

Stable and efficient expression of brain-derived neurotrophic factor in tobacco chloroplasts

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Stable and efficient expression of brain-derived neurotrophic factor in

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- Abstract: Brain-derived neurotrophic factor (BDNF) is a factor with many functions such as 20 acceleration of cell proliferation and differentiation and is therefore widely used in clinical 21 applications. In this study, an expression vector named pWYP23402 having a codon-optimized 22 BDNF gene was constructed and transferred into chloroplasts of tobacco by gene-gun. After 23 three or four rounds of selection with proper spectinomycin, BDNF was integrated into the 24 25 chloroplast genome of homoplastomic plants confirmed by PCR and Southern hybridization. ELISA assay indicated that *BDNF* fused with GFP represented approximately $15.72\% \pm 0.33\%$ 26 of total soluble protein in the leaves of transplastomic plants. Moreover, the chloroplast-derived 27 BDNF displayed similar biological activity to the commercial product. This is the first case 28 report of BDNF expression by chloroplast transformation in this model plant, supplying an 29 additional pathway for the production of chloroplast-expressed therapeutic proteins. 30
- 31 Key words Tobacco, Chloroplast, Genetic transformation, Brain-derived neurotrophic factor,
- 32 Green fluorescent protein

33 Abbreviations

34	BDNF	Brain-derived neurotrophic factor
35	ELISA	Enzyme-linked immunosorbent assay
36 37	GFP	Green fluorescent protein
38	LTP	Long-term potentiation
39	MCS	Multiple cloning site
40	NGF	Nerve growth factor
41	PBS	Phosphate-buffered saline
42	PMP	Plant-made pharmaceuticals
43	SCI	Spinal cord injury

SDS-PAGE Sodium dodecyl sulfate-polyacrylamide gel electrophoresis

45 TrkB Tropomycin receptor kinase B

Introduction

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BDNF (brain-derived neurotrophic factor) is a growth factor studied intensively and belongs to the member of the neurotrophin family of secret proteins that involving in neurotrophin-3 (NT3), NGF, and neutorophin-4 (NT4) [1-5]. It was the first neurotrophic factor explored and purified from pig brain in the year of 1982 since the discovery of NGF. This NGF-like substance was found to be expressed at an extremely low level in the brain [6]. The human BDNF gene is composed of 11 exons [7] and its full primary structure showed highly conservative, with 85.9 to 100 % identity among the genes of various vertebrates and humans as well [8]. BDNF is first synthesized into pre-pro-BDNF in cell bodies of neurons and glia and then cleaved to become mature BDNF. The mature BDNF will be transported to the terminals where it is released 72 [9]. However, BDNF can also be delivered to the dendrites where the activity-dependent local translation of BDNF takes place [10]. The factor binding with high affinity to the tropomycin receptor kinase B (TrkB) receptor can activate at least three intracellular signaling pathways 75 downstream [11, 12]. 76 BDNF gene was located in No. 11q13 chromosome of the human genome and No. 12 chromosome 77 of rat genome and shared the 50 to 60 percentage of similarity of sequence over other members of the 78 79 neurotrophic factor family. It is one of the most widely distributed neurotrophic factors in the brain and is mainly expressed in the central nervous system, especially with the highest levels in the 80 regions of the hippocampus and cortex. The mRNA for BDNF was found predominantly in the central nervous system and its sequence of protein indicated that BDNF and NGF are related closely both in structure and function [13]. As a multi-functional chemical molecule, BDNF has been proved to play a critical role in synaptic plasticity, LTP, learning, and memory [14-17, 13].

The more growing evidence demonstrated that BDNF was involved in a wide range of neurophysiological processes [18]. For instance, over the last decade, BDNF has been consistently highlighted as a key linker between the antidepressant drug and the neuroplastic changes, resulting in the improvement of depressive symptoms [19]. It was reported that local hydrogel delivery of BDNF with controlled dosing and duration to the injured cervical spinal cord can repair the respiratory neural circuitry, and this delivery was a safe and robustly-effective strategy to restore diaphragm function after SCI [20].

To be the ideal platform for protein expression, chloroplasts have been regarded to be the favorite organelle for the production of therapeutic proteins including growth factors like bFGF, EGF, and antimicrobial peptides like MSI-99 [21-23].

Tobacco was the firstly engineered flowering plant via chloroplast transformation more than three decades ago [24], and still served as the model plant due to its easy plastom genetic manipulation. Compared with existing prokaryotic and eukaryotic expression systems which have technical and economic limitations, plants have created substantial interest in developing new expression platforms for the effective production of much safer and biologically active complex therapeutic proteins or antigens. More importantly, chloroplast transformation has several advantages like higher expression, generally soluble proteins, and mostly maternal inheritance over nuclear transformation. No sterile laboratories or related equipment is required for protein production as well [25-27].

In this experiment, we engineered tobacco plants to firstly express BDNF via chloroplast transformation. These results will offer solid support for potentially producing BDNF production via growing plants on a large scale to meet the growing clinical demand for therapeutic proteins.

Materials and Methods

Plant materials

Tobacco (*Nicotiana tabacum* cv. Petit Havana) seeds were surface-sterilized and grown on a half-strength MS medium supplemented with 8 g/L agar and 30 g/L sucrose under a 16/8 h light/dark (L/D) cycles. Fully-expanded young leaves were used for subsequent chloroplast DNA extraction and genetic transformation.

Vector construction

- According to the nucleotide sequence in the NCBI database, the *BDNF* gene was optimized under the precondition of keeping the amino acid sequence unchanged. The codons were optimized and replaced by Gene Designer based on the codon bias of the tobacco chloroplast genome. The modified gene sequence was analyzed by DNAMAN7.0 and Vector NTI 10.0 to find out enzyme site information. Meanwhile, the optimal codon ATT for serine amino acid was replaced by ATA to eliminate *Xba* I enzyme sites. Two enzyme sites, *Cla* I and *Xba* I, were flanked by the two ends of the optimized gene to fuse with *the gpf* gene.
- The expression cassette was chemically synthesized in which a linker amino acid sequence (GSSSGSSSGSSSGSSSDDDDK) combined reporter gene *gfp* with target gene *BDNF* and selectable gene *aadA*, all three cluster genes were driven by promoter *Prrn-T7g10* and terminator *Trps16*.
- This artificial cassette was confirmed by sequencing and inserted into the site between *trn I* gene and *trn A* gene to form the expression vector namely pWYP23402.
 - Tobacco Chloroplast transformation and homoplastomic selection
 - The tobacco chloroplast transformation was carried out according to the protocol previously described [21]. Young leaves were placed on the MS medium with the adaxial side down, cultured in the dark for 4 h, and then bombarded by gene-gun. The bombarded leaves were cultured at 26°C for 3 days in the dark condition before being cut into 5 mm × 5 mm pieces and transferred to a selective medium that was the MS medium containing 500 mg/L spectinomycin, 2 mg/L 6-BA, 0.1 mg/L NAA,

30 g/L sucrose, and 8 g/L agar at the first round of selection. The medium was renewed every three weeks until the appearance of resistant transformants. During the selection, a UV lamp was used to detect the intensity of GFP expression in the leave cells of transformants, and the plants with high intensity were picked up to do PCR analysis. The PCR-confirmed leaves of resistant transformants were cut into pieces and subjected to additional cycles of selection until the homoplastomic plants were achieved.

Molecular analysis

(1) PCR analysis

For the preliminary identification of homoplastomic plants, total DNA was extracted from young leaves of both transplastomic and wild-type tobacco plants to carry out PCR analysis. The P1 and P2 marked in Fig.1 were designed to check the expression cassette.

(2) Southern blot

Integration of the foreign expression cassette into the tobacco plastid genome was characterized by Southern blot hybridization. The total plastome DNA extracted from young leaves of the transgenic plant was digested with two endonucleases *Bam*H I and *Kpn* I (New England Biolabs, UK) at 37°C overnight and subjected to 0.8% agarose gel electrophoresis, then the digested DNA in gel was transferred onto positive-charged Hybond-N⁺ nylon membrane (Roche, Switzerland). DNA was fixed by UV-cross-linking twice for 30 s each time at 2 min intervals. Pre-hybridization and hybridization were carried out by using the DIG High Prime DNA Labeling and Detection Starter Kit II (Roche, Switzerland). The DNA-fixed membrane was hybridized with a ~1.4kb probe in an oven for 16 h at 42°C before washing and visualization. The untransformed wild-type plant was used as the negative control.

Western blot and ELISA assays

To measure the expression level of BDNF, Western blot analysis, and ELISA assay were performed. Briefly, the leaves were grounded by liquid nitrogen, homogenized in PBS buffer (137 mM NaCl, 2.7 mM KCl, 10 mM Na₂HPO₄, 2 mM KH₂PO₄) to extract the crude total soluble protein from leaves of PCR-positive plants, separated by a 15% SDS-PAGE gel, then transferred to a nitrocellulose membrane (0.22 μm, Millipore, USA) via a semi-dry trans-blot electrophoretic transfer cell (200 mA, 30 min, Bio-Rad, USA). The nitrocellulose membrane was then saturated with 5% BSA (w/v) in TBS-Tween 20 (25 mM Tris-HCl pH 7.6, 0.15 M NaCl, 0.05% (w/v) Tween-20) for 1 h. The membrane was incubated with a BDNF-directed antibody (1:100, Santa Cruz Biotechnology Inc. USA), before incubation with the secondary antibody (1:3000, goat anti-mouse IgG alkaline phosphatase conjugate, Novagen, Germany). Labeled proteins were visualized by reaction with BCIP/NBT as a substrate (Roche, Switzerland).

According to the Kit instruction (Abcam, Cambridge, MA, USA), ELISA was performed using commercial anti-BDNF antibodies and commercial BDNF as standard. Wild-type plant was exploited to be the negative control. Three independent duplicates were set to collect data for the statistical

Elimination of tag from GFP-fusion protein

analysis.

Purified GFP-fusion protein was digested by enterokinase according to the instruction of the Recombinant Enterokinase Kit (Novagen, Germany) to separate GFP and BDNF. Purification is carried out on AKTA pure150. Equilibrate 3-5 column volumes with balance solution (pH8.0, 20mmol/L Tris-HCl, 0.1mol/L NaCl) for CM Sepharose FF column. Loading raw BNDF on an ion-exchange column. Inflow balance solution 3-5 column volumes to baseline. Elution CM Sepharose FF at 1.5 mol/L NaCl with elution buffer (pH8.0, 20mmol/L Tris-HCl, 2.0mol/L NaCl) and collected effluent. The biological activity was confirmed by the 3T3 cell proliferation assay.

3T3 cell proliferation analysis

NIH3T3 cells were put in 96-well plates with 8×103 cells/well. They were cultured in Dulbecco's Modified Eagle Medium (DMEM; Gibco, China) containing 40ng/mL BDNF for 24 hours. An equal volume of PBS buffer and C-BDNF was added as the blank and positive group, and NIH3T3 cells cultured under the same conditions as the experimental group served as the control and positive group. Cellular proliferative activity was determined via Cell Counting Kit-8 (CCK-8) assays, in which cell survival rate = (A experimental group- A blank group)/(A control group - A blank group). Three independent duplicates were set to perform the statistical analysis. Statistical significance was analyzed with a one-way ANOVA test and the number of asterisks indicates the level of significance (* for p \leq 0.05 and ** for p \leq 0.01).

Results

Construction of expression vector

An expression vector named pWYP23402 for chloroplast transformation of tobacco was constructed based on the protocol previously [28] with some modifications (Fig. 1). The homologous fragment from alfalfa was replaced with fragment from tobacco plastid genome, and then the codon-optimized *BDNF* gene was inserted into the MCS of the expression vector. The vector was reconfirmed by sequencing.

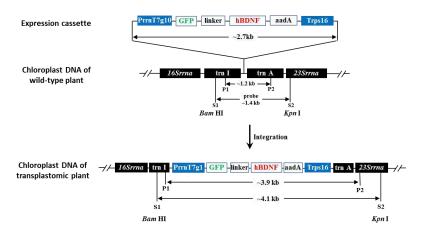


Fig. 1 Construction of expression vector pWYP23402 for chloroplast transformation. pWYP23402 vector flanked with sequences of 16S-trnI and trnA-23S comprised a synthetic expression cassette which contains gfp reporter gene, BDNF target gene, and aadA selectable marker gene. These three genes were jointly driven by the Prrn-T7g10 promoter and the terminator was Trps16. P1/P2 indicated the sites of primers used for PCR analysis and S1/S2 indicated the sites of digestion by BamH I and Kpn I for Southern hybridization. A \sim 3.9 kb fragment was expected to amplify for transplastomic plants, but only a \sim 1.2 kb fragment for wild-type plants. Similarly, total DNA was digested by BamH I and Kpn I for Southern blot analysis. A \sim 3.9 kb signal was hybridized for transplastomic plants, but only a \sim 1.4 kb for untransformed plants. The linker was inserted between the gfp gene and the BDNF gene and digested by enterokinase for BDNF purification subsequently. The amino acid sequence of the linker is GSSSGSSSGSSSDDDDK.

Tobacco homoplastomic selection and molecular characterization

Young leaves from PCR-positive shoots were cut into small pieces and cultured for two to three more additional cycles to obtain the homoplastomic plants (Fig. 2a).

Total DNA was extracted from young leaves of both transgenic and wild-type plants and the integration of the *BDNF* gene into the chloroplast genome was confirmed by Southern blot. As indicated in Fig. 2b, the single ~4.7 kb band in lanes 1, 3 and 4 implied homoplastomic status,

whereas lane 2 displayed an additional ~1.2 kb weak fade band, identically amplified from wild-type plants, indicating its heteroplastomic status.

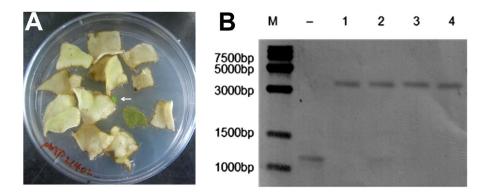


Fig. 2 Homoplastomic resistant callus after selection and molecular testing. **a** Resistant shoots (indicated by white arrow) appear while most leave pieces bleached on antibiotic medium; **b** Southern blot analysis. M, DNA molecular marker, Lane **1-4** The leaves of putative transgenic plants. The single 4.7 kb band in lanes 1, 3 and 4 indicated homoplastomic status, whereas lane 2 with an additional ~1.2 kb band, indicated heteroplastomic status. Line '-' is untransformed tobacco as the negative control.

Assessment of BDNF expression in transgenic plants

The SDS-PAGE gel was used to test the expression of GFP-fused recombinant BDNF. The results showed in Fig. 3a. Approximately 26.9 kD of GFP (lane P) and 40.1 kD of GFP-fused BDNF (lane 1-4) appeared clearly in the gel. In contrast, no such signal was observed in the lane wt which the crude protein was extracted from the untransformed plant.

For the western blot assessment, the results revealed approximately 40.1 kDa signals in lanes 1 to 4, representing BDNF expression in the transgenic lines (Fig. 3b). As expected, no signal was observed for wild-type plants.

The ELISA assay indicated that BDNF fused with GFP represented approximately $15.72\% \pm 0.33\%$ of total soluble protein, 2.417 ± 0.051 g kg⁻¹ of fresh leaf.



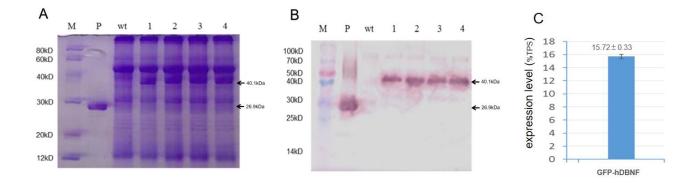


Fig. 3 SDS-PAGE analysis (**a**), Western blot (**b**) and ELISA assay (**c**) for determining the BDNF expression in transplastomic plants. **a** M, protein marker; **p** GFP as the positive control, wt, crude protein from the untransformed plant, **1-4**, crude proteins from four transgenic plants. **b** M, protein marker. Expression of GFP-fused BDNF (40.1 kD) was observed in the leaves of four transgenic plants (lanes 1-4), but absent in the leaves of a wild-type plant (**wt**). Lane **p** indicated the GFP as the positive control. **c** the expression level of GFP-hDBNF in the transgenic plants.

in vivo detection of GFP in transplastomic plants

As the reporter gene, *gfp* gene was inserted in the expression vector, together with the *BDNF* gene, to investigate the integration and the expression of target gene. As the same promoter and terminator was used to drive the *gfp* and *BDNF* genes which connected by a linker, indicating that the *in vivo* detection of GFP in the plant directly reflect the expression of target *BDNF* gene at the same time. The results demonstrated that the transplastomic plant emit strong green light under UV light (Fig. 4d), while the wild-type plant showed red color due to chlorophyll autofluorescence (Fig. 4c).

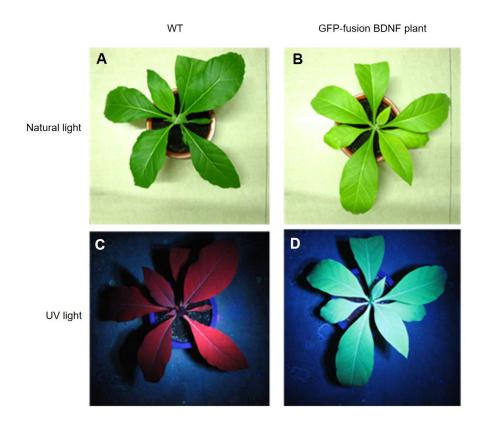


Fig. 4 *in vivo* detection of green fluorescent protein (GFP) in transplastomic plant. **a, b** The growth condition of wild-type and GFP-fusion BDNF plants under natural light, Wild-type plant (**c**) showed red autofluorescence whereas green fluorescence from transplastomic plant (**d**) under UV light.

Isolation and purification of BDNF and the evaluation of its biological activity

To purify the total soluble protein from the leave, the mature leave was ground in liquid nitrogen, add 5ml PBS buffer per gram. After extraction filtration by 0.22 µm filter membrane, the liquor was put into the 50KD tube, centrifuge for 30 min at 4000 ×g, the outflow was put into fresh 30KD tube, centrifuge for 45 min at 4000 ×g again. The outflow will be the primary product of GFP-fused BDNF in N terminal. The GFP-fused BDNF was digested by enterokinase to separate the GFP from BDNF. The target BDNF was purified by heparin affinity chromatography column. The results showed that the BDNF showed much higher biological activity then the negative control PBS buffer (Fig. 5b),

indicating the potential application in the market of medical proteins of raw materials. Meanwhile, the there is no significant difference between this chloroplast-derived BDNF and the commercial product as standard positive control purchased from Abcam (Cambridge, MA, USA) (See Fig. 5c).

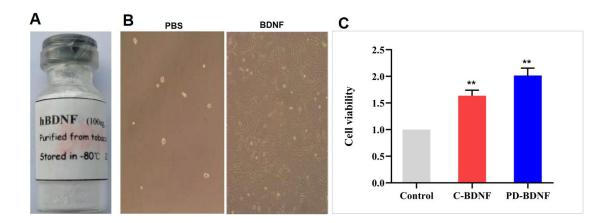


Fig. 5 Purified laboratory product of chloroplast-expressed BDNF and the its cell viability. **a** Cold-dry protein powder of BDNF expressed in chloroplast, **b** 3T3 cell proliferation promoted by BDNF growth factor. The result showed the biological activity of BDNF and the PBS buffer was set as negative control, **c** Comparison of cell viability of three different groups. Control indicated the negative group, C-BDNF indicated the biological activity of commercial product purchased from the company, PD-BDNF indicated the biological activity of BDNF derived from transplastomic tobacco plants. Both C-BDNF (red bar) and PD-BDNF (blue bar) were great significant difference (two asterisks) with the negative control (gray bar).

Discussion

Among existing expression systems of recombinant proteins including microorganisms [29-33], animals [34], and plants [25, 21, 22], plants have shown specific advantages over the other two production systems. The PMPs produced by both nuclear and chloroplast transformation offer some unique advantages over microorganism and mammalian systems like production in large-scale growing plants, cost-efficient, and lack of endotoxins and pyrogens [35, 36].

Current evidence strongly implicates that BDNF is a key and versatile regulator of neural circuit development and function. The secreted BDNF enables to mediate multiple processes in the mammalian brain including sleep deprivation, memory, mode disorder, neuronal differentiation and growth, synapse formation, and plasticity [11, 37-39]. Due to the multiple physiological functions of BDNF in the healthcare industry such as Parkinson's disease [40, 41], the growing supply-demand gap will be a great challenge for the scientists to fill in and the PMPs will be the batter option. For the plant-based expression system, chloroplast transformation has emerged as a new alternative to generate transplastomic plants due to some attractive features over nuclear transformation. The potential for high-level protein expression of this technology makes it feasible to produce therapeutics, vaccines, antigens, and commercial enzymes [42].

Originated from cyanobacterial cells around 1.5 billion years ago, chloroplasts are organelles only existing in the cells of plants. The incredibly high copy number of a cell, reaching up to 10000 copies per mesophyll cell [43], makes chloroplasts feasible to be the bioreactor for the production of therapeutic proteins with high expression levels [35, 44, 36]. Unlike nuclear transformation, gene silencing, variable expression level, and epigenetic regulation have not been observed during the practice of chloroplast transformation.

The expression level is always the concern of the producer. In this research, the expression level of GFP fusion BDNF can reach up to a rather high level of 15.7%. Such a high level of expression maybe benefit from the highly stable characteristics of GFP. Moreover, the target factor is still soluble and shows biological activity after separation with GFP. Considering commercial production in the future, two things should be addressed. One is the expression manner of BDNF. In this research, GFP fusion protein is expressed in tobacco chloroplasts. Subsequently, the enzyme digestion of fusion protein to separate the GFP and BDNF will greatly increase the overall cost and lower the yield of the BDNF. How to promote the yield of BDNF in the tobacco chloroplasts without fusion of GFP will be the challenge in the next step. The second one is the host of expression. The

seed-based system exploits the natural storage properties and therefore has additional merits such as easy transportation and storage, batch processing, highly stability [45]. The large biomass of tobacco mature leaves, however, will be the advantage with an average yield of 2.7 tons per hectare [46]. Moreover, it is practicable to harvest five to seven times annually under field conditions depending on the growing season.

It was reported that the production cost of the tobacco expression system is only one over fifty of the *E. coli* fermentation system [47]. Just recently, there is a report of launching a tobacco leaf-based pectinase production of the commercial-scale platform by Denill's Lab [48], indicating that engineered chloroplasts can offer attractive potential for industrial application just using crude leaf extracts without fermentation, purification, cold-chain transportation that microbial-enzyme products generally have. Moreover, the enzyme can maintain the full function in dried leaves making it much easier during its storage and crude enzyme extract, the competitive low cost, therefore, will be a remarkable feature for this platform. Similarly, plastid-derived growth factors like value-added BDFN will be also expected to develop such a platform to protect jobs and increase income of tobacco-growing farmers.

Compliance with Ethical Standards

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329 **Author contributions** Yunpeng Wang, Jieying Fan and Nuo Xu were responsible for material preparation and conducting 330 331 the experiments. Zhengyi Wei and Wen Xin were responsible for data collection and analysis, 332 Shaochen Xing and Yunpeng Wang were responsible for the draft of the manuscript. All authors read 333 and approved the final manuscript. 334 **Competing interests** The authors declare that there are no conflicts of interest. 335 **Ethics** approval 336 This article does not contain any studies with human participants or animals performed by any of the 337 authors. 338 **Consent to participate** 339 Informed consent was obtained from all individual participants included in the study 340 **Consent to publish** 341 All the authors have given the permission for the publication of the manuscript. 342 343 344 345 346

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