

Supporting Information

Figure Legends for Supporting Information

Fig. S1. Multiple amino acid alignments among tissue-type (*A*) and venom-type (*B*) VEGFs.

Identical residues with *Vaa*-VEGF-A₁₆₆ (*A*) and barietin (*B*) are shaded and conserved cysteine residues are highlighted in black. Putative signal peptides and possessed sequences are shown in *italic-style lowercase*. The numbers at the top refer to residue numbers of Hs-VEGF-A₁₆₅ (*A*) and barietin (*B*) respectively. The secondary structural elements are shown as arrows for β -strands and cylinders for α -helices, and loops are labeled. C-terminal putative coreceptor-binding regions are boxed with dotted lines, and the reported heparin-binding sequences are marked with underlines. *B*, The six residues which are predicted to form a basic cluster in barietin are in bold letters. Note that the sequences of HF and ICPP are shown as mature proteins, because the cDNAs encoding HF and ICPP have not been reported (29,30).

Fig. S2. Nucleotide sequences of tissue- and venom-type VEGF genes from *T. flavoviridis*

A. Nucleotide sequence of tissue-type VEGF gene from *T. flavoviridis* (*Tf*-VEGF-A). Sequences of exons and introns are shown in bold capitals and lowercase, respectively.

B. Nucleotide sequence of venom-type VEGF gene from *T. flavoviridis* (*Tf*-svVEGF). Additive sequences that are only seen in the venom-type VEGF gene are shaded in gray.

C. Sequence alignment of introns of genes encoding tissue- and venom-type VEGFs from *Trimeresurus flavoviridis*. Identical nucleotides between tissue- and venom-type genes are shaded in gray.

Fig. S3. Sequence alignment of amino acids and nucleotides of tissue- and venom-type VEGFs from *Trimeresurus flavoviridis*

Identical nucleotides between tissue- and venom-type VEGF cDNAs are shaded in gray. The amino acid sequence of tissue-type (*Tf*-VEGF-A) is shown at the top, and that of venom-type (*Tf*-svVEGF) is on the bottom. Each exon is divided with a vertical bar, and numbered with Roman numerals. -; gap.

Fig. S4. Characterization of barietin, novel venom-type VEGF from the venom of *Bitis arietans*

A, Purification of barietin, novel venom-type VEGF from the venom of *Bitis arietans*. Barietin was purified by using three steps of chromatography: Superdex 200-pg gel-filtration, Q Sepharose High Performance, and Hi-Trap Heparin FPLC columns. Figure S4A shows a chromatogram of heparin affinity chromatography, the final step of purification. The fractions containing barietin were determined by anti-vammin antiserum reactivity (●), indicated with a bar on the top. SDS-PAGE (10% acrylamide) of purified barietin, with or without reducing agent, is indicated by *R* or *NR*.

B, Amino acid sequence deduced from the nucleotide sequence encoding barietin. Putative signal peptide and propeptide are indicated in lowercase italics. The numbers at the top refer to residue numbers of purified barietin. The determined internal amino acid residues by amino acid analysis are indicated with lines and arrows. *K*: endoprotease Lys-C; *D*: endoprotease Asp-N.

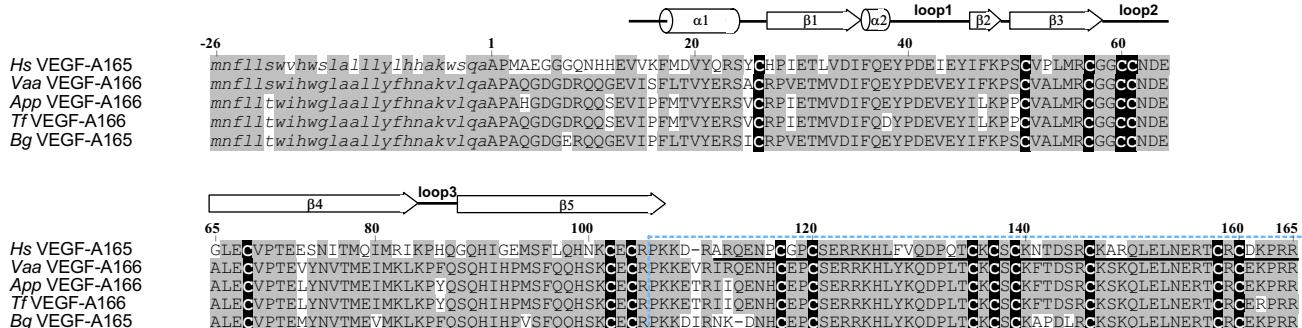
C, D, The molecular masses of purified barietin (*C*) and peptide-K2 (*D*) as analyzed by MALDI-TOF MS (average \pm SD, *n*=3).

Fig. S5. Multiple alignment of cDNA sequences encoding venom-type VEGFs

Conserved nucleotides among cDNAs are shaded in light (UTR) and dark gray (ORF), and the initial-codon (ATG) and stop-codon (TGA) are highlighted in black. The number on the left indicates nucleotides starting at the initial-codon. Each exon is divided with a vertical bar, and numbered with Roman numerals (*I* to *VI*). The nucleotide sequence of the 3'-UTR of *Tf*-svVEGF cDNA (10) was determined in this

study. The complete sequences of 5'- and 3'-UTRs of Pm-VEGF cDNA have been not reported (11). -; gap. Gaps in the C-terminal tail-coding region are highlighted in yellow.

A Tissue-type



B Venom-type

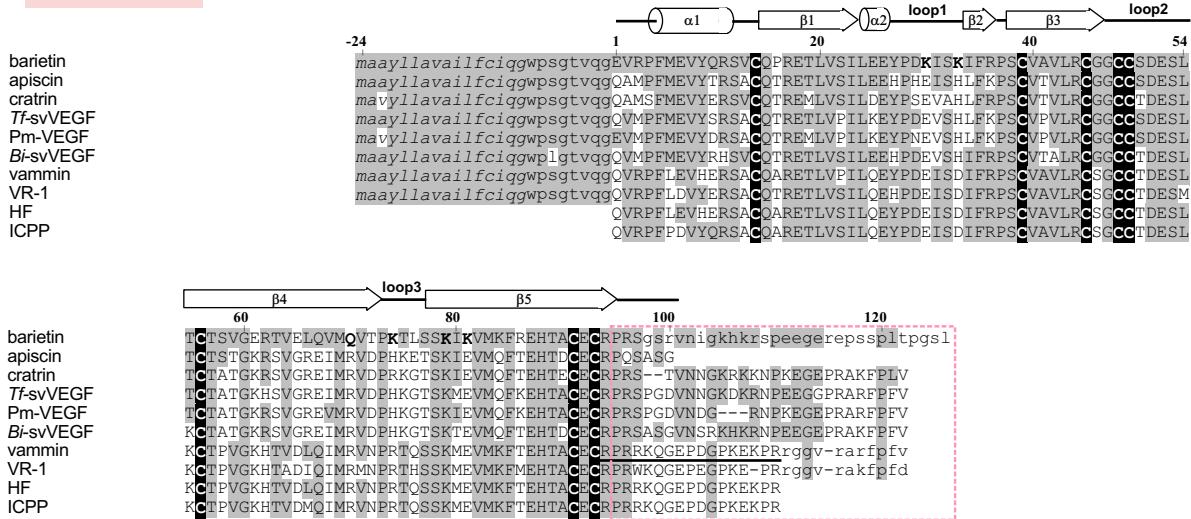


Fig. S1.

Fig. S2A. Tissue-type VEGF (Tf VEGF-A₁₉₀) gene

Fig. S2A. Tissue-type VEGF (*Tf* VEGF-A₁₉₀) gene (continued)

Fig. S2A. Tissue-type VEGF (*Tf*VEGF-A₁₉₀) gene (continued)

		(aa)
1	GGGAATGGACTGCCAGGCCCTGCCATGCTTCTGGGATTGCCCCACTCTGTGCCTGCTGTT GTCCACCGCCTTCGCTCTCTCTGCTGTTAACTCACCAACCCACCCACCC AGTCATGAAATTGCCCCAGCTCAGATCTCCCCATCTCTCTGAGCAGTGTGAAG CCAGGAGAGATAAGGCATGGCTGCCGACCTCTGAGGAGATCTCTGATCC m a a y l l a v a i l f c i	Exon I
	AGgtgagaaaactctgggttccctggatcttgctggatcaactctcacaggc q	-10
	aatttcagccaataagggggaaaaacagattaaatctgttatttagtcagggtaagc aataaaagtacttctaagggttggagaattcagcacccctgtcgatggataaaa ttacctctgttaggttgtgcagocctccactaacacacgtggccacccacccgggt ctggcaggatgtggatgcacagacgtgggtctgtggggagaagagcgcacccgt tgccccctgcaggagaccgcggcaggatgtggccgtcatccctctgttcactcgt 601 agcatagaagcttgcatttcaaaagttagccctgcattgtgaccactggcaggagc cccccaggacaaggcaactggcggagggttaaaacatccacaccacgtttcaaaat tggcacttgggtgtgtggacttccatccacccgtgtactggggatttgggaaattga agtccacattaaatgtggctaaattgtggatggatggggacttggggatggggact agaaccgtggctgggtgcagatgcattttgttgcacttctgtggttccctgt tcctgtggccatctgtgtatgtccgggtgcagatgtggccatctgtggggatccct ccttcttctgtcagGGCTGGCATCAGGGACAGTGCAGGGACAAGGcaagttccct g w p s g t v q g o	1
	tctaaagatcttatctggtaatgggggtgtgtccagagcatgcgtcttgccttggaaaa gagccggggggcagaaaaggggaccccccacccgtccgtccgtccggccagaacccttgg gtgagatccctggggacacacgcggccgtccacccgtcagatgtggggacaccc ctccctggggggggggcagaaagggttgcagggttgcagacgcgttcccccgggt ggggctggggggggggacatagccggctggagaagaggaggggactaccggccccccacc tgtttttccccaacccaggaaagggtgtgcgttgcggtaaaggcaagaggaggatctg aaggcgaggcagcgtctgacttttcccccctctcgtTGATGCCCTTATGGAAAGT V M P F M E V	II
	GTACAGCGCAGCGCTGCCAGACCCAGGGAGACACTAGTGCCTCATCTCAAAAGACTACCC Y S R S A C Q T R E T L V P I L K E Y P CGATGAAGTTTCCCACCTCTCAAGGCCCTCTGTGCTCTGTGTGGATGCCGTGGCTG D E V S H L F K P S C V P V L R C G G C CTGAGCGACGAAAGCCTCACGTGCACCGCTACGGGAAAGCACTCAGTCGGTGGAGgt C S D E S L T C T A T G K H S V G R E 67	III
	caggggctgttgcgtggatgtggggatgggggtgggggtgtcaggatgtcttgggg ggggcaggaaaggcaggccgcgttgcgtgggggtgtcaggatgtgggggggggggg cttcccttggtagatCATGCGGGTGGATCCCAACAGGGACTTCGAAGATGGGGT I M R V D P H K G T S K M E V	IV
1801	GATGCAATTCAAGGAGCACAGCCTGTGAATGCCAGtggagaaaggggggcgaaacagg M Q F K E H T A C E C R 94	Exon IV
	ctgtccctgggtggctgtcagagggtggggaaatgggggagaattgtgggggggggg atgacaagcagtcttagttcacacaacccctccctccctgttttctgtgttgc gttctcccaactttttatatttttttttttttttttttttttttttttttttttt ctattt caaaacgtcacccggctatttccacactgggggtctgcagggtctctgagcagg cagggtgggggtccacacgtgtcaggacccctctcaggggcagggttttgc acttctttctggagccccccacgcggggggccaaatctgtggctcaggatgt tgcaactgggttatt P R S P G D V N N G K	Exon V
	GGACAAGAGGtacatggggggggggggctggggtaacagaataagagatgttaaggga D K R 108	Exon V
2401	cctggggagggtttcttagtccaaacccctggctcaggttggctggggcaggctgggggg ggggcaggaggggcaagatggctgtgggggggggggggggggggggggggggggg AACCAGAGGAAGGGGGGCCAGAGGCCAGGGTCCCTTGTGACCAGCTTGTACTCC N P E G G P R A R F P V * 122	Exon VI
	AGACCCCTTGTGAGCTTCAACGCCCAACCAAGTGGGGAGGCTCTGTCTGCAAAGCCAGC TGGGGACGGCCCTGGTCCCTGTTCTCTCTTGTGACTGCTGGGGTGGGGAAAAGGG AGGCATCTCCAACATCTGGAGAAGTGTCTATCTACATCTACACTCTATGACACCCGG GCCTGGCTGGCCCTTCCTCATGTGGTGTGACCTGCACAAACACATCCTCCAGGTGCAA GGCCAAAGTCAAGGAGCGAACGGCAGCTCCCTCTCCAGTAACCTACCAATCAGGTGCAA TTTCAGGCTACGAAAGCCTCTTGACCAAGCTAATCTCTCTAAGGACTTGGCATT TTGTGTCAGGCTTAATGGCTTTAGTAAATTCGTTGTAAATACACAATTTTAAATT AAGATTGATATCTCCGGACCCCTCATCGGGGGAGCTGGGGAAAACAGTTCCTTTAA GAATGTGGATTAATGCTCTTGTGGGGCTGGGGGGGGGGGGGGGGGGGGGGGGGGGG AATAACGGAGTAAACT (3,137 bp)	Exon VI

Fig. S2B. Venom-type VEGF (*Tf*-svVEGF) gene

Intron 1 Tissue- vs venom-type genes

Fig. S2C.

Intron 1 Tissue- vs venom-type genes (*continued*)

	(bp)	
tissue-type	3071	cagtgggtcacccctcccttttagtcatgcagctgaagggtggaaatcttctccgataaaaaaa tggatgttcagccctttcgatgtatccatgtatcccattggccatgcgttttttccttagtgcata tgccatgtcagaaatccctccattggccatgcgttttttccttagtgcata ggggggaggaggaggaggagaactgcagcattaggcagaaaaaaaattagaggagggtttctcc cccccccccccccgcacacttaggcattgggtttactctgtatcccaggggccctccaaag tggctggattgttgcgtccagtc当地aaacagttacaagctttatgtgaggctgtgagatta aggggagggggaggaggaaaagctagaaaagaatgtttatgtgaaagaataggaaatgtgt gtttgtgtgttatgcgtgtgttagaagcaatgtaccaggtaaccttacttgctt catataggaatgagtaggatataagaatagaattgcagctccctgc当地actgtgtgctcaa gccttaaaagagatattctctactgtctgtctgaaaactctaccaggagacaattaa tgtcttaatggggaaacaatgagtaaaacccatcaggccaggaaaaagagatggtaacaagcaat tgacccatc当地acccatgttgc当地gagaaaaagggtttgtcttc当地ttgtggaa caggggacatattgc当地ccatctaagccctgtatctgggttaacggggatggccagaaaaat gttaattcatcagcatggggcagagaaaaacctcagattctcaatggatcaaggtgactct gctgtataaaatagctcttagattcaaaaggccaaacggcatgagcaatgtgtgaaatagattgtac tgc当地ggcttaatggggttctttc当地ttgtctgatgggtgaaatatactgttccagttctgctaa tcggggaaatagtagtctagccctgacaggcagttgaaagagagaaaaagagacagatggac agacatc当地ccatgttgc当地atcttctgtatggatcttctgtatggatcttct aatgatc当地ccacacacacacacacacacacacacacacacacacacac acacacacacagggotttgagagaaaagagagaaaagctgtttggacaaggtaagaaa cagcagcaagaatgttccatcaggccaaactttagggccagaagtaaggaaacagttctagctgatgc cgcaggacccaagcagggttgc当地gagaaatgaaatgtgggggtgtctccccc当地ggccctgac caagttatggcagggtctcatgttgc当地gtctcagtc当地ggggacttgc当地ggggcagcatgttgc cagttctcatctctc当地gtatggggatccccc当地ggggcagcatgttgc当地ggggcagcatgttgc tacgtctcatc当地gtatggggatccccc当地ggggcagcatgttgc当地ggggcagcatgttgc ggccactc当地ggggatccccc当地ggggcagcatgttgc当地ggggcagcatgttgc ctgatgttgc当地gtatggggatccccc当地ggggcagcatgttgc当地ggggcagcatgttgc gtttgttaaagttactgtgaccttccatccc当地ggggatccccc当地ggggcagcatgttgc tgc当地ccatgttgc当地gtatgactc当地gttgc当地gtatccatgttgc当地 caatagatc当地gttccatccc当地ggggatccccc当地ggggcagcatgttgc当地 gaaatgtctccaaatc当地ggggatccccc当地ggggcagcatgttgc当地 aacccctgttccatgttgc当地gtatggggatccccc当地ggggcagcatgttgc gattccagactccatataatttgc当地aaatc当地ggggatccccc当地ggggcagcatgttgc ggtaatgtgacccctgttgc当地gtatggggatccccc当地ggggcagcatgttgc tccatcttttagatgtccatattttcttccatccc当地ggggatccccc当地ggggcagcatgttgc gtttgtatgtttaatcttataataatgttccatggggatccccc当地ggggcagcatgttgc tgagagtgtatccggagaaaacttttgc当地gtatggggatccccc当地ggggcagcatgttgc ataagcttaatttgc当地gtatggggatccccc当地ggggcagcatgttgc ttc当地ctgtatggggatccccc当地ggggcagcatgttgc当地 tc当地gtatgttgc当地gtatggggatccccc当地ggggcagcatgttgc tagggactgaaatgttgc当地gtatggggatccccc当地ggggcagcatgttgc ctgaatc当地aaatc当地gtatggggatccccc当地ggggcagcatgttgc atttaaaccccttttagaaaaacaggtaagccacagtttgc当地acttgc当地 tcaaaaggggatccccc当地ggggatccccc当地ggggcagcatgttgc atttacaggatattttgc当地gtatggggatccccc当地ggggcagcatgttgc tttccaaaggaaatgttgc当地gtatggggatccccc当地ggggcagcatgttgc gtttagaaacataatttgc当地gtatggggatccccc当地ggggcagcatgttgc ttctgc当地gtatggggatccccc当地ggggcagcatgttgc当地 qqctqagggcaccqatttgc当地gtatggggatccccc当地ggggcagcatgttgc当地
	3786	
	4436	
	5086	
	5736	
	6191	

Fig. S2C. (*continued*)

Intron 2 Tissue- vs venom-type genes

Fig. S2C. (*continued*)

Intron 2 Tissue- vs venom-type genes (*continued*)

Fig. S2C. (*continued*)

Intron 3 Tissue- vs venom-type genes

Fig. S2C. (continued)

Intron 4 Tissue- vs venom-type genes

Fig. S2C. (*continued*)

Intron 5 Tissue- vs venom-type genes

Fig. S2C. (*continued*)

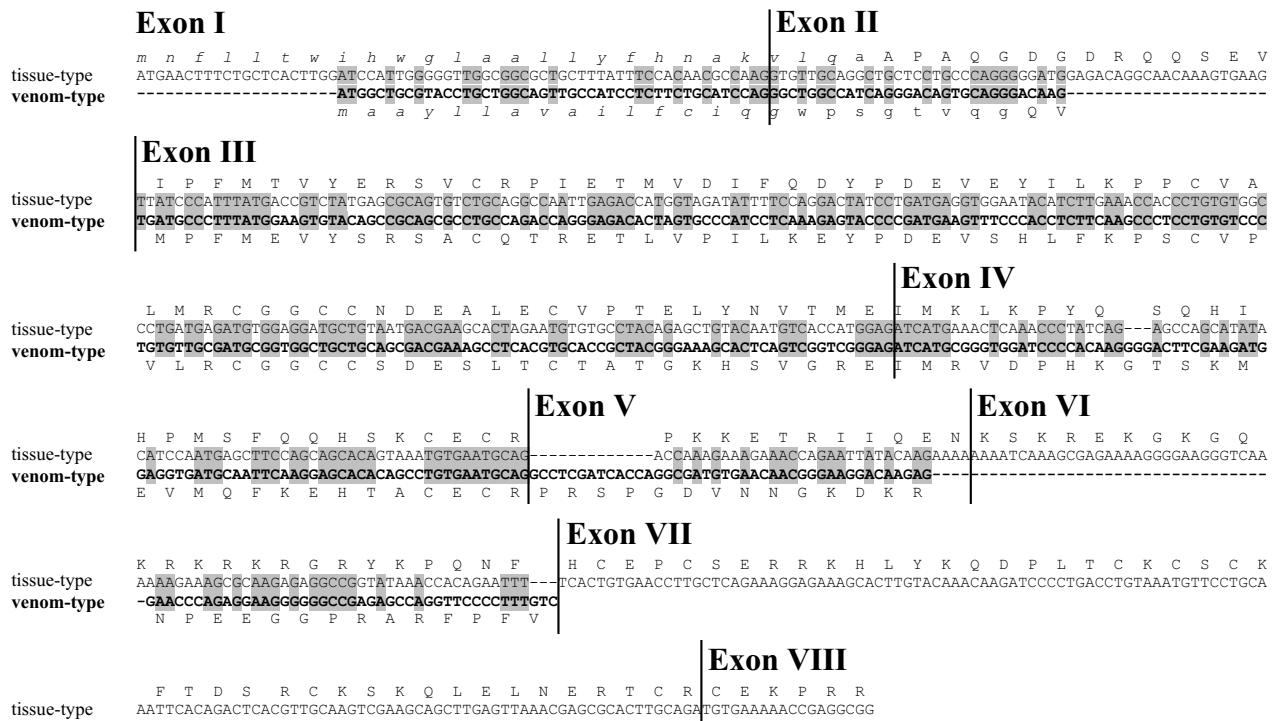


Fig. S3.

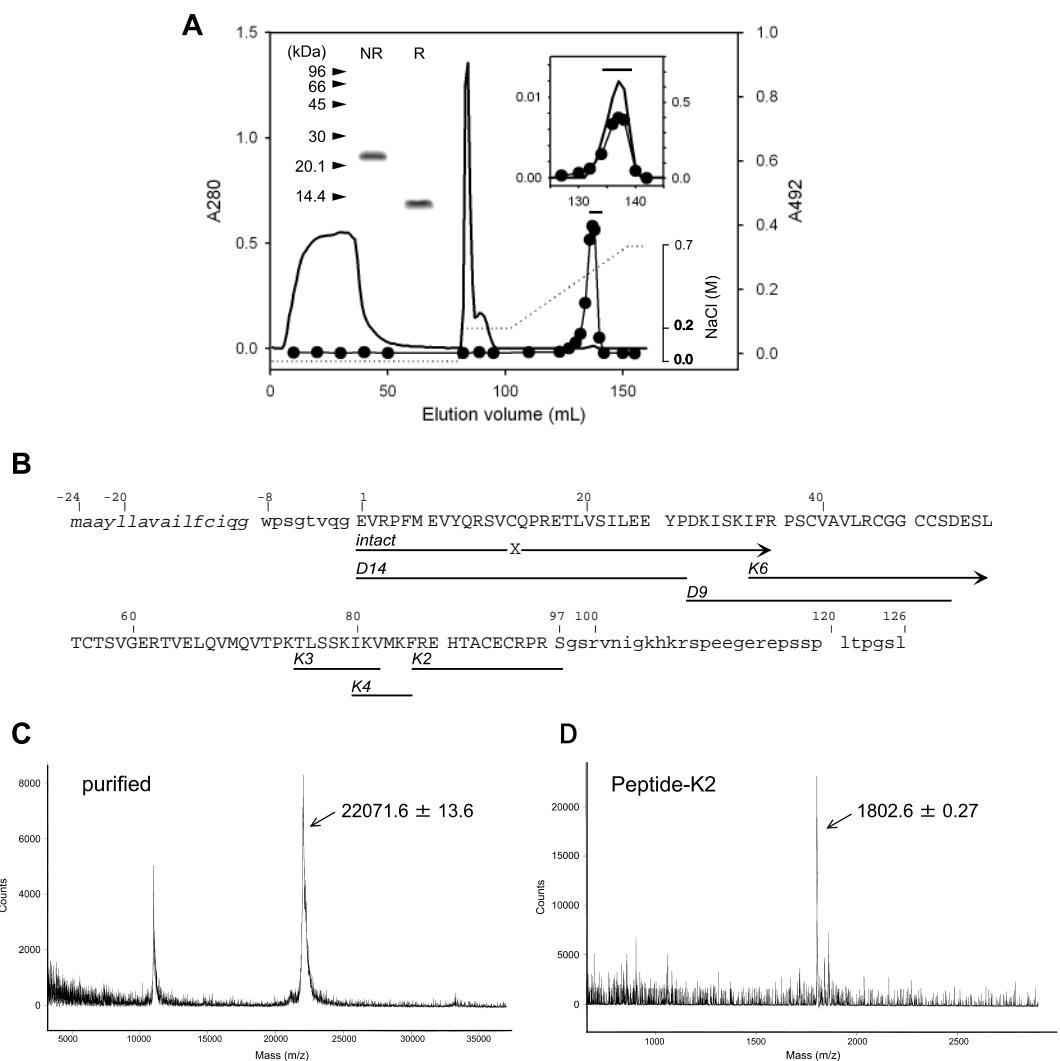


Fig. S4.

Exon I

barietin	-145	GGAGTAGACCGCAGGGAAACGGACGCCAGGCC-----TTGCCCCACTCTGTGCTGCTGTGTCACCCGGCTCTGCCT
apiscin	-184	ACTGGCAGCCCCGTCGA-GCTTCTGGCTTTGCCACTCTGTGCTGCTGTGTCACCCGGCTCTGCCT
cratrin	-217	CAGACAGCAGATCCAGGGAAACGGACTGCCAGGGCTCCA-GCTTCTGGCTTTGCCACTCTGTGCTGCTGTGTCACCCGGCTCTGCCT
Tf-svVEGF	-197	GGGAATGGACTGCCAGGCCCTGCCATGCTCTGGGATTGCCACTCTGTGCTGCTGTGTCACCCGGCTCTGCCT
Pm-VEGF	-173	CA-GCTTCTGGCTTTGCCACTCTGTGCTGCTGCTGTGTCACCCGGCTCTGCCT
Bi-svVEGF	-185	GCCAGGCCCTGCCA-GCTTCTGGCTTTGCCACTCTGTGCTGCTGCTGTGTCACCCGGCTCTGCCT
vammin	-134	TGCCCCACTCTGTGCTGCTGTGTCACCCGGCTCTGCCT
VR-1	-131	TGCCCCACTCTGTGCTGCTGTGTCACCCGGCTCTGCCT

Exon II

barietin	-47	CTCTCTTGTCTGTG-----AGATCTCACCCCATCTCTCTGAGCAGCTGTGAAGCCAGGAGAAGATAGGCC
apiscin	-115	CTCTCTTGTCTGTAAACCTCACCAACCAACCA-----CCCAGCTGAAATTGCCCAGCTGATGATCACCCCATCTCTTC
cratrin	-120	TTCTCTTGTCTGTAAACCTCACCAACCAACCAACCA-----CCCAGCTGAAATTGCCCAGCTGATGATCACCCCATCTCTTC
Tf-svVEGF	-119	CTCTCTTGTCTGTAAACCTCACCAACCAACCAACCA-----CCCAGCTGAAATTGCCCAGCTGATGATCACCCCATCTCTTC
Pm-VEGF	-119	CTCTCTTGTCTGTAAACCTCACCAACCAACCAACCA-----CCCAGCTGAAATTGCCCAGCTGATGATCACCCCATCTCTTC
Bi-svVEGF	-119	CTCTCTTGTCTGTAAACCTCACCAACCAACCAACCA-----CCCAGCTGAAATTGCCCAGCTGATGATCACCCCATCTCTTC
vammin	-71	CTCTCTTGTCTGTAAACCTCACCAACCAACCAACCA-----AGATCTGCCCATCTCAACCTCTGTGAGCAGCTGTGAAGCCAGGAGAAGATAGGCC
VR-1	-71	AGATCTCACCCCATCTCTCTTGTGAGCAGCTGTGAAGCCAGGAGAAGATAGGCC

Exon III

barietin	1	ATGCCTCGTACCTGCTGGCAGTGGCATCCTCTTCATCCAGGGTGGCCATCAGGGACAGTGCAAGGAGAAGTGAGGCCCCCTTATGGAAGTGTACAGGCCAGCTCTGCAGGCC
apiscin	1	ATGCCTCGTACCTGCTGGCAGTGGCATCCTCTTCATCCAGGGTGGCCATCAGGGACAGTGCAAGGAGAAGCGATGCCCTTATGGAAGTGTACAGGCCAGCTCTGCAGGCC
cratrin	1	ATGCCTCGTACCTGCTGGCAGTGGCATCCTCTTCATCCAGGGTGGCCATCAGGGACAGTGCAAGGAGAAGCGATGCCCTTATGGAAGTGTACAGGCCAGCTCTGCAGGCC
Tf-svVEGF	1	ATGCCTCGTACCTGCTGGCAGTGGCATCCTCTTCATCCAGGGTGGCCATCAGGGACAGTGCAAGGAGAAGCGATGCCCTTATGGAAGTGTACAGGCCAGCTCTGCAGGCC
Pm-VEGF	1	ATGCCTCGTACCTGCTGGCAGTGGCATCCTCTTCATCCAGGGTGGCCATCAGGGACAGTGCAAGGAGAAGCGATGCCCTTATGGAAGTGTACAGGCCAGCTCTGCAGGCC
Bi-svVEGF	1	ATGCCTCGTACCTGCTGGCAGTGGCATCCTCTTCATCCAGGGTGGCCATCAGGGACAGTGCAAGGAGAAGCGATGCCCTTATGGAAGTGTACAGGCCAGCTCTGCAGGCC
vammin	1	ATGCCTCGTACCTGCTGGCAGTGGCATCCTCTTCATCCAGGGTGGCCATCAGGGACAGTGCAAGGAGAAGCGATGCCCTTATGGAAGTGTACAGGCCAGCTCTGCAGGCC
VR-1	1	ATGCCTCGTACCTGCTGGCAGTGGCATCCTCTTCATCCAGGGTGGCCATCAGGGACAGTGCAAGGAGAAGCGATGCCCTTATGGAAGTGTACAGGCCAGCTCTGCAGGCC

Signal peptide

VHD

Exon IV

barietin	121	AGGGAGACACTCTGTCCATACTTGAAAGTAGCTCCCTGATAAAAATTCTAAGATCTTCAGGGCTCCTGTGTGCTGTGTTGCCATCGGGTGGCTGCTGCTGCTGACGAAAGCTTGACGTGCG
apiscin	121	AGGGAGACGCTACTGTCCATCTCTGATAAGGAGCACCCCTGATAAAATTCTCAGGACCTCTGTGTCACCGTGTGCAATTGCGCTGTGCGAGGAGAACGCTTCACGTGCG
cratrin	121	AGGGAGATCTACTGTCCATCTCTGATAACTTCCCACTCTGATACTGCCCCACTCTTCAGGCCCCCTCTGTGTCACCGTGTGCAATTGCGCTGTGCGAGGAGAACGCTTCACGTGCG
Tf-svVEGF	121	AGGGAGACACTGTGCCCCATCTCTGATAAGGAGCACCCCTGATAAAATTCTCAGGACCTCTGTGTCACCGTGTGCAATTGCGCTGTGCGAGGAGAACGCTTCACGTGCG
Pm-VEGF	121	AGGGAGACACTGTGCCCCATCTCTGATAAGGAGCACCCCTGATAAAATTCTCAGGACCTCTGTGTCACCGTGTGCAATTGCGCTGTGCGAGGAGAACGCTTCACGTGCG
Bi-svVEGF	121	AGGGAGACACTGTGCCCCATCTCTGATAAGGAGCACCCCTGATAAAATTCTCAGGACCTCTGTGTCACCGTGTGCAATTGCGCTGTGCGAGGAGAACGCTTCACGTGCG
vammin	121	AGGGAGACGCTCTGTGCCATACTTCAAGAGTACCTGTGATAAAATTCCGACATCTTCAGGGCCCTCTGTGTCACCGTGTGCAATTGCGCTGTGCGAGGAGAACGCTTCACGTGCG
VR-1	121	AGGGAGACGCTCTGTGCCATACTTCAAGAGTACCTGTGATAAAATTCCGACATCTTCAGGGCCCTCTGTGTCACCGTGTGCAATTGCGCTGTGCGAGGAGAACGCTTCACGTGCG

VHD (continued)

Exon V

barietin	241	ACCTCTGGAGAGCGCACTTCGAACTGCAAGGTCTGAACTCCAAAACACTGCTTCCAAAATCAAGGTGATGAAATTCAAGGGACACACGCCGTGTAATGCAAGGCCCTCGCA
apiscin	241	ACCTCTACGGAAAGCGCTCCGTCGTCGGAGATCTGCGGGTGGATCCCACAAAGGGACTCTGGCAAGATGAGGTGATGCAATTCAAGGGACACACGCCGTGTAATGCAAGGCCCTCGCA
cratrin	241	ACCCCTACGGAAAGCGCTCCGTCGTCGGAGATCTGCGGGTGGATCCCACAAAGGGACTCTGGCAAGATGAGGTGATGCAATTCAAGGGACACACGCCGTGTAATGCAAGGCCCTCGCA
Tf-svVEGF	241	ACCCCTACGGAAAGCGCTCCGTCGTCGGAGATCTGCGGGTGGATCCCACAAAGGGACTCTGGCAAGATGAGGTGATGCAATTCAAGGGACACACGCCGTGTAATGCAAGGCCCTCGCA
Pm-VEGF	241	ACCCCTACGGAAAGCGCTCCGTCGTCGGAGATCTGCGGGTGGATCCCACAAAGGGACTCTGGCAAGATGAGGTGATGCAATTCAAGGGACACACGCCGTGTAATGCAAGGCCCTCGCA
Bi-svVEGF	241	ACCCCTACGGAAAGCGCTCCGTCGTCGGAGATCTGCGGGTGGATCCCACAAAGGGACTCTGGCAAGATGAGGTGATGCAATTCAAGGGACACACGCCGTGTAATGCAAGGCCCTCGCA
vammin	241	ACCCCTACGGAAAGCGCTCCGTCGTCGGAGATCTGCGGGTGGATCCCACAAAGGGACTCTGGCAAGATGAGGTGATGCAATTCAAGGGACACACGCCGTGTAATGCAAGGCCCTCGCA
VR-1	241	ACCCCTGTGGGAAAGCACACCGTCGATAATTGCAAGATCATGCGGTGAATCCCACAGTCAGTCGCGGATGAATTCCCGACACATTCTTCGAAAATGGAGGTGATGAAATTCACTGGGACACACGCCGTGTAATGCAAGGCCCTCGCA

VHD (continued)

Exon VI

barietin	361	TCAGGAACGAGGGTGAACATCGGGAAAGCAAGA-GAAGCCCGAGAGGAAGGGGAGC-GAGAGCCAAGGTTCCCTTTGACT-----ECAGGATECCCTTTCGAAGCTTCACAG
apiscin	361	TCAGCAAGGGG-TCAACAGCGGGAAAGCGCAAGAGGA-CCCAGAGGAAGGGGAGCCGAGGCCAACTGTTCTGATCACGCTTTTGACTCCGGCTTGAAGGCTTCACAG
cratrin	361	-----ACCAAGCTGAAACACGGGAACGCCAAAGAA-CCAAAGGGAGGGGAGGCCAGAGGCCAACTGTTCTGCTGATCACGCTTTTGACTCCGGCTTGAAGGCTTCACAG
Tf-svVEGF	361	TCACCAAGGGGATGAAACACGGGAACACAAGGAA-CCCAGAGGAAGGGGAGCCGAGGCCAGGTTCCCTTTGTCGAACAGCTTTTGACTCCGGCTTGAAGGCTTCACAG
Pm-VEGF	361	TCACCAAGGGGATGAAACACGGGAACACAAGGAA-CCCAGAGGAAGGGGAGCCGAGGCCAGGTTCCCTTTGTCGAACAGCTTTTGACTCCGGCTTGAAGGCTTCACAG
Bi-svVEGF	361	TCACCAAGGGGATGAAACACGGGAACACAAGGAA-CCCAGAGGAAGGGGAGCCGAGGCCAGGTTCCCTTTGTCGAACAGCTTTTGACTCCGGCTTGAAGGCTTCACAG
vammin	361	-----AGGAAGCAGGGGTAACCTGAGGGCCGAAAGA-GAAGCCCGAGAGGAAGGGGAGT-GAGAGCCAAGGTTCCCTTTGTCGAACAGCTTTTGACTCCGGCTTGAAGGCTTCACAG
VR-1	361	-----TCGAACAGGGGTAACCTGAGGGCCGAAAGA-GAAGCCCGAGAGGAAGGGGAGT-GAGAGCCAAGGTTCCCTTTGTCGAACAGCTTTTGACTCCGGCTTGAAGGCTTCACAG

C-terminal tail (continued)

Exon VII

barietin	465	CCACACCAGGTGGGAGGCTCT-GGTCTGCAAGAC-AGATGGGGATGGGCCCTGGCTCTGATGCTGGGGTGGCTGGGAGAGGGAGGCATCTCAACCTCTGGAG
apiscin	480	CCACCGAGGTGGGAGGCTCTGGCTCAAGGCGACTGGGACGCCCTGGTCCCTGTCCTCTGATGCTGGGGTGGCTGGGAAAGGGAGGCATCTCAACATCTGGAG
cratrin	474	CCCAACAAAGATGGGAGGCTCT-GGTCTGCAAGGCGACTGGGACGCCCTGGTCCCTGTCCTCTGATGCTGGGGTGGCTGGGAAAGGGAGGCATCTCAACATCTGGAG
Tf-svVEGF	479	CCCAACAAAGGTGGGAGGCTCT-GGTCTGCAAGGCGACTGGGACGCCCTGGTCCCTGTCCTCTGATGCTGGGGTGGCTGGGAAAGGGAGGCATCTCAACATCTGGAG
Pm-VEGF	471	CCACCGAGGTGGGAGGCTCT-GGTCTGCAAGGCGACTGGGACGCCCTGGTCCCTGTCCTCTGATGCTGGGGTGGCTGGGAAAGGGAGGCATCTCAACATCTGGAG
Bi-svVEGF	480	CCACACCAGGTGGGAGGCTCT-GGTCTGCAAACTTCAAGCTGGGGAGGCCCTGGTACCTCTGTCCTCTGATGCTGGGGTGGCTGGGAAAGGGAGGCATCTCAACATCTGGAG
vammin	477	CCACACCAGGTGGGAGC-----AAAGCCGAGATGGGGAGACGCCCTGGTCCCTGTCCTCTGATGCTGGGGTGGCTGGGAAAGGGAGGCATCTCAACATCTGGAG
VR-1	474	TCACACCAGGTGGGAGC-----AAAGCCGAGATGGGGAGACGCCCTGGTCCCTGTCCTCTGATGCTGGGGTGGCTGGGAAAGGGAGGCATCTCAACATCTGGAG

C-terminal tail (continued)

Fig. S5.

barietin 583 AGGGTGTCTTGACATGTCTCACACTTACGGCAGCCGGGCGTC-----AAAAACACATC-CTCCCAGGCTGCAAGGCCAGAACTGAAGAGCGAAG
apiscin 600 AAGTTGCCATGTATCCCATCTACACTTCCATGACAGCCGGGCGTCGGCTCTTCCATGTTGACCTGAAAACACATCAGCCCAGGCTGCAAGGCCAGAACTGAAGAGCGAAG
cratrin 593 AAGTTGCTATGTATCCATCTACACTTCTATGACAGCCGGGCGTCGGCTCTTCCATGTTGACCTGAAAACACATCAGCCCAGGCTGCAAGGCCAGAACTGAAGAGCGAAG
Tf-svVEGF 598 AAGTTGCTATGTATCCATCTACACTTCTATGACAGCCGGGCGTCGGCTCTTCCATGTTGACCTGAAAACACATCAGCCCAGGCTGCAAGGCCAGAACTGAAGAGCGAAG
Pm-VEGF 590 AAGTTGCTATGTATCCATCTACACTTCTATGACAGCCGGGCGTCGGCTCTTCCATGTTGACCTGAAAACACATCAGCCCAGGCTGCAAGGCCAGAACTGAAGAGCGAAG
Bi-svVEGF 599 AAGTTGCTATGTATCCATCTACACTTCTATGACAGCTGGCGCTGCTGGCGCTCTTCCATGTTGACCTGAAAACACATCAGCCCAGGCTGCAAGGCCAGAACTGAAGAGCGAAG
vammin 583 AGGGTGTCTTGATATCCATCTAACATTCTGTGACAGCCGGGCGTC-----AAAAACACATCAGCCCAGGCTGCAAG-CCAGAGCTGAAGAGTGAAAG
VR-1 579 AGGGTGTCTTGATATCCATCTAACATTCTGTGACAGCCGGGCGTC-----AAAAACACATCAGCCCAGGCTGCAAGGCCAGAGCTGAAGAGCGAAG

barietin 673 GCAGCTTCCCTTCCAAATAACTCAGCAATCGAGTTGAATATCTGGCATCTTTTCTTACATCCGAAAGCCAGGTGCCTCTTGACCAGCTAATCATCTCCCT-----GTTTGCCTT
apiscin 720 GCAGCTTCCCTCAGTAACTCAGGAATCGAGTTGAATTTCTGGCATC-----CGAAAGCC-----TCTTGACCAGCTAACTC-----CTCA-GAGTTTGCCATT
cratrin 713 GCAGCTTCCCTCAGTAACTCAGGAATCGAGTTGAATTTCTGGCATC-----TGAAAGCC-----TCTTGACCAGCTAACTC-----CTCA-GAGTTTGCCATT
Tf-svVEGF 718 GCAGCTTCCCTCAGTAACTCAGCAATCGAGTTGAATTTCTGGCATC-----CGAAAGCC-----TCTTGACCAGCTAACTCAGTCTCCCTCAAGAGTTTGCCATT
Pm-VEGF 710 GCAGCTTCCCTCAGTAACTCAGCAATCGAGTTGAATTTCTGGCATC-----CGAAAGCC-----TCTTGACCAGCTAACTCAGTCTCCCTCAAGAGTTTGCCATT
Bi-svVEGF 719 GCAGCTTCCCTCAGTAACTCAGGAATCGAGTTGAATTTCTGGCATC-----CGAAAGCC-----TCTTGACCAGCTAACTC-----CTCA-GAGTTTGCCATT
vammin 673 GCATCTTCCCTTACCAAATAACTCAGCAATCGAGTTGAATATCTGGCTCTTTTCTTACATCCGAAAGCCAGGTGCCTCTGACCAACTAATCATCTCCCT-----GTTTGCCTT
VR-1 670 GCAGCTTCCCTTCCAAATAACTCGCGATCGAGTTGAATATCTGGCATCTTTTCTTACATCCGAAAGCCAGGTGCCTCTGACCAACTAATCATCTCCCTCA-----GTCAGCCATT

barietin 788 TTGTGTCAAGCTTTAGGCTTTAGTAAATTG-----AATACACAATTAAAATAAGATTGATATCTCCGACCCCTCATCAGGGGAGCTGGGGAAAAACATTGTTTCTT
apiscin 812 TTGTGTCAAGT-CTTAATGCTTTAGTAAATTGCTGTAAATCACATTAAAATAAGATTGATATCTCCCAACCCCTCATCAGGGGAGCTGGGGAAAAACATTAGTTTCTT
cratrin 805 TTGTGTCAAGC-CTTAATGGCTTTAGTAAATTGCTGTAAATCACATTAAAATAAGATTGATATCTCCGACCCCTCATCAGGGGAGCTGGGGAAAAACATTAGTTTCTT
Tf-svVEGF 817 TTGTGTCAAGC-CTTAATGGCTTTAGTAAATTGCTGTAAATCACATTAAAATAAGATTGATATCTCCGACCCCTCATCAGGGGAGCTGGGGAAAAACAGTTTCTT-----
Pm-VEGF 809 TTGTGTCAAGC-CTTAATGGCTTTAGTAAATTGCTGTAAATCACATTAAAATAAGATTGATATCTCCGACCCCTCATCAGGGGAGCTGGGGAAAAACAGTTTCTT-----
Bi-svVEGF 811 TTGTGTCAAGC-CTTAATGGCTTTAGTAAATTGCTGTAAATCACATTAAAATAAGATTGATATCTCCGACCCCTCATCAGGGGAGCTGGGGAAAAACATTAGTTTCTT
vammin 788 TTGTGTCAAGC-CTTAATGGCTTTAGTAAATTGCTGTAAATCACATTAAAATAAGATTGATATCTCCGACCCCTCATCAGGGGAGCTGGGGAAAAACATTAGTTTCTT
VR-1 787 TTGTGTCAAGC-CTTAATGGCTTTAGTAAATTGCTGTAAATCACATTAAAATAAGATTGATATCTCCGACCCCTCATCAGGGGAGCTGGGGAAAAACATTAGTTTCTT

barietin 902 TTAAAGAATGTGGAATTCTGTCTTTAGCCC-ACCTG-----CATGAGTTCACGGAGCCCTGGAAATAA-----CGGAGTAAATT-polyA
apiscin 931 TTAAAGAATGTGGAATTATGTCTTTAGCCCCGCTGGATCTGCATGACTTACAGAGCCCTGGAAATAA-----CGGAGTAAACTAAAAAGC-polyA
cratrin 924 TTAAAGAATGTGGAATTATGTCTTTAGCCCCGCTGGATCTGCATGACTTCA-GAGCCTTGGAAATAA-----CGGAGTAAACC-polyA
Tf-svVEGF 933 -TAAAGAATGTGGAATTATGTCTTTAGCCCCGCTGGATCTGCATGACTTACAGAGCCCTGGAAATAA-----CGGAGTAAACT-polyA
Pm-VEGF 926 -TAAAGAATGTGGAATTATGTCTTTAGCCCCGCTGGATCTGCATGACTTACAGAGCCCTGGAA
Bi-svVEGF 930 TTAAAGAATGTGGAATTATGTCTTTAGCCCCAATCTG-----CATGACTTACAGAGCCCTGGAAATAAATAACGGAGTAAACT-polyA
vammin 907 TTAAAGAATGTGGAATTATGTCTTTAGCCCCAATCTG-----CATGACTTACAGAGCCCTGGAAATAA-----GGAGTAAACC-polyA
VR-1 906 TTAAAGAATGCCGAATTATGTCTTTAGCCCCAATCTG-----CATGACTTACAGAGCCCTGGAAATAA-----GGAGTAAACTC-polyA

Fig. S5. (continued)