

**Molecular cloning and functional characterization of an H⁺-pyrophosphatase
from *Iris lactea***

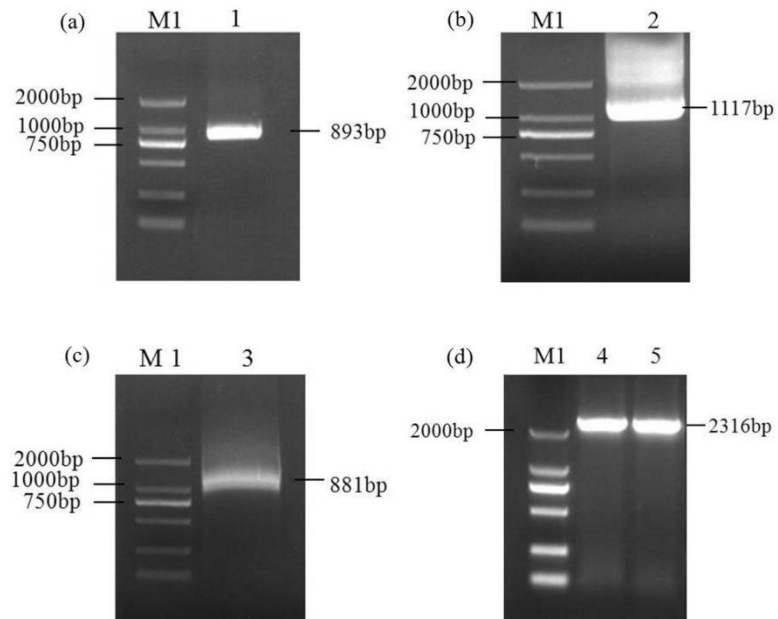
Authors: Lin Meng*, Shanshan Li, Jingya Guo, Qiang Guo, Peichun Mao, Xiaoxia Tian

Institutional addresses: Beijing Research and Development Centre for Grass and Environment, Beijing Academy of Agriculture and Forestry Sciences, Beijing 100097, P. R. China

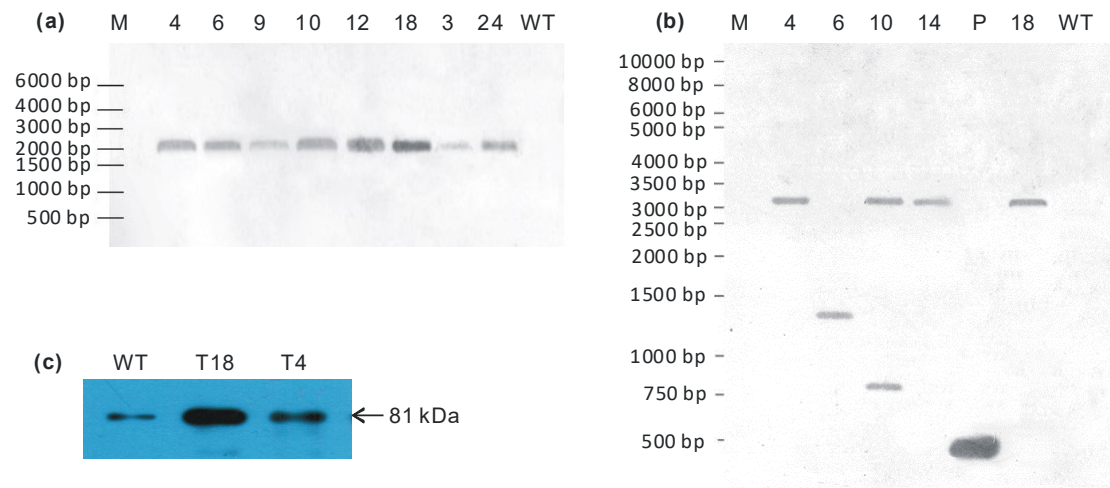
***Corresponding author:** Lin Meng, E-mail: menglin9599@sina.com

Tel: +86-10-51503345

Fax: +86-10-51503297



Supplementary Figure 1 Electrophoresis of *IIVP* cDNA clones. (a) 1: RT-PCR products of the *IIVP* fragment; (b) 2: 3' RACE product of the *IIVP* gene; (c) 3: 5' RACE product of the *IIVP* gene; (d) 4, 5: PCR product of the full-length *IIVP* gene.



Supplementary Figure 2 **a.** Northern blot analysis of *IlVP*-transgenic tobacco plants. Lines 4, 6, 9, 10, 12, 18, 3, and 24 and WT correspond to individual transgenic tobacco lines and the wild type, respectively. **b.** Southern blot analysis of *IlVP*-transgenic tobacco plants. Lines P (positive control) 4, 6, 10, 14, 18 and WT correspond to individual transgenic tobacco lines and the wild type, respectively. **c.** Western blot analysis of *IlVP* protein in WT and transgenic tobacco lines (T4 and T18) under 200 mM NaCl treatment for 10 d, respectively.

Supplementary Table 1 Primer sequences used in the experiments

Primer	Sequence (5'-3')
P1	TATGGTGATGAYTGGGAAGG
P2	GCAATRCCWCCAGCATTRTCA
P3	CAACATCAGCAGCTTTAGTGTAGATA
P4	TATGGCCCCATCAGTGACAATGCT
P5	TCCCCCGGGATGGTGGCGGCGATGCT
P6	<u>CGAGCTCT</u> TAGAAGATCTTGAAGAGGATGCC
P7	CGAACTGACTGCTATGATGTACCC
P8	CCAATAACAACCGCAATACCA
A1	TATTGTGCTGGATTCTGGTGATG
A2	GGAGGATAGCATGGGGAAGAG