Supplementary figures



Figure S1. Pericyte staining in human melanoma tissues.

Representative images of immunohistochemical staining for NG2 (upper panels, brown) and desmin (lower panels, brown) in VM+ human melanoma tumor tissues. The panels on the right show negative controls (no primary antibodies used) and positive controls (muscle tissue).







Figure S3. Quantification of perivascular coverage (α SMA+ or α SMA-) of blood vessels (CD31+) in tumors from VM+ (C8161) or VM- (OCM-1) cell lines.



Figure S4. PAS loop formation in cultured tumor cells in the presence or absence of HVSCs.

(A) PAS stained co-cultures of both VM+ (left panels) and VM- (middle and right panels) cell lines in the absence (upper panels) or presence (middle panels) of HVSC perivascular cells. (B) PAS stained co-cultures of VM+ cells combined with VM- cells or HVSCs alone.



Figure S5. Vascular-like network formation in cultured tumor cells in the presence or absence of HVSCs.

(A) Formation of vascular-like networks on Matrigel by VM+ and VM- cell lines. (B) Co-localization of CFSE-labeled HVSCs (green) with vascular-like networks formed by C8161 cells at day 4. (C) Effect of different co-cultures on the stability of vascular-like networks formed on Matrigel by VM+ cell line C8161. Bottom two panels show absence of network formation by VM- cells even in the presence of HVSC. (D) Effect of normal culture medium or FCS-free conditioned medium of HVSCs on vascular-like network formation by C8161 cells.



Figure S6. Effect of hypoxia and matrix on PDGFB expression in VM+ and VM- cells.

(A) The levels of *PDGFB* mRNA in different VM+ and VM- cell lines (C81-61, C8161). (B) The level of secreted PDGFB protein as assessed by ELISA in culture medium of VM+ and VM- cell lines (C81-61, C8161). (C) Representative images of PDGFB immunostaining staining (brown; left panels) and HIF1alpha immunostaining (brown; right panels) in consecutive sections of tumor tissue from patients with cutaneous melanoma. No apparent colocalization of PDGFB and HIF1alpha could be observed. (D) The levels of *PDGFB* mRNA (left bar chart) or *VEGFA* (right bar chart) in VM+ (C8161) and VM- (OCM-1) cells cultured under normoxic (white bars) or hypoxic (black bars) conditions for 3 days. (E) Similar to (D) for cells cultured on Matrigel for 24 h. (F) Similar to (D) for cells cultured on Matrigel for 48 or 72 h in the presence of HSVCs.



Figure S7. The effect of conditioned medium (CM) from VM- C81-61 or VM+ C8161 cutaneous melanoma cells on cellular organization of actin stress fibers in HVSCs.





(A) Formation of vascular-like networks by B16F10 melanoma cells on Matrigel (right panel). (B)

Quantification of CD31+ endothelial cells in tumors from wildtype B16F10 cells or PDGFB

overexpressing B16F10 cells.





(A) Tumor volumes of VM+ (C8161) and VM- (OCM-1) xenograft tumors. * p<0.05. (B) Quantification of blood lakes and microvessel density in VM+ (C8161) and VM- (OCM-1) xenograft tumors. * p<0.05. (C) Representative pictures and quantification of CD31 staining in VM+ (C8161) and VM- (OCM-1) xenograft tumors treated with PBS or STI-571. * p<0.05. (D) quantification of CD31+ vessels not associated with α SMA+ perivascular cells in VM+ (C8161) xenograft tumors treated with PBS, STI-571 or blocking antibodies targeting either PDGF receptor alpha or PDGF receptor beta. * p<0.05 treatment versus PBS.