
HBEGF	0.15	0.01	24.0 (20.6-28.3)*	25.2 (21.5-39.9)	26.4 (22.0-35.2)
<b>EC growth factors</b>					
HGF	-0.23	1.3E-4	113 (93-138)*	97 (72-125)	92 (75-109)
PIGF	-0.22	2.9E-4	192 (167-248)*	167 (135-231)	155 (134-187)
VEGF	-0.16	0.007	1468 (1220-1898)	1342 (1029-1761)	1269 (1038-1593)

All biomarkers except hsCRP are expressed as arbitrary units and values are shown as median and IQR. Correlations are shown as Spearman rank correlation coefficients. Differences between IMT change categories (regression and progression versus no change) were calculated with one-way ANOVA and Scheffe's post hoc test. \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001. If correcting for multiple comparisons in the present table, the threshold for significance is p<0.004. SMC; smooth muscle cell, EC; endothelial cell.

**Supplemental table 9.** Relationships between biomarkers and change in CCA IMT in subjects with T2D. Related to table 4.

	Correlation	P	Regression (n=122)	No change (n=256)	Progression (n=186)
<b>Inflammatory</b>					
hsCRP (mg/L)	0.05	ns	1.40 (0.69-3.60)	1.45 (0.66-3.10)	1.67 (0.86-3.00)
IL-6	0.01	ns	41.3 (26.0-64.7)	36.8 (27.3-55.7)	38.6 (26.3-59.3)
MCP-1	-0.08	ns	13.4 (10.2-16.8)	11.5 (9.1-15.9)	12.2 (9.4-15.7)
MIP-1 $\alpha$	0.01	ns	4.86 (3.77-5.83)	4.72 (4.00-5.66)	4.72 (3.97-5.84)
<b>Matrix proteases</b>					
MMP-3	-0.04	ns	2.68 (2.04-3.32)	2.31 (1.91-3.05)	2.51 (1.94-3.16)
MMP-7	0.12	0.004	399 (193-696)**	449 (308-686)	495 (347-718)
MMP-12	0.01	ns	1347(101-209)	149 (105-207)	147 (103-215)
<b>Apoptosis</b>					
TNFR-1	-0.06	ns	6608 (5557-8452)	6700 (5176-8135)	6562 (5221-8023)
TRAILR-2	-0.18	<0.0001	4.04 (3.11-5.06)**	3.43 (2.46-4.53)	3.27 (2.48-4.35)
Fas	-0.02	ns	215 (175-267)	208 (175-244)	212 (171-266)
<b>Growth factors</b>					
PDGF	0.09	0.037	88.0 (41.1-215.3)*	144.0 (67.2-247.7)	153.3 (72.0-245.6)
EGF	0.02	ns	42.5 (16.7-111.3)	55.3 (24.4-111.4)	58.1 (21.5-108.8)
HBEGF	0.06	ns	24.1 (18.5-33.4)	25.3 (20.5-32.7)	26.2 (20.7-33.6)
<b>EC growth factors</b>					
HGF	-0.04	ns	117 (93-143)	111 (92-136)	111 (91-141)
PIGF	-0.01	ns	184 (145-243)	171 (144-212)	167 (144-221)
VEGF	-0.03	ns	1520 (1209-1868)	1489 (1168-1898)	1389 (1128-1758)

All biomarkers except hsCRP are expressed as arbitrary units and values are shown as median and IQR. Correlations are shown as Spearman rank correlation coefficients. Differences between IMT change categories (regression and progression versus no change) were calculated with one-way ANOVA and Scheffe's post hoc test. \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001. If correcting for multiple comparisons in the present table, the threshold for significance is p<0.004. SMC; smooth muscle cell, EC; endothelial cell.

**Supplemental table 10.** Relationships between biomarkers and change in CCA IMT in subjects without T2D. Related to table 4.

	Correlation	P	Regression (n=59)	No change (n=122)	Progression (n=113)
<b>Inflammatory</b>					
hsCRP (mg/L)	-0.04	ns	0.96 (0.63-1.69)	0.98 (0.60-2.20)	0.95 (0.60-2.12)
IL-6	-0.04	ns	35.3 (21.6-50.6)	29.2 (18.3-46.4)	31.3 (20.0-47.8)
MCP-1	0.02	ns	11.8 (8.9-14.8)	11.2 (8.5-14.3)	11.6 (8.7-14.8)
MIP-1 $\alpha$	-0.01	ns	4.79 (3.93-5.80)	4.77 (4.02-6.02)	4.69 (3.92-5.82)
<b>Matrix proteases</b>					
MMP-3	0.06	ns	2.35 (1.93-2.99)	2.40 (1.79-3.04)	2.34 (1.90-3.35)
MMP-7	-0.12	0.05	501 (301-664)	392 (292-614)	383 (276-541)
MMP-12	-0.04	ns	1347(101-209)	149 (105-207)	147 (103-215)
<b>Apoptosis</b>					
TNFR-1	-0.06	ns	6383 (4873-7697)	5894 (4999-7345)	5873 (4837-7082)
TRAILR-2	-0.07	ns	3.48 (2.41-4.18)	2.85 (1.97-3.86)	2.85 (2.32-3.72)
Fas	-0.02	ns	202 (159-236)	195 (162-238)	194 (164-239)
<b>Growth factors</b>					
PDGF	0.05	ns	122.8 (43.0-200.9)	145.0 (58.3-254.2)	115.4 (57.3-240.5)
EGF	-0.03	ns	50.2 (16.6-108.8)	44.9 (20.6-115.2)	47.5 (18.9-94.4)
HBEGF	0.01	ns	25.6 (21.5-33.4)	25.6 (21.4-32.8)	24.9 (21.9-34.5)
<b>EC growth factors</b>					
HGF	-0.01	ns	101 (82-123)	96 (78-122)	96 (78-123)
PIGF	-0.04	ns	177 (146-226)	174 (140-208)	170 (134-218)
VEGF	0.01	ns	1389 (1105-1652)	1342 (1044-1675)	1389 (1053-1687)

All biomarkers except hsCRP are expressed as arbitrary units and values are shown as median and IQR. Correlations are shown as Spearman rank correlation coefficients. Differences between IMT change categories (regression and progression versus no change) were calculated with one-way ANOVA and Scheffe's post hoc test. \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001. If correcting for multiple comparisons in the present table, the threshold for significance is p<0.004. SMC; smooth muscle cell, EC; endothelial cell.

**Supplemental table 11.** Relationships between baseline vascular measurements and change in carotid bulb IMT. Related to table 5.

	<b>T2D</b>				
	Correlation	P	Regression	No change	Progression
<b>CCA</b>			n=126	n=125	n=269
Mean IMT (mm)	0.03	ns	0.89 (0.80-1.02)	0.88 (0.77-0.98)	0.86 (0.74-0.99)
<b>Bulb</b>					
Mean IMT (mm)	0.02	ns	1.23 (1.01-1.63)***	1.06 (0.89-1.23)	1.11 (0.94-1.37)
GSM baseline	0.16	0.001	43 (31-69)***	65 (49-84)	67 (46-85)
GSM follow-up	0.23	6.9E-5	51 (35-68)**	67 (49-91)	60 (46-83)
<b>Arterial stiffness</b>					
PWV (m/s)	0.12	ns	11.2 (9.3-13.3)	10.2 (8.9-12.3)	10.7 (9.2-12.6)
<b>Endothelial function</b>					
RHI	0.00	ns	1.97 (1.68-2.45)	2.11 (1.66-2.58)	2.04 (1.65-2.57)
	<b>Non-T2D</b>				
<b>CCA</b>			n=65	n=75	n=143
Mean IMT (mm)	0.00	ns	0.86 (0.80-1.02)	0.83 (0.74-0.984)	0.87 (0.75-0.98)
<b>Bulb</b>					
Mean IMT (mm)	-0.16	0.01	1.21 (1.01-1.49)***	0.99 (0.87-1.16)	1.01 (0.87-1.26)
GSM baseline	0.25	1.9E-4	45 (31-78)**	68 (49-94)	69 (56-87)
GSM follow-up	0.23	0.007	59 (43-75)	70 (55-87)	72 (50-108)
<b>Arterial stiffness</b>					
PWV (m/s)	-0.07	ns	10.2 (9.2-12.6)**	9.0 (7.9-10.4)	9.7 (8.3-11.4)
<b>Endothelial function</b>					
RHI	-0.03	ns	2.31 (1.91-2.83)	2.36 (2.04-2.71)	2.38 (1.90-2.75)

Grey scale median (GSM) value is the median of all grey levels of the pixels in the plaque. PWV; pulse wave velocity, RHI; reactive hyperemia index. Correlations are shown as Spearman rank correlation coefficients. Differences between IMT change categories (regression and progression versus no change) were calculated with one-way ANOVA and Scheffe's post hoc test. Values are shown as median and IQR. \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001.

**Supplemental table 12.** Relationships between baseline vascular measurements and change in CCA IMT. Related to table 5.

	<b>T2D</b>				
	Correlation	P	Regression (n=122)	No change (n=256)	Progression (n=186)
<b>CCA</b>					
Mean IMT (mm)	-0.23	4.2E-8	0.98 (0.86-1.15)***	0.88 (0.77-0.99)	0.86 (0.74-1.00)
<b>Bulb</b>					
Mean IMT (mm)	-0.02	ns	1.13 (0.94-1.45)	1.06 (0.93-1.34)	1.11 (0.94-1.37)
GSM baseline	0.13	0.007	51 (37-79)	62 (44-84)	67 (46-85)
GSM follow-up	0.12	0.034	53 (42-73)	68 (43-91)	60 (46-83)
<b>Arterial stiffness</b>					
PWV (m/s)	-0.09	ns	11.6 (9.9-13.7)	10.5 (8.9-13.1)	10.7 (9.2-12.6)
<b>Endothelial function</b>					
RHI	0.05	ns	1.97 (1.70-2.43)	2.03 (1.67-2.49)	2.06 (1.75-2.47)
	<b>Non-T2D</b>				
			Regression (n=126)	No change (n=125)	Progression (n=269)
<b>CCA</b>					
Mean IMT (mm)	-0.24	3.1E-5	0.94 (0.83-1.12)***	0.85 (0.75-0.98)	0.83 (0.73-0.92)
<b>Bulb</b>					
Mean IMT (mm)	0.03	ns	1.05 (0.90-1.35)	1.03 (0.88-1.29)	1.06 (0.87-1.30)
GSM baseline	-0.03	0.001	63 (37-78)	70 (46-8)	60 (40-82)
GSM follow-up	0.03	6.9E-5	62 (48-75)	70 (48-92)	64 (49-86)
<b>Arterial stiffness</b>					
PWV (m/s)	-0.12	0.04	10.2 (9.2-11.5)	9.5 (8.3-11.6)	9.5 (8.2-11.0)
<b>Endothelial function</b>					
RHI	0.00	ns	2.26 (1.83-2.74)	2.44 (1.99-2.80)	2.35 (1.99-2.74)

Grey scale median (GSM) value is the median of all grey levels of the pixels in the plaque. PWV; pulse wave velocity, RHI; reactive hyperemia index. Correlations are shown as Spearman rank correlation coefficients. Differences between IMT change categories (regression and progression versus no change) were calculated with one-way ANOVA and Scheffe's post hoc test. Values are shown as median and IQR. \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001.

**Supplemental table 13.** Biomarkers in T2D subjects without and with insulin treatment. Related to table 3.

	No insulin (n=419)	Insulin (n=143)	P
<b>Inflammatory</b>			
hsCRP (mg/L)	1.43 (0.67-2.94)	1.80 (0.84-3.51)	0.04
IL-6	36.6 (25.4-56.2)	41.6 (31.8-64.9)	0.007
MCP-1	11.9 (9.2-15.9)	12.6 (10.0-16.3)	ns
MIP-1 $\alpha$	4.68 (3.95-5.67)	4.92 (4.00-5.90)	ns
<b>Matrix proteases</b>			
MMP-3	2.45 (1.97-3.05)	2.27 (1.87-3.27)	ns
MMP-7	428 (290-665)	532(356-733)	0.001
MMP-12	140 (101-205)	170 (115-232)	0.003
<b>Apoptosis</b>			
TNFR-1	6427 (5221-7968)	7231 (5595-8964)	0.001
TRAILR-2	3.40 (2.52-4.53)	3.76 (2.71-5.31)	0.02
Fas	211 (175-256)	208 (168-263)	ns
<b>SMC Growth factors</b>			
PDGF	137.2 (57.3-242.2)	160.9 (71.0-240.5)	ns
EGF	51.1 (19.8-104.0)	61.0 (29.4-124.5)	0.045
HBEGF	25.5 (19.8-33.1)	25.3 (20.0-32.7)	ns
<b>EC growth factors</b>			
HGF	110 (92-137)	120 (83-144)	ns
PIGF	171 (143-214)	177 (146-228)	ns
VEGF	1428 (1160-1833)	1563 (1209-1859)	ns

All biomarkers except hsCRP are expressed as arbitrary units and values are shown as median and IQR. Comparisons between the groups were done using the Mann-Whitney U test. If correcting for multiple comparisons in the present table, the threshold for significance is  $p < 0.004$ . SMC; smooth muscle cell, EC; endothelial cell.



**Supplemental table 14.** Vascular measurements in T2D subjects without and with insulin treatment. Related to table 5.

	No insulin (n=419)	Insulin (n=143)	P
Bulb baseline IMT (mm)	1.06 (0.93-1.31)	1.22 (1.00-1.56)	<0.001
CCA baseline IMT (mm)	0.89 (0.77-1.03)	0.90 (0.81-1.03)	ns
Progression bulb IMT (mm)	0.06 (-0.03-0.17)	0.06 (-0.11-0.20)	ns
Progression CCA IMT (mm)	0.02 (-0.04-0.08)	0.00 (-0.05-0.06)	(0.05)
GSM baseline	64 (44-89)	59 (39-79)	ns
GSM follow-up	61 (44-84)	62 (43-85)	ns
PWV (m/s)	10.5 (8.9-12.8)	12.0 (9.9-14.0)	<0.001
RHI	2.05 (1.73-2.51)	1.97 (1.65-2.35)	ns

Grey scale median (GSM) value is the median of all grey levels of the pixels in the plaque. PWV; pulse wave velocity, RHI; reactive hyperemia index. Comparisons between the groups were done using the Mann-Whitney U test.