SUPPLEMENTARY FIGURES

Supplementary Figure 1

(A) DNA-sequence of mini-gp130-ELP: normal print: cDNA coding for the LeB4-signal peptide, bold: cDNA coding for mini-gp130-ELP; Primer and restriction sites are indicated. The ATG start codon is underlined.

(B) Protein sequence of mini-gp130-ELP. Grey: LeB4-signal peptide; red: sgp130 domains 1-3; red, cursive: amino acid sequence determined by Edman-degradation of the mature purified mini-gp130-ELP; blue: c-myc-tag; green: 100 repeats of ELP; brown: ER-retention-signal.

Supplementary Figure 2

Protein stability of purified mini-gp130-ELP at 37°C and 4°C.

(A) 1 μg purified mini-gp130-ELP was diluted in 1 ml PBS and incubated for 48 h at 37°C. Aliquots of originally 10 ng mini-gp130-ELP were separated by SDS/PAGE and blotted onto a PVDF membrane. Proteins were detected with the c-myc specific antibody 9E10 and visualized by ECL detection. Lane 1+2: 0 h; lane 3+4: 12 h; lane 5+6: 24 h; Lane 7+8: 48 h.
(B) 1 μg purified mini-gp130-ELP was diluted in 1 ml PBS and incubated for 26 d at 4°C. Aliquots of originally 10 ng mini-gp130-ELP were separated by SDS/PAGE and blotted onto a PVDF membrane. Proteins were detected with a c-myc specific antibody and visualized by ECL detection. Lane 1: 0 d; lane 2: 7 d; lane 3: 19 d; Lane 4: 26 d.

Supplementary Figure 3

Expression of sgp130-variants in transgenic tobacco.

Leaves of transgenic lines were extracted in a mortar under liquid nitrogen in 50 mM Tris-HCl, 200 mM NaCl, 5 mM EDTA, 0.1% Tween 20, pH 8.0. The homogenate was centrifuged for 5 min at 4°C and 16,000 g. Proteins were separated on a 10% SDS polyacrylamide gel, blotted and c-myc-tag containing proteins were detected by Western blotting and ECL.

(A) Lane 1: 40 µg extract of sgp130Fc-ELP transgenic tobacco plants; lane 2: 40 µg extract of sgp130Fc transgenic tobacco plants; lane 3: 40 ng c-myc-tagged scFv control protein. M: molecular mass marker (kDa).

(**B**) Lane 1: 5 µg extract of mini-gp130-ELP transgenic tobacco plants; lane 2: 5 ng control protein; lane 3: 10 ng c-myc-tagged single chain Fv control protein; lane 4: 5 µg extract of mini-gp130-ELP transgenic tobacco plants. M: molecular mass marker (kDa).

Supplementary Figure 4 Calibration of size exclusion chromatography.

(A) For calibration the high molecular mass standard from Amersham Pharmacia Biotech was used. Peak 1: Thyroglobin 669 kD; Peak 2: Ferritin 490 kDa; Peak 3: Katalase 232 kDa; Peak 4: Aldolase 158 kDa.

(B) The known molecular masses of the purchased standard proteins and their respective elution volumes were subject to linear regression.

Supplementary Figure 5 Absorption spectra of pure mini-gp130-ELP

(A) The absorption spectrum of purified mini-gp130-ELP was recorded in the range of 240-320 nm.

(B) The protein concentration of purified mini-gp130-ELP (after inverse transition cycling and size exclusion chromatography) was calculated as indicated from the absorption at 280 nm.

1 ATGGCTTCCA AACCTTTTCT ATCTTTGCTT TCACTTTCCT TGCTTCTCTT

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Primer: 5´gp130-∆signal

	A	<i>f</i> /111			
51	TACAAGCACA	TGTTTAGCAG	AGCTGCTGGA	TCCTTGCGGC	TATATCTCCC
101	CTGAGTCTCC	TGTGGTGCAG	CTGCATTCTA	ACTTCACCGC	CGTGTGTGTG
151	CTGAAGGAAA	AGTGCATGGA	CTACTTCCAC	GTGAACGCCA	ACTACATCGT
201	GTGGAAAACC	AACCACTTCA	CCATCCCCAA	GGAGCAGTAC	ACCATCATCA
251	ACCGGACCGC	TTCTTCTGTG	ACCTTCACCG	ATATCGCCTC	CCTGAATATC
301	CAGCTGACCT	GCAACATCCT	GACCTTTGGA	CAGCTGGAGC	AGAATGTGTA
351	CGGCATCACC	ATCATCTCTG	GCCTGCCTCC	AGAGAAGCCT	AAGAACCTGT
401	CCTGCATCGT	GAATGAGGGC	AAGAAGATGA	GGTGTGAGTG	GGATGGCGGC
451	AGAGAGACAC	ATCTGGAGAC	CAACTTCACC	CTGAAGTCTG	AGTGGGCCAC
501	CCACAAGTTT	GCCGACTGCA	AGGCCAAGAG	AGATACCCCT	ACCTCTTGCA
551	CCGTGGACTA	CTCCACCGTG	TACTTCGTGA	ACATCGAGGT	GTGGGTGGAG
601	GCTGAGAATG	CTCTGGGCAA	GGTGACCTCT	GACCACATCA	ACTTCGACCC
651	CGTGTACAAG	GTGAAGCCTA	ACCCTCCTCA	CAACCTGTCC	GTGATCAACT
701	CTGAGGAGCT	GTCCTCTATC	CTGAAGCTGA	CCTGGACCAA	CCCTTCCATC
751	AAGTCCGTGA	TCATCCTGAA	GTACAACATC	CAGTACAGGA	CCAAGGATGC
801	TTCTACCTGG	TCTCAGATCC	CTCCTGAGGA	TACCGCTTCC	ACCAGATCCA
851	GCTTCACAGT	GCAGGACCTG	AAGCCTTTTA	CCGAGTACGT	GTTCAGGATC
901	CGGTGCATGA	AGGAGGATGG	CAAGGGCTAT	TGGTCTGACT	GGTCTGAGGA

Primer: 3´gp130-Nael

Nael

►

951	GGCTTCTGGC	ATCACCTACG	AGGACAGAGC	CGGCGGACAA	GCGGCCGCAG
1001	AACAAAAACT	CATCTCAGAA	GAGGATCTGA	ATGGGGCCGT	CGAGATGGGC
1051	CACGGCGTGG	GTGTTCCGGG	CGTGGGTGTT	CCGGGTGGCG	GTGTGCCGGG
1101	CGCAGGTGTT	CCTGGTGTAG	GTGTGCCGGG	TGTTGGTGTG	CCGGGTGTTG
1151	GTGTACCAGG	TGGCGGTGTT	CCGGGTGCAG	GCGTTCCGGG	TGGCGGTGTG
1201	CCGGGCGTGG	GTGTTCCGGG	CGTGGGTGTT	CCGGGTGGCG	GTGTGCCGGG
1251	CGCAGGTGTT	CCTGGTGTAG	GTGTGCCGGG	TGTTGGTGTG	CCGGGTGTTG
1301	GTGTACCAGG	TGGCGGTGTT	CCGGGTGCAG	GCGTTCCGGG	TGGCGGTGTG
1351	CCGGGCGTGG	GTGTTCCGGG	CGTGGGTGTT	CCGGGTGGCG	GTGTGCCGGG
1401	CGCAGGTGTT	CCTGGTGTAG	GTGTGCCGGG	TGTTGGTGTG	CCGGGTGTTG
1451	GTGTACCAGG	TGGCGGTGTT	CCGGGTGCAG	GCGTTCCGGG	TGGCGGTGTG
1501	CCGGGCGTGG	GTGTTCCGGG	CGTGGGTGTT	CCGGGTGGCG	GTGTGCCGGG
1551	CGCAGGTGTT	CCTGGTGTAG	GTGTGCCGGG	TGTTGGTGTG	CCGGGTGTTG
1601	GTGTACCAGG	TGGCGGTGTT	CCGGGTGCAG	GCGTTCCGGG	TGGCGGTGTG
1651	CCGGGCGTGG	GTGTTCCGGG	CGTGGGTGTT	CCGGGTGGCG	GTGTGCCGGG
1701	CGCAGGTGTT	CCTGGTGTAG	GTGTGCCGGG	TGTTGGTGTG	CCGGGTGTTG
1751	GTGTACCAGG	TGGCGGTGTT	CCGGGTGCAG	GCGTTCCGGG	TGGCGGTGTG
1801	CCGGGCGTGG	GTGTTCCGGG	CGTGGGTGTT	CCGGGTGGCG	GTGTGCCGGG
1851	CGCAGGTGTT	CCTGGTGTAG	GTGTGCCGGG	TGTTGGTGTG	CCGGGTGTTG
1901	GTGTACCAGG	TGGCGGTGTT	CCGGGTGCAG	GCGTTCCGGG	TGGCGGTGTG
1951	CCGGGCGTGG	GTGTTCCGGG	CGTGGGTGTT	CCGGGTGGCG	GTGTGCCGGG
2001	CGCAGGTGTT	CCTGGTGTAG	GTGTGCCGGG	TGTTGGTGTG	CCGGGTGTTG
2051	GTGTACCAGG	TGGCGGTGTT	CCGGGTGCAG	GCGTTCCGGG	TGGCGGTGTG
2101	CCGGGCGTGG	GTGTTCCGGG	CGTGGGTGTT	CCGGGTGGCG	GTGTGCCGGG
2151	CGCAGGTGTT	CCTGGTGTAG	GTGTGCCGGG	TGTTGGTGTG	CCGGGTGTTG
2201	GTGTACCAGG	TGGCGGTGTT	CCGGGTGCAG	GCGTTCCGGG	TGGCGGTGTG
2251	CCGGGCGTGG	GTGTTCCGGG	CGTGGGTGTT	CCGGGTGGCG	GTGTGCCGGG
2301	CGCAGGTGTT	CCTGGTGTAG	GTGTGCCGGG	TGTTGGTGTG	CCGGGTGTTG
2351	GTGTACCAGG	TGGCGGTGTT	CCGGGTGCAG	GCGTTCCGGG	TGGCGGTGTG
2401	CCGGGCGTGG	GTGTTCCGGG	CGTGGGTGTT	CCGGGTGGCG	GTGTGCCGGG
2451	CGCAGGTGTT	CCTGGTGTAG	GTGTGCCGGG	TGTTGGTGTG	CCGGGTGTTG
2501	GTGTACCAGG	TGGCGGTGTT	CCGGGTGCAG	GCGTTCCGGG	TGGCGGTGTG
2551	CCGGGCGGGC	TGGCGGCCGC	AGAACCCAAA	GACGAACTCT	AG

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cleavage site of the signal peptide

LeB4-signal peptide sgp130-ELP MASKPFLSLLSLSLLLFTSTCLAELLDPCGYISPESPVVQLHSNFTAVCV 50 LKEKCMDYFHVNANYIVWKTNHFTIPKEQYTIINRTASSVTFTDIASLNI 100 QLTCNILTFGQLEQNVYGITIISGLPPEKPKNLSCIVNEGKKMRCEWDGG 150 RETHLETNFTLKSEWATHKFADCKAKRDTPTSCTVDYSTVYFVNIEVWVE 200 AENALGKVTSDHINFDPVYKVKPNPPHNLSVINSEELSSILKLTWTNPSI 250 KSVIILKYNIQYRTKDASTWSQIPPEDTASTRSSFTVQDLKPFTEYVFRI 300 RCMKEDGKGYWSDWSEEASGITYEDRAGGQAAAEQKLISEEDLNGAVEMG 350 HGVGVPGVGVPGGGVPGAGVPGVGVGVGVGVGVGVGGGVPGAGVPGGGV 400 PGVGVPGVGVPGGGVPGAGVPGVGVPGVGVPGVGVPGGGVPGAGVPGGGV 450 PGVGVPGVGVPGGGVPGAGVPGVGVPGVGVPGVGVPGGGVPGAGVPGGGV 500 PGVGVPGVGVPGGGVPGAGVPGVGVGVGVGVGVGVGGGVPGAGVPGGGV 550 PGVGVPGVGVPGGGVPGAGVPGVGVPGVGVPGVGVPGGGVPGAGVPGGGV 600 PGVGVPGVGVPGGGVPGAGVPGVGVPGVGVPGVGVPGGGVPGAGVPGGGV 650 PGVGVPGVGVPGGGVPGAGVPGVGVPGVGVPGVGVPGGGVPGAGVPGGGV 700 PGVGVPGVGVPGGGVPGAGVPGVGVPGVGVPGVGVPGGGVPGAGVPGGGV 750 PGVGVPGVGVPGGGVPGAGVPGVGVPGVGVPGVGVPGGGVPGAGVPGGGV 800 PGVGVPGVGVPGGGVPGAGVPGVGVGVGVGVGVGVGGGVPGAGVPGGGV 850 PGGLAAAEPKDEL*

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