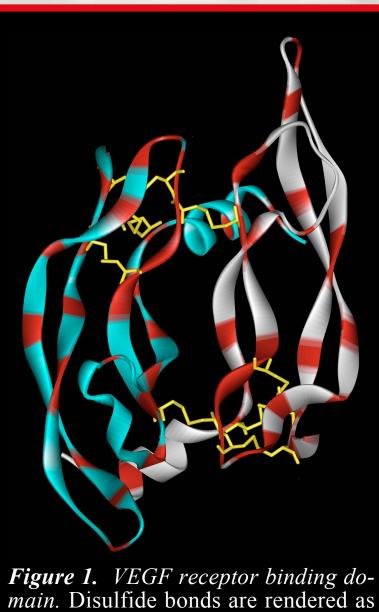
# Summary

Upon reaching a size of several millimeters a developing embryo cannot meet its metabolic needs by diffusion alone. Solid tumors face the same problem, when expanding to a similar magnitude. Both embryo and tumor can only continue growing by establishing a circulatory system to supply them with oxygen and nutrients. The obvious similarity between these processes is reflected on the molecular level: Vascular endothelial growth factor (VEGF, Fig. 1) has been identified as a key regulator of blood vessel growth in both embryos and tumors.

Recently we described two proteins homologous to VEGF: VEGF-C and VEGF-D. The structural similarity be-



yellow lines. Amino acids matching a superimposed VEGF-C molecule are shown in red.

tween VEGF-C and VEGF-D define a VEGF subfamily. Similar to Platelet-derived growth factor (PDGF) mature VEGF-C and D are generated by the cleavage of a precursor. VEGF-C plays an important role in the development of the lymphatic vascular system. The role of VEGF-D is less well defined.

Topics of our research include the further characterization of VEGF-C (biosynthesis, structure, determinants of receptor specificity) and the identification of the function of VEGF-D.

# **VEGF-D** Function

### **Transgenic mice with tissue-specific** overexpression of VEGF-D

To identify the function of VEGF-D we are establishing animal models with tissue-specific overexpression of VEGF-D analogous to the existing VEGF-C models using the keratin 14 and the rat insulin promoter.

## Recombinant **Growth Factors**

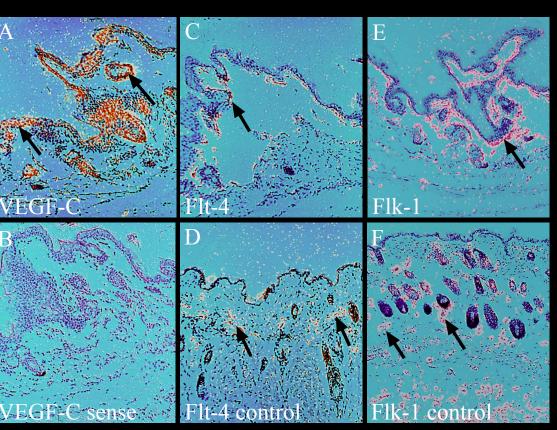
#### Baculoviral and **E.coli proteins**

Recombinant VEGF-C produced in the baculovirus system has been used in several studies (PEPPER, 1998; OH, 1997; JELTSCH, 1997; KUKK, 1996). Presently VEGF-C and VEGF-D are produced in E.coli for the production of monoclonal

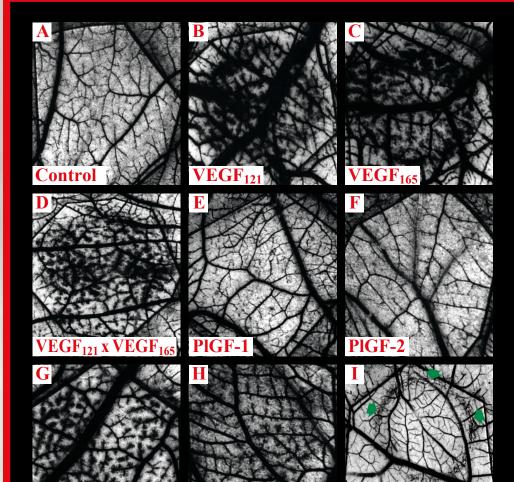


Figure 6. The CAM and its

developmental origin.

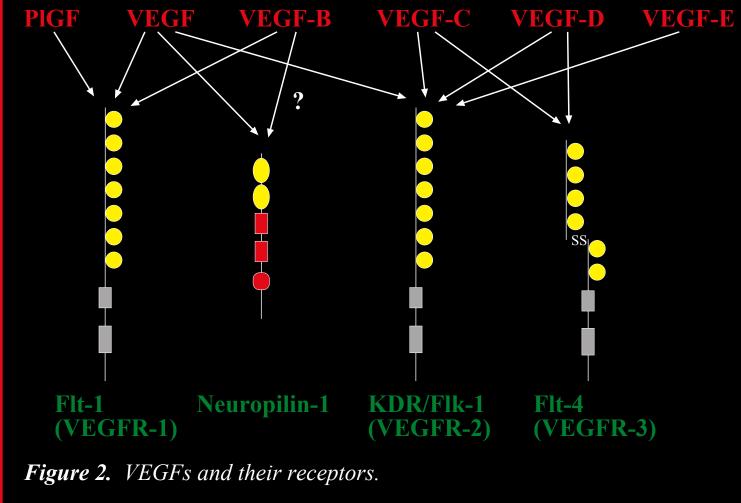


**Figure 5.** In-situ hybridisation analysis of K14-VEGF-C transgenic mice skin. A B: VEGF-C antisense and sense probes. C D: Flt-4. E F: Flk-1. Arrows in A indicate basal keratinocytes; in C through E, they point out the lymphatic endothelium; and in F, arrows show endothelial cells.



## **VEGFs and Their Receptors**

VEGF is an important regulator of endothelial cell proliferation and migration in embryonic vasculogenesis (the in-situ differentiation of endothelial cell precursors) and angiogenesis (the growth of new blood vessels from preexisting ones).



We lately cloned two factors homologous to VEGF (JOUKOV, 1997; ACHEN, 1998), designated as VEGF-C (or VRP) and VEGF-D (or FIGF). Both VEGF-C and D are ligands for KDR (the main

proliferationinducing receptor on blood vessel endothelial cells) and Flt-4, a receptor tyrosine kinase whose expression becomes restricted to lymphatic endothelium during developdevelopment (Fig. 2).

# **Completed Studies**

Hyperplasia of lymphatic vessels in VEGF-C transgenic mice

antibodies, in-vivo assays and crystallization purposes.

# **Structure & Function**

### **Random mosaic molecules and X-ray** crystallography

The crystal structure of VEGF has been resolved as well as the structure of the VEGF-Flt-1 complex. The interaction between VEGF and KDR has not been resolved at structural level and almost nothing is known about the KDR-VEGF-C interaction. To pinpoint the important elements of receptor specificity we started the determination of VEGF-C structure by X-ray crystallography and the creation of a mammalian expression library of 512 different VEGF-VEGF-C mosaic molecules, whose receptor specificities are correlated to the individual composition of the clone. Key features of this approach are the exploitation of the high homology of VEGF to VEGF-C in the core region to create the library and the use of an efficient screening system, that evaluates the receptor affinities of the individual clones.



Figure 7. Effect of VEGFs on CAM. A Carrier one doesn't induce vascular alterations. VEGF<sub>121</sub>, VEGF<sub>165</sub> and VEGF<sub>121</sub>xVEGF<sub>165</sub> het-erodimers are angiogenic. **E F** PlGF-1 and 2 are not angiogenic. **G H** VEGFxPlGF heterodimers induce angiogenesis. **I** VEGF-C doesn't induce gross changes of the vascular system, its effect shows only in the area of highest growth factor concentration (arrows). Magnification: x12.

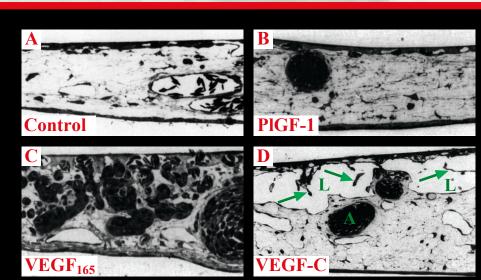
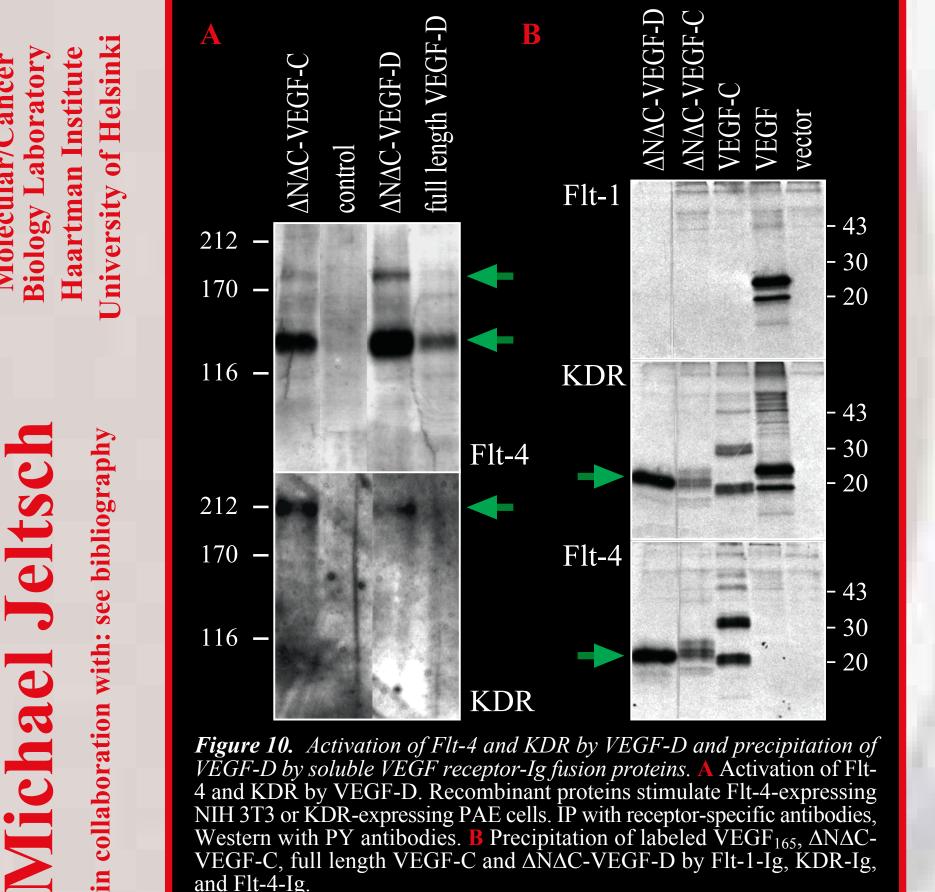


Figure 8. Semithin sections of treated CAM. capillaries (arrows) are found in the chorionic epithelium; arteries, veins, and lymphatics in the CAM stroma. B PIGF-1 has no effect. C VEGF<sub>165</sub>; note the amount of blood-filled capillaries. **D** VEGF-C: new lymphatics locate immediately under the chorionic epithelium. Note many intralu-minally located endothelial cells (arrows), indicative of plexus formation. A artery, L lymphatics,



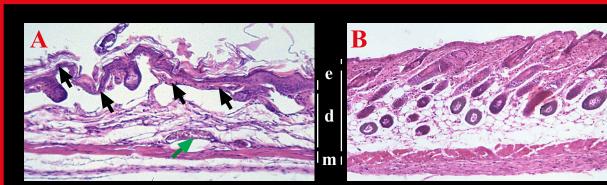


Figure 3. Skin of K14-VEGF-C transgenic mice. A HE-stained section of the skin of a 8-week-old transgenic mouse. Hyperkeratotic epidermis (e) showed underlying vessel spaces lined with endothelium (black arrows) but devoid of red cells (compare with the dermal vein shown with a green arrow). The dermis (d) is atrophic compared with the control littermate skin (**B**) and the muscle layer (m) is also reduced. Scale bar: 250 µm

VEGF-C overexpression in the basal epidermis of mice promotes the growth of lymphatic structures in the dermis, including large vessel lacunae histologically similar to lymphangioma (Figs. 3, 4, 5; JELTSCH, 1997).

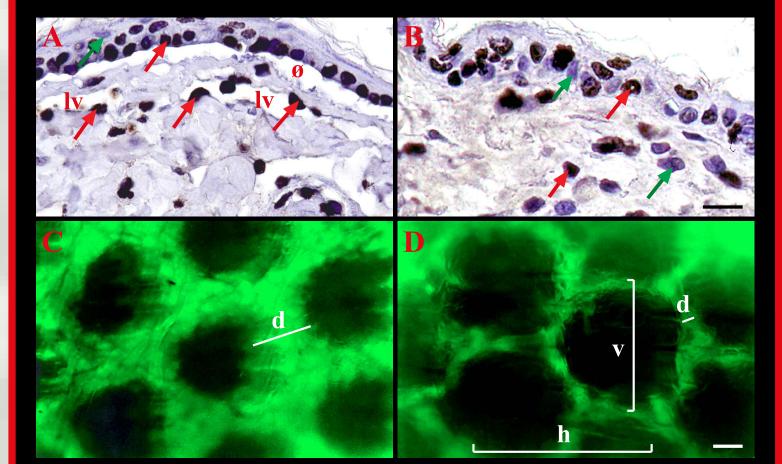
### Angiogenesis and lymphangiogenesis in the chorioallantoic membrane (CAM)

Due to the accessibility and regularity of its vascular system the CAM has been used for many years to evaluate potential angiogenic factors (Fig. 6). Less well documented is, that the CAM is drained by a dense network of lymphatic vessels.

The effects of VEGF, PIGF, VEGFxPIGF heterodimers and VEGF-C on the differentiated avian CAM were studied (Figs. 7, 8). Only VEGF-C is chemoattractive for lymphatic endothelial cells, induces their proliferation and the development of new lymphatic sinuses (OH, 1997).

### **VEGF-D** is a ligand of KDR/Flk-1 and Flt-4

VEGF-D was identified by computer-based homology search. It is most

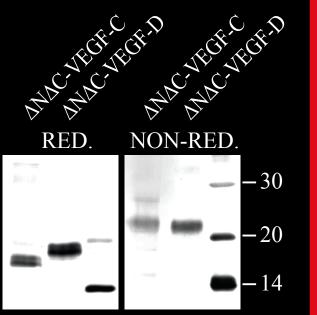


closely related VEGF-C (Fig. 9). VEGF-D activates KDR and

**Figure 10.** Activation of Flt-4 and KDR by VEGF-D and precipitation of VEGF-D by soluble VEGF receptor-Ig fusion proteins. A Activation of Flt-4 and KDR by VEGF-D. Recombinant proteins stimulate Flt-4-expressing NIH 3T3 or KDR-expressing PAE cells. IP with receptor-specific antibodies, Western with PY antibodies. **B** Precipitation of labeled VEGF<sub>165</sub>, ΔNΔC-VEGF-C, full length VEGF-C and ΔNΔC-VEGF-D by Flt-1-Ig, KDR-Ig, and Elt 4 Lg and Flt-4-Ig.

**More about** 

VEGF-C & D



**Figure 11.** Analysis of recombinant VEGF-D by Ag staining. Proteins were produced in HF cells and purified from conditioned medium by Ni<sup>2+</sup>NTA affinity chromatography

hVEGF-B <sub>167</sub>	PVSQP	D	APGHQRKVVS	WID.VYTRAT	COPREVVVPL	
hPlGF-2	VPPQQW	ALSAG	NGSSEVEVVP	FQE.VWGRSY	CRALERLVDV	
	141				190	
hVEGF-C	GKEFGVATNT	FFKPPCVSVY	RCGGCCNSEG	LOCMNTSTSY	LSKTLFEITV	
mVEGF-C	GKEFGAATNT	FFKPPCVSVY	RCGGCCNSEG RCGGCCNSEG	LOCMNTSTGY	LSKTLFEITV	
hVEGF-D	ASELGKSTNT	FFKPPCVNVF	RCGGCCNEES	LICMNTSTSY	ISKQLFEISV	
mVEGF-D	ASELGKTTNT	FFKPPCVNVF	RCGGCCNEES RCGGCCNEEG	VMCMNTSTSY	ISKOLFEISV	
hVEGF <sub>165</sub>	FOEYPDEIEY	IFKPSCVPLM	RCGGCCNDEG	LECVPTEESN	ITMOIMRIKP	
hVEGF-B <sub>167</sub>	TVELMGTVAK	<b>OLVPSCVTVO</b>	RCGGCCNDEG RCGGCCPDDG	LECVPTGOHO	VRMOILMIRY	
hPlGF-2			RCTGCCGDEN			
	191				239	
hVEGF-C	DI COCDUDIT	ISFANHTSCR	CMSKLDVYRQ	VHSTIRRSLP		
mVEGF-C	PLSOGPKPVT	ISFANHTSCR	CMSKLDVYRO	VHSIIRRSLP	ATLPOCOAA	
hVEGF-D	PLTSVPELVP	VKVANHTGCK	CLPTAP RH	PYSTIRRSIO	IPEEDRCSHS	
	PLTSVPELVP	VKTANHTGCK	CMSKLDVIRQ CLPTAPRH CLPTGPRH	PUSTTERSTO	TPEEDECPHS	
hVEGF	PLISVPELVP HQGQHIGE PSSQLGE	MSFLOHNKCE	CRPKKDR	ARO	ENPCGPCSER	
hVEGE-B	PSSO	MSLEEHSOCE	CRPKKKD	SAVKPDSP	RPL CPRCTOH	
hPlGF-2	C DPPSVVF	I.TESOHVECE	CRPLREK	MKD	FPPPP	
111 101 -2	G. DIGDIVE	HII BYIII CH	ON LINER.		Lidde	
	240				289	
hVEGF-C		NNHTOPOLAO	EDFMFSSDAG	DDSTDCTUDT		
mvEGF-C			QDFIFYSNVE			
hVEGF-D			EENPL.AGTE			
mvegf-D mvegf-D			DETPL.PGTE			
hVEGF <sub>165</sub>						
	но					
11VEGF - D <sub>167</sub>	ng		•••••	•••••		
	290				339	
hVEGF-C		PDASCOUVE	LDRNSCQCVC	WWWI FROCC		
mvegf-c			LDRDSCQCVC			
hVEGF-D						
mvegf-d mvegf-d	•••••			PTLCG		
mvege –D	• • • • • • • • • •	•••••	•••••	FILCG	PHMIP DEDRU	
	340				383	
hVEGF-C		NOPLNPCKCA	C.ECTESPQK	CLLKCKKENN		
mvegf-c			C.ECTENTQK			
hVEGF-C			CFECKESLET			
mVEGF-D			CFECKESLES			
			CSCKNTD.SR			
hVEGF <sub>165</sub>	· · · · · · · · · · · · ·					
hPlGF-2			KGRG KR	REEKORPIDC		
	384			419		
hVEGF-C		KACEDOECUC	EEVCRCVPSY			
			EEVCRCVPSY			
mVEGF-C						
hVEGF-D			KEKRAAQGPH			
mVEGF-D			KETR. AQGLY			
hVEGF <sub>165</sub>			• • • • • • • • • •			
hPlGF-2	.AV			••••PRR*		

Figure 9. Comparison of VEGFs. Color coding: utative) signal peptide, propeptides,

Jeltsch M (1997): Functional analysis of VEGF-B and VEGF-C. M. Sc. Thesis. University of Helsinki. http://www.helsinki.fi/~mjeltsch /msc.html

Jeltsch M, Kaipainen A, Joukov V, Meng X, Lakso M, Rauvala H, Swartz M, Fukumura D, Jain RK and K Alitalo (1997): Hyperplasia of lymphatic vessels in VEGF-C transgenic mice. Science 276, 1423-5.

Joukov V, Sorsa T, Kumar V, Jeltsch M, Claesson-Welsh L, Cao

**Figure 4.** Immunohistochemical analysis of endothelial proliferation and intravital fluorescence microlymphography of K14-VEGF-C transgenic mice. **A B**: Staining of cells in the S phase of the cell cycle (BrdU). In 14-day-old transgenic mice, the nuclear staining was observed in many endothelial cells of the lymphatic vessels (Iv) as well as in keratinocytes (red arrows in A). In nontransgenic littermates, mainly nuclei of epidermis keratinocytes and some dermal cells are stained (red arrows in B); unstained nuclei were observed in both cases (green arrows). **C D**: Fluorescence microscopy of lymphatic vessels of transgenic and control skin. The measured parameters are diameter (d) (in C, D) and horizontal (h) and vertical (v) mesh sizes (D only). Scale bars: A, B 5 μm; C, D 250 μm.

Flt-4. A short form of VEGF-D demonstrated receptorbinding resides in the VEGF homology domain which corresponds to the mature form of VEGF-C (Fig. 11; ACHEN, 1998).

#### Y, Saksela O, Kalkkinen N and K Alitalo (1997): Proteolytic processing regulates receptor specificity and activity of VEGF-C. EMBO J 16, 3898-911.

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Pepper MS, Mandriota SJ, Jeltsch M, Kumar V and K Alitalo (1998): Vascular endothelial growth factor (VEGF)-C synergizes with basic fibroblast growth factor and VEGF in the induction of angiogenesis in vitro and alters endothelial cell extracellular proteolytic activity. J Cell Physiol 177, 439-52.