ESM Table 2

| Gene | Type 2 diabetes without MVD | Type 2 diabetes with PAD | Tye 2 diabetes with CAD | Healthy control | Non-type 2 diabetes with PAD | Non-type 2 diabetes with CAD | p-value |
|--------|--------------------------------|--------------------------------|-------------------------------|--------------------|------------------------------------|------------------------------------|---------|
| SMC | | | | | | | • |
| ACTA2 | 1.4 ± 0.4 | $0.8\pm0.1\;\Delta$ | $0.6\pm0.1\;\Delta$ | 1.0 ± 0.3 | 0.4 ± 0.2 | $0.7\pm0.2~\Delta$ | NS |
| SMT | 2.0 ± 0.6 | 1.9 ± 0.6 | 1.6 ± 0.4 | 1.0 ± 0.2 | 1.1 ± 0.3 | 1.6 ± 0.3 | NS |
| SRF | 1.1 ± 0.2 | 1.0 ± 0.1 | 1.0 ± 0.6 | 1.0 ± 0.1 | 0.8 ± 0.1 | 1.1 ± 0.2 | NS |
| KLF4 | 1.4 ± 0.4 | 1.2 ± 0.2 | 1.7 ± 0.6 | 1.0 ± 0.2 | 0.9 ± 0.2 | 0.9 ± 0.2 | NS |
| SMAD4 | 1.5 ± 0.2 | 1.1 ± 0.1 | 1.1 ± 0.1 | 1.0 ± 0.04 | $0.9\pm0.04~^a$ | 1.0 ± 0.03 | < 0.05 |
| TGFB1 | 1.3 ± 0.2 | 1.3 ± 0.1 | 1.7 ± 0.2 | 1.0 ± 0.1 | 1.0 ± 0.1 | $1.1\pm0.04~\Delta$ | NS |
| TGFBR1 | 1.5 ± 0.4 | 1.1 ± 0.1 | 1.1 ± 0.2 | 1.0 ± 0.1 | 1.0 ± 0.1 | 1.1 ± 0.2 | NS |
| TGFBR2 | 1.3 ± 0.2 | 0.9 ± 0.1 | 1.0 ± 0.1 | 1.0 ± 0.1 | 0.8 ± 0.1 | 1.1 ± 0.1 | NS |
| BMP4 | 1.1 ± 0.4 | 0.5 ± 0.1 | 0.3 ± 0.1 | 1.0 ± 0.3 | 0.4 ± 0.1 | 0.5 ± 0.1 | NS |
| BMP6 | 3.4 ± 1.3 | $5.9\pm1.4^{\text{b}}$ | $0.7\pm0.4~\Delta$ | 1.0 ± 0.4 | 1.3 ± 0.4 | 4.0 ± 1.8 | < 0.05 |
| PDGFBB | 1.7 ± 0.3 | 1.4 ± 0.3 | 1.0 ± 0.3 | 1.0 ± 0.2 | 1.2 ± 0.1 | 1.2 ± 0.2 | NS |
| EC | | | | | | | |
| KDR | $0.5 \pm 0.2 \Delta$ | 0.2 ± 0.1 | 0.5 ± 0.2 | 1.0 ± 0.4 | 0.3 ± 0.1 | 0.3 ± 0.2 | NS |
| VEGFA | $0.9\pm0.2\;\Delta$ | 1.1 ± 0.1 | 1.3 ± 0.2 | 1.0 ± 0.2 | 1.2 ± 0.3 | 1.5 ± 0.4 | NS |
| NOS3 | $0.8\pm0.3\;\Delta$ | 1.4 ± 0.5 | $0.4\pm0.1\;\Delta$ | 1.0 ± 0.3 | $0.8\pm0.2\;\Delta$ | 1.1 ± 0.2 | NS |
| CDH5 | 2.9 ± 1.0 | 8.3 ± 3.6 | $1.5\pm0.6~\Delta$ | 1.0 ± 0.5 | 1.6 ± 0.4 | 1.4 ± 0.7 | NS |
| PECAM1 | 2.0 ± 0.7 | 1.0 ± 0.1 | 1.0 ± 0.1 | 1.0 ± 0.1 | 0.9 ± 0.1 | 1.1 ± 0.1 | NS |

Table 2. Gene expression of SMC and EC lineage genes in cultured CACs.

Data are expressed as relative mRNA expression +/- SEM compared to the healthy control group. N=6 per group unless indicated by Δ (N=5). *ACTA2*, α -Smooth muscle actin; *SMT*, Smoothelin, *SRF*, Serum response factor; *KLF4*, Krüppellike factor 4; *SMAD4*; Mothers against decapentaplegic homolog 4; *TGFB1*, Transforming growth factor- β ; *TGFBR1*, Transforming growth factor- β receptor 1; *TGFBR2*, Transforming growth factor- β receptor 2; *BMP4*, Bonemorphogenetic protein 4; *BMP6*, Bone-morphogenetic protein 6; *PDGFBB*, Platelet-derived growth factor-bb; *KDR*, Kinase domain receptor; *VEGFA*, Vascular endothelial growth factor A; *NOS3*, Endothelial nitric oxide synthase; *CDH5*, *Vascular endothelial cadherin; PECAM1*; Platelet endothelial cell adhesion molecule 1

Statistically significant with ANOVA compared to: ^anon-T2DM with PAD vs. non-T2DM with CAD ^bT2DM with PAD vs. T2DM with CAD.