

The LIM domain protein BmFHL2 inhibits egg production in female silkworm, *Bombyx mori*

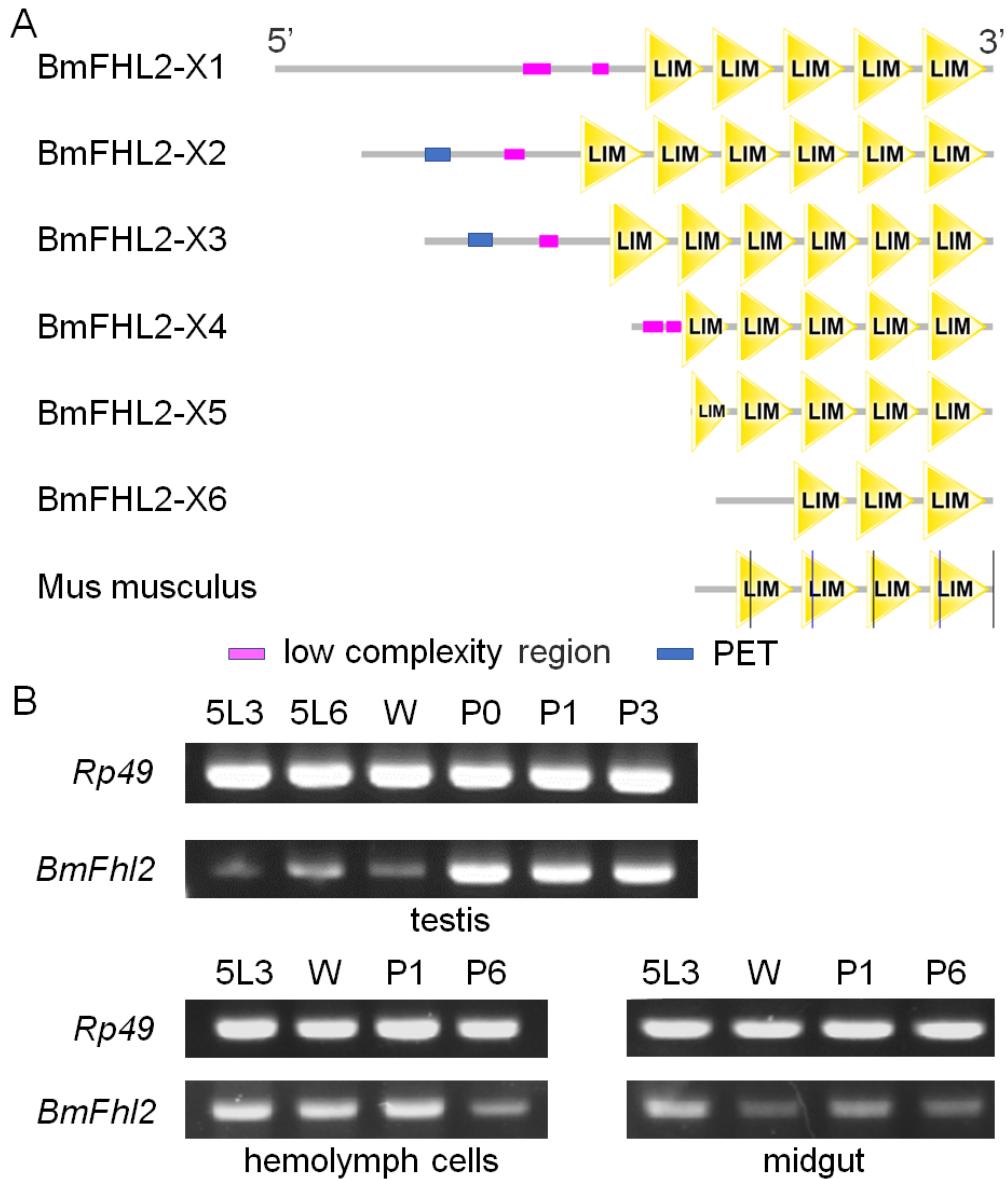


Figure S1. Schematic diagram of the domain analysis of BmFHL2. (A) Schematic diagram of the structure of different shears of silkworm FHL2 and mouse FHL2. (B) *BmFhl2* expression in testis, hemolymph cells and midgut at different time points. Total RNA was extracted from testis, hemolymph cells and midgut at different time points for RT-PCR analysis. *Rp49* was used as a reference gene. All samples were collected and mixed for three silkworm populations. 5L3: 5th instar 3 days, 5L6: 5th instar 6 days, W: wandering stage, P0: pupal stage 0 days, P1: pupal stage 1 day, P3: pupal stage 3 days, P6: pupal stage 6 days.

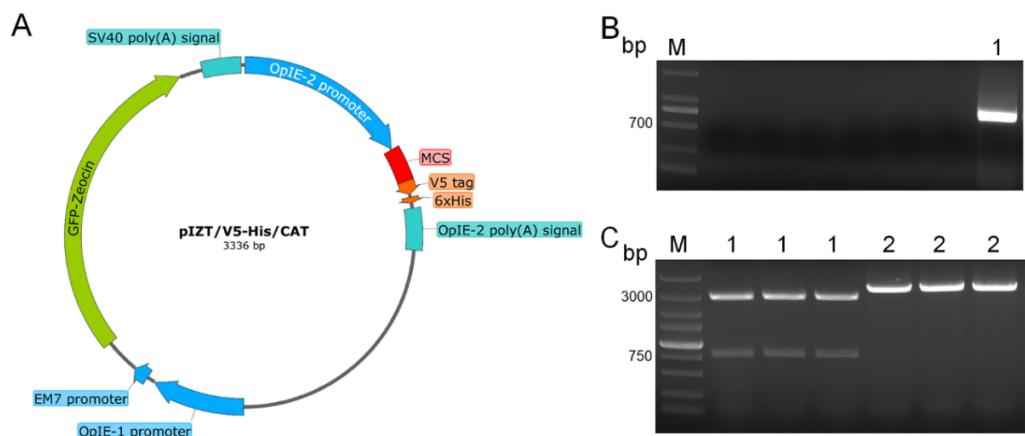


Figure S2. Overexpression of *BmFhl2* in BmN cells. (A) Schematic diagram of the *BmFhl2.A.pIZT/V5*-His plasmid overexpressed in BmN cells. (B) *BmFhl2-X6* was amplified from the cDNA of *B. mori* ovary at 5L3, and the product was subjected to agarose gel electrophoresis. M: Marker; 1: Target fragment. (C) Schematic diagram of electrophoresis of enzyme digestion products. M: Marker; 1: Not I and Kpn I enzyme digestion products of T vector; 2: Not I and Kpn I enzyme digestion products of empty vector.

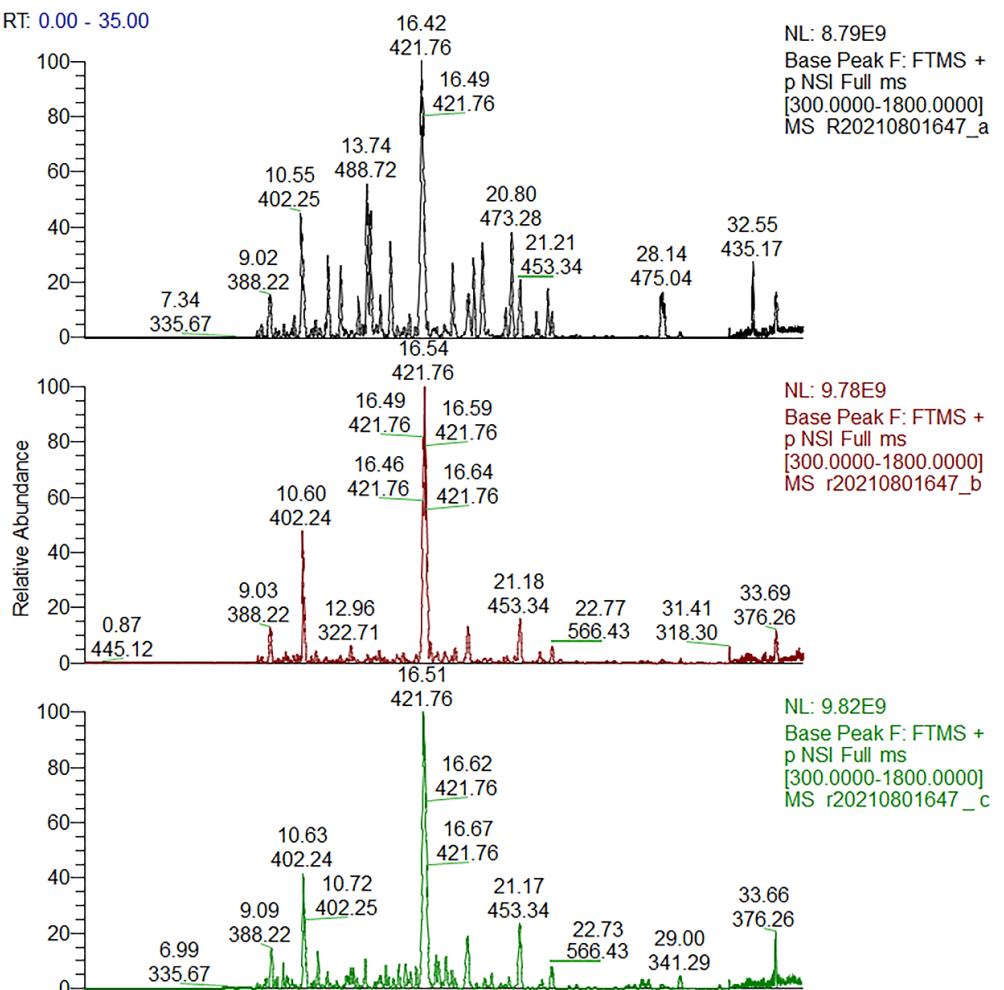


Figure S3. Co-IP mass spectrometry identification.

Table S1. Identification of interacting proteins of BmFHL2 by mass spectrometry.

NO.	Protein name	MW (kDa)	PepCount/ Unique Pepcount	Cover percent	sequence
a-1	26S proteasome non-ATPase regulatory subunit 11	47.2	15/15	34.92%	TYKPELEEDAVVR
a-2	proliferation-associated protein 2G4	42.1	8/8	21.9%	VLEQVIAK
a-3	26S proteasome non-ATPase regulatory subunit 13	43	7/7	20.52%	YVG CADGEALP ATER
a-4	probable citrate synthase 2	51.8	8/8	18.49%	VVP PILTELGK
a-5	FAD-dependent oxidoreductase domain-containing protein 1	43.5	6/6	15.35%	IIDTENN VINAK
a-6	BTB/POZ domain-containing protein At1g21780	41.9	3/3	7.48%	EETIESSQK
b-1	glutamate dehydrogenase	61.4	25/25	43.86%	IIAEAANGPTTPAA DK
b-2	ATP synthase subunit beta, mitochondrial	54.8	15/15	40.94%	FTQAGSEVSALLG R
b-3	low molecular lipoprotein 30K pBmHPC-12	29.2	3/2	14.84%	VVYGGNSADSTR
b-4	low molecular lipoprotein 30K pBmHPC-21	30.2	2/1	10.27%	VIFGTNTADTTR
b-5	low molecular lipoprotein 30K pBmHPC-19	27.4	2/2	6.33%	LIFAENAIK
c-1	polyubiquitin	17.2	3/3	39.22%	TITLEVEPSDTIENV K
c-2	ecdysteroid-phosphate phosphatase	161.1	2/2	1.4%	AELPLAALQR

Table S2. The sequence of primers.

Gene name	Primer name	Sequence (5'-3')
BmRp49	Rp49-q-F Rp49-q-R	CAGGC GGTCAAGGGTCAATAC TACGGAATCCATTGGGAGCAT

<i>BmRp49</i>	Rp49-RT-F Rp49-RT-R	GACCTGTTACAGGCCGACA ACTCTGATGCTGAGCTGCTG
<i>BmFHL2</i>	FHL2-q-F FHL2-q-R	TTCTTGACCAAGCCCATCA CCTGTCCGTCGGTGATGAAG
<i>BmFHL2</i>	FHL2-RT-F FHL2-RT-R	TTCCGTGCTGGAACCAAGAA CGCAGATGAAGCAGTCGTTG
<i>Vg</i>	Vg-F Vg-R	CTTGTGCCATCGATAGAACAG GTCGATATTGCATCCCCATC
<i>30Kc19</i>	30Kc19-F 30Kc19-R	GCGGTTGAAAAGAGCAAGCA ACTGGGAAACAATCCGGAC
<i>VgR</i>	VgR-F VgR-R	GAGTGCCTGGCGAGGATGT CTGAGCGTCTGGCTTGTGA
<i>EcR</i>	EcR-F EcR-R	GTGCTGGATGTGGTTGCTGAGGT CCGCTGGTCTAATAACGGTGGCT
<i>USP</i>	USP-F USP-R	ATGTCACCGCCCCGAAATGAAGCC CTGCTGGATGAGCGTCTTCTGT
<i>E75a</i>	E75a-F E75a-R	TCGGTCGAGCTTGAGTGAGA TAGCACAGGAGTGCATGAGC
<i>siBmFHL2-405</i>	F R	CCGGCUGCUACGAGGACAATT UUGUCCUCGUAGCAGCCGGTT
<i>siBmFHL2-777</i>	F R	GCGCCAAGCAGAAGCUAUTT AUGAGCUUCUGCUUGGCGCTT
<i>si NC</i>		UUCUCCGAACGUGUCACGUTT
<i>BmFHL2-X6</i>	F R	GGTACCATGTCAGCGGACGTCTTGAG (Kpn I) GCGGCCGCTCATGAGCTTGCTGGC (Not I)