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# Gigantol from *Dendrobium chrysotoxum* Lindl. binds and inhibits aldose reductase gene to exert its anti-cataract activity: An *in vitro* mechanistic study



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# ARTICLE INFO

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Gigantol Aldose reductase (AR) gene Interaction UV–visible spectroscopy Atomic force microscopy (AFM) Surface-enhanced Raman spectroscopy (SERS)

# ABSTRACT

*Ethnopharmacological relevance: Dendrobium. chrysotoxum* Lindl is a commonly used species of medicinal Dendrobium which belongs to the family of Orchidaceae, locally known as "Shihu" or "Huangcao". *D. chrysotoxum* Lindl is widely known for medicinal values in traditional Chinese medicine as it possesses anti-inflammatory, anti-hyperglycemic induction, antitumor and antioxidant properties.

*Study aim:* To characterize the interaction between gigantol extracted from *D. chrysotoxum* Lindl and the AR gene, and determine gigantol's efficacy against cataractogenesis.

*Materials and methods:* Human lens epithelial cells (HLECs) were induced by glucose as the model group. Reverse transcription polymerase chain reaction (RT-PCR) was used to assess AR gene expression. Then, the mode of interaction of gigantol with the AR gene was evaluated by UV–visible spectroscopy, atomic force microscope (AFM) and surface-enhanced Raman spectroscopy (SERS). The binding constant was determined by UV–visible.

*Results:* Gigantol depressed AR gene expression in HLECs. UV–visible spectra preliminarily indicated that interaction between the AR gene and gigantol may follow the groove mode, with a binding constant of  $1.85 \times 10^3$ L/mol. Atomic force microscope (AFM) data indicated that gigantol possibly bound to insert AR gene base pairs of the double helix. Surface-enhanced Raman spectroscopy (SERS) studies further supported these observations.

*Conclusion:* Gigantol extracted from *D. chrysotoxum* Lindl not only has inhibitory effects on aldose reductase, but also inhibits AR gene expression. These findings provide a more comprehensive theoretical basis for the use of *Dendrobium* for the treatment of diabetic cataract.

# 1. Introduction

Diabetic cataract (DC) is a major cause of blindness in patients with diabetes (Minassian and Mehra, 1990; Wild et al., 2004; Pascolini and Mariotti, 2011; Rao et al., 2011; Bahmani et al., 2012; Pokharel et al., 1998; Nasir et al., 2013). While the development and progression of DC is caused by multifactorial mechanisms, the polyol pathway has been

recognized as one of the most important mechanisms (Agte and Gite, 2014; Srinivasan et al., 2012; Miyamoto, 2002; Yabe-Nishimura, 1998). Aldose reductase (AR) is the first and rate-limiting enzyme in the polyol pathway; thus, genetic polymorphisms of this enzyme impose genetic susceptibility to DC (Chung and Chung, 2003; Demaine, 2003; Wang et al., 2014; Li et al., 2015). High blood glucose levels can induce AR gene expression in human lens epithelial cells,

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Abbreviations: DC, diabetic cataracts; AR, aldose reductase; HLECs, human lens epithelial cells; APTES, aminopropyl triethoxysilane; UV spectra, Ultraviolet-visible spectroscopy; ETOH, Ethyl Alcohol; ETOAC, ethyl acetate; MEOH, Methyl alcohol; CC, column chromatography; AFM, Atomic force microscope; SERS, surface-enhanced raman spectroscopy; RT-PCR, real-time reverse transcription polymerase chain reaction; MEM, minimum essential medium; FBS, Fetal bovine serum; PBS, phosphate-buffered saline; IC50, half maximal inhibitory concentration

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and lead increased sorbitol production from glucose via the polyol pathway (Gudipudi et al., 2016; Susen and Buyukbingol, 2003; Kinoshita, 1990; Judy and Stanley, 1986). The resulting sorbitol production increases osmotic pressure, changes cell membrane permeability, and promotes the development of cataracts (Henry et al., 1993).

In traditional Chinese medicine, Shihu is a famous and precious Chinese herbal medicine derived from different species of Dendrobium, including D. chrysotoxum Lindl., D. nobile Lindl., and D. fimbriatum Hook. Shihu has been used to prepare various medicinal and health products in many Asian countries. D. chrysotoxum Lindl. is a commonly used species of medicinal Dendrobium, indexed in the Chinese Pharmacopoeia (Commission of Chinese Pharmacopoeia, 2010). Many ancient books e.g. "Shennong's Herbal", "Compendium of Materia Medica", "Supplement to compendium of materia medica", have been recorded its traditional treatment of nourishing yin and clearing heat, tonifying stomach and promoting fluid, improve eyesight (Pan et al., 2014). While the therapeutic potential for cataracts may be attributed to an additive/synergistic effect of constituents in D. chrysotoxum Lindl (Fang et al., 2015). Gigantol (Yang et al., 2006; Fan et al., 2014; Bhummaphan and Chanvorachote, 2015; Charoenrungruang et al., 2014; Wu et al., 2015), extracted from D. chrysotoxum Lindl., was found to exert anti-DC effects through the inhibition of AR activity down-regulation of AR gene expression in a galactose-induced cataract rat (Fang et al., 2015). However, the mechanisms underlying gigantol and AR gene interaction remain unclear. In this study, the human lens epithelial cells (HLECs) were cultured in 50 mM glucose medium to further validate gigantol curative effects. Reverse transcription polymerase chain reaction (RT-PCR) analysis showed that gigantol suppressed AR gene transcription in HLECs, and various biophysical tools were used to further monitor direct interaction of gigantol and the AR gene, including Ultravioletvisible spectroscopy (UV spectra), atomic force microscopy (AFM), and surface-enhanced Raman spectroscopy (SERS). Our findings provide insights on the mechanisms of gigantol and AR gene at the molecular level, as well as a comprehensive theoretical basis for the use of Dendrobium as anti-DC.

#### 2. Materials and methods

#### 2.1. Reagents and instruments

Glucose and y- aminopropyl triethoxysilane (APTES) were purchased from Aladdin Reagent Co. (Shang Hai, China); the HLECs (Ibarki et al., 1998) line SRA01/04 was a gift from the Ophthalmology Center of Sun Yat-Sen University (P.R. China). Minimum essential medium (MEM), Penicillin-Streptomycin and 0.25% Trypsin were purchased from Gibco (Grand Island, USA); Fetal bovine serum (FBS) from Biological Industries (Israel), mica from Tosai (Japan), and TRIzol reagent from Invitrogen (New Zealand). The plasmid pcDNA3.1 (-)-N1 was provided by Guangxian Zhang, Ph.D. (GuangZhou University of Chinese Medicine, China). The Endo Free Plasmid Maxi kit and DNA Gel Extraction kit was purchased from Sagene Biotech Co. (Guangzhou, China). The Prime STAR<sup>™</sup> HS DNA Polymerase kit was purchased from Takara (Japan). The plasmid pET28a-AR was purchased from Landbiology Co. (Guangzhou, China), the T4 DNA Ligase and the restriction enzymes BgIII and EcoRI from Takara Biotechnology, Co., Ltd (Japan). All-in-One™ qPCR Mix was purchased from GeneCopoeia (MD, USA). UV spectra was recorded using an Ultrospec 3300 pro UV-visible spectrophotometer (Biochrom Ltd, England), AFM images were taken using a Bruker MultiMode 8 scanning probe microscope (Germany), and SERS were recorded using a Micro-Raman spectrometer (H.J.Y. Company, French). The centrifugation in the experiment was used High speed refrigerated centrifuge, HR/T16M (Changsha, China).

#### 2.2. Gigantol extraction

The stems of Dendrobium chrysotoxum Lindl were collected from Puer municipal, Yunnan province, P.R. China and identified by Professor Hong Yu of the Yunnan University, Kunming, P.R. China. Voucher specimens (NO. Zsh-2 and Zsh-5) were preserved at the State Key Laboratory of Phytochemistry and Plant Resource in West China, Kunming Institute of Botany, Chinese Academy of Sciences. P.R. China. Gigantol was extracted from Dendrobium chrysotoxum Lindl, according to the method of Hu et al. with some minor modifications (Hu et al., 2012). The air dried stems of *Dendrobium chrusotoxum* (7.6 kg) were powdered and extracted with 95% aqueous EtOH (38 L×3) under reflux: the EtOH extract (105 L) was evaporated under reduced pressure and fractionated successively into EtOAc soluble (250 g) and n-BuOH soluble (249 g) fractions; EtOAc extract was subjected to column chromatography (CC) (silica gel (200-300 mesh, 3000 g), petroleum ether/Me2CO (7:3-0:10)) to give fractions I-VI; Fraction III (26.0 g) was subjected to CC (silica gel, petroleum ether/ Me2CO(4:1)) to afford fractions a-e, then the fraction c (5.4 g) was purified by CC (sephadex LH-20, CHCl3/MeOH, 1:1, v/v) to afford compounds gigantol (3.5 g) (Hu et al., 2012).

# 2.3. HLECs culture and experimental groups

HLECs were cultured in MEM supplemented with 20% FBS and 1% penicillin streptomycin at 37 °C in a humidified environment containing 5% CO<sub>2</sub>. The culture medium was changed after 24 h and then every alternate day. Confluent cells were washed with phosphate-buffered saline (PBS), harvested and seeded at a density of  $1-2\times10^{-5}$  cell/well. Twenty-four hours after seeding, the treated cells were grouped with control group (5.56 mmol/l glucose-MEM, A), model group (50 mmol/l glucose-MEM, B), and gigantol group (50 mmol/l glucose-MEM +1 µg/mL gigantol, C) for 24 h. Experiments were performed in triplicate.

# 2.4. RT-PCR

Total RNA was extracted and purified from HLEC using the TRIzol reagent, according to the manufacturer's instructions. RNA counts were quantified by measuring the absorbance at 260 nm. Samples with OD260/ 280 nm greater than 1.8 were used. RT-PCR was performed using All-in-One<sup>™</sup> qPCR Mix. In general, total RNA (1 µg) was added to a 20 µL reaction mixture for the reverse transcription; all PCR procedures were performed according to the manufacturer's instructions. The primers used this studv 5'CTTCCCTCCCTATCTA were: AR, in ACCCT3' and 5'TCTAAACTTCCCGTGGCATAA3'; β-actin, 5'-GAGACCTTCAACACCCAGCC-3' and 5'-GCGGGGCATCGGAACCGTCA-3'. The cycling condition was: 30 cycles at 98 °C for 10 s (denaturation), 58 °C for 15 s (annealing), and 72 °C for 1 min (extension). Gene expression was normalized to β-actin mRNA content.

# 2.5. Preparation of AR gene solution

The AR gene was prepared according to the method of Shao (Shao et al., 2015). The purified PCR products of AR gene which was amplified from the pET28a-AR plasmid and the plasmid pcDNA3.1 (-)-N1 were digested by the BgIII and EcoRI restriction enzymes for six hours at 37 °C, respectively. The digestion products were purified and isolated, and the target gene (AR gene) was then subcloned into the pcDNA3.1 (-)-N1 vector using T4 DNA ligase, forming the recombinant plasmid pcDNA3.1 (-)-N1-AR. DNA was extracted from plasmid bacteria pcDNA3.1 (-)-N1-AR by use of the Endo Free Plasmid Maxi kit. The target AR gene was separated from plasmid on 0.8% for agarose gel electrophoresis, taking the DL 1000 DNA Marker as the reference, then recycled gel with the AR gene. PCR amplification was conducted in a 25  $\mu$ L volume, using Prime STAR<sup>TM</sup> HS DNA

#### Table 1

Oligonucleotide primer pairs used for PCR assay.

Target gene	Primer designation	Nucleotide sequence 5 ' $\rightarrow$ 3 '
AR gene	AR-BgIII (F) Ard-EcoRI (R)	GAAGATCTATGGCAAGCCGTCTCCTG CGGAATTCTAAACTCTTCATGGAAGGGGT

Polymerase kit according to the manufacturer's instructions. Each assay contained 1  $\mu$ L plasmid DNA. Primers were designed by Sangon Biotech Co., Ltd (Shanghai, China) and were selected to amplify AR gene fragments (Table 1). After the plasmid DNA was denatured for 1 min at 94 °C, amplification conditions consisted of 30 cycles of 10 s at 98 °C and 15 s at 58 °C followed by extension at 72 °C for 1 min. The reaction was concluded with a final extension at 72 °C for 10 min. At last, the purified AR gene solution was obtained by use of the DNA Gel Extraction kit.

### 2.6. UV-visible spectroscopy

UV–visible spectra of gigantol and AR gene were monitored at 220–320 nm (Banitaba et al., 2011). AR gene solution  $(10^{-7} \text{ M})$  in 0.01 M Tris-Hcl buffer (PH7.4) was added to 4.5 mL colorimetric tubes and then mixed with 2 µL of selected concentrations of gigantol. 0.01 M Tris-Hcl buffer (PH7.4) was used as the blank control. The binging constant between AR gene and gigantol was calculated by the double-reciprocal formula (1) (Ouameur et al., 2005); and the Gibbs free energy ( $\Delta$ Gm) of the reaction was estimated from Eq. (2), and the reaction could proceed spontaneously and exergonic if the  $\Delta$ rGm is negative (Daikos et al., 2016).

$$1/(A0 - A) = 1/A0 + 1/(Ka \times A0 \times cDNA)$$
(1)

$$\Delta r G m = -R T ln K a \tag{2}$$

where A0 and A are the absorbance value of AR gene at 280 nm in the absence and presence of gigantol, respectively; Ka is referred to the binding constant; Ka is the binding constant at the corresponding temperature, and R is the gas molar constant.

#### 2.7. Atomic force microscopy

Fresh APTES modified mica, was prepared, according to previous studies (Prasad et al., 2016). AR gene solution was diluted with ultrapure water to 5 ng/µL; gigantol was diluted with ultrapure water to  $10^{-4}$ ,  $10^{-5}$ , and  $10^{-6}$  M solution. A total of 2 µL AR gene solution was added to 8 µL of gigantol solution. The final concentration of AR gene was 1 ng/µL. The resulting mixtures were incubated for 30 min at 37 °C. At the end of the incubation, the reaction mixture was dispensed onto the APTES modified mica plate, followed by the incubation for 10 min at room temperature. After rinsed for 5 ~ 6 times with ultrapure water, the plates were air dried and stored in a dry container until analysis. Experiments were carried out at room temperature and detected using tapping mode imaging.

#### 2.8. Surface enhanced Raman spectroscopy

The microwave heating method was used for the preparation of silver colloid (Kang and Zhou, 2012; Nahid Shahabadi et al., 2011; Zhang et al., 2005), using citric acid sodium to reducing silver nitrate to yield silver colloids with uniform particle size. The silver gum solution was obtained after centrifugation for 5 min at 12,000 rpm. The supernatant was discarded and the particles were then resuspended in deionized water, followed by ultrasonic cleaning for 5 min. The nano silver particles were washed with deionized water for three times to generate the final product with a higher concentration and purity. The AR gene and gigantol mixes were prepared by mixing 10  $\mu$ L of 0.9  $\mu$ g/L AR gene with 10  $\mu$ L gigantol at the concentration of 10<sup>-5</sup>, 10<sup>-6</sup> or 10<sup>-7</sup>



**Fig. 1.** Expression of AR gene in HLECs. The relative levels of AR gene expression in different group, and it was measured by real-time PCR. Results are shown as mean  $\pm$  SD (n=6);\*(P < 0.05), \*\*(P < 0.01).

M. To the resulting mixtures, Ag glue solution was added at the equal volume. Experimental excitation wavelength of 633 nm was applied for 5 s with 1 time exposure and a 50×magnification.

#### 2.9. Statistical analysis

Statistical analyses and data processing were performed using SPSS 22.0 software. All data are expressed as mean  $\pm$  SD. P < 0.05 was considered statistically significant.

#### 3. Results

#### 3.1. AR gene expression in HLECs

AR gene expression was increased by glucose by 2.2 fold compared to the control group (Fig. 1). But gigantol can significantly down-regulated the AR gene expression in HLECs by 1.35 fold compared to the model group (Fig. 1).

#### 3.2. Mode of interaction between gigantol and AR gene

We used UV-visible spectroscopy to investigate the absorbance spectra of AR gene in the presence of increasing concentrations of gigantol. The characteristic peaks of AR gene were observed in the range of near 262 nm. But on subsequent addition of gigantol to AR gene, the absorbance was gradually increased, indicating the hyperchromicity (Fig. 2).

AR gene and gigantol interactions in UV–visible spectra appear as red shift in maximum absorption peak. At fixed AR gene concentrations, gradual addition of gigantol results in 260 nm absorbance change (Fig. 3), binging constant between AR gene and gigantol was derived as Ka= $1.85 \times 10^3$  L/mol; and,  $\Delta rGm=-1.86 \times 10^3$  J/mol, which means that the reaction between gigantol and AR gene could proceed spontaneously. Form the red shift in UV–visible spectra indicated that the intercalative binding is most likely the dominant mode of the interaction between AR and gigantol.

# 3.3. Effect of gigantol on surface morphology of AR gene

To understand the cause of the interaction signal between gigantol and the AR gene in the UV-visible spectroscopic upon AR genegigantol binding, a series of tapping mode AFM images of AR gene were recorded with increasing concentrations of gigantol (Fig. 4 and Fig. 5). We can clearly saw that the AR gene was adsorbed on mica sheet surface by linear winding, the average height of AR gene computation for 1.2 nm (Fig. 4). After the AR gene interacted with gigantol, AR gene was cut into many fragments of varying sizes (Fig. 5-



**Fig. 2.** UV–visible spectroscopy of AR gene in 0.01 M Tris-Hcl buffer (PH7.4) at room temperature in the presence of increasing amounts gigantol. The concentration of AR gene was  $10^{-7}$  mol/L, and the concentration of gigantol was 0, 0.33, 0.66, 1.32, 2, 2.67,  $3.33 \times 10^{-9}$  µM from bottom to the top. Arrows indicated the change in absorbance upon increasing the gigantol concentration.



**Fig. 3.** The plot of 1/(A-A0) vs. 1/[AR gene] for AR and its complexes with gigantol, where A0 is the initial absorption band of free AR gene (280 nm) and A is the recorded absorption band at different gigantol concentrations.

A1), With increasing of gigantol concentrations, part of the AR gene started winding, forming spiral rings, even remaining together. Average height increased by about 0.3-0.6 nm s (Fig. 5-B1, C1).

3.4. The results of SERS for interaction between gigantol and AR gene

The interaction with gigantol and AR gene was characterized by SERS. And the SERS spectrum for gigantol and AR gene is shown in Fig. 6. The main bands were observed at Raman shifts: 422.63, 1143.29, 1264.98, 1391.20, 1576.39, 2116.81, 2920.27 cm<sup>-1</sup>. From these, the modes at 2920 cm<sup>-1</sup> are assigned to stretching vibrations of aliphatic CH groups. And the strongest SERS peak was located 2122.10 cm<sup>-1</sup> and 2122.86 cm<sup>-1</sup> for the AR gene and gigantol. In this experiment, the SERS peaks of AR gene had practically no signals, but there were appearing some peaks in the absence of gigantol. Once gigantol was added, the typical peaks of AR gene were detected. With the increasing amounts gigantol, the peaks of AR gene were greatly enhanced.

# 4. Discussion

In this work, it provided the evidence of binding mode of AR gene and gigantol to reveal the interaction between AR gene and gigantol. It indicated that the dominant binding mode of AR gene and gigantol was the intercalate binding. Our findings on the interactions of gigantol with AR gene had been studied using UV–visible spectra, AFM and SERS.

As we know, UV–visible spectra is a classical method for studying interactions between small molecules and DNA (Guo et al., 2007, Ouameur et al., 2005). In this experiment, we found that the red-shift was happened at the interaction of AR gene and gigantol. And the role of DNA and small molecules in UV–visible spectra will cause changes in maximum absorption peak and intensity change to induce the red shift/blue shift, or hyperchromicity/hypochromicity, which can help uncover the interaction mode between small molecules and DNA (Dehkordi et al., 2012; Jaumot and Gargallo, 2012). In addition, the red shift would appear at the intercalative binding between small molecules and DNA (Niu et al., 2015; Pakravan and Masoudian, 2015; Tysoe et al., 1993). This might indicate that gigantol was inserted into the AR gene base pairs. In addition, from the binding constant Ka  $(1.85 \times 10^3 \text{ L/mol})$  and  $\Delta r\text{Gm}$  ( $-1.86 \times 10^3 \text{ J/mol}$ ), indicated that the reaction of gigantol and AR gene was spontaneous (Gao and Liu, 2002).



Fig. 4. The surface topography of AR gene was assessed by AFM under the natural state. A, B were the images of AR gene at different amplification factor: A  $1 \times 1 \mu m$ ; B  $332 \times 332 nm$ . The AR gene has some winding in natural sedimentation and APTES adsorption, and the arrows indicated the AR gene.



Fig. 5. The images of AR gene with Gigantol at different concentrations interaction assessed by AFM. A and A1, B and B1, C and C1 were the images of gigantol at different concentrations of  $10^{-6}$ M,  $10^{-5}$ M,  $10^{-4}$ M, respectively. A, B, C, and A1, B1, C1were the images of AR gene at different amplification factor: A, B, C 1×1 µm; A1 378×378 nm, B1 350.2×350.2 nm and C1 503.9×503.9 nm. All images were acquired in tapping mode in liquid. Arrows indicated the AR gene.



**Fig. 6.** Raman spectra of the AR gene in gigantol at different concentrations gigantol.A, B, C, D and E were the images of AR gene, gigantol, and AR gene in gigantol at different concentrations of  $10^{-7}$ M,  $10^{-6}$ M, and  $10^{-5}$ M, respectively.

To further characterize the interaction of gigantol and the AR gene, AFM was employed to investigate the surface morphology change of AR. AFM could observed the DNA molecular structure *in vitro* at room temperature and dry air condition (Xu et al., 2016; Zheng, 2006). From the images of AFM, we observed that AR gene was forming many fragments of varying sizes after the interaction of AR gene and gigantol. The results of AFM provided great help in further identifying the binding interaction of the small molecule with DNA (Zhang et al., 2013; Zhou et al., 2016). Previous findings demonstrated that when linear DNA interacts with the drug, the latter can be inserted into the DNA double helix between the base pairs; this could be followed by the formation of a number of spiral DNA molecular chains, which can be cut into many fragments (Nahid et al., 2011; Salerno et al., 2010; Hou et al., 2009). Combine the results of UV–visible spectra, it indicated that the insertion binding was a major role in the interaction between gigantol and AR gene.

And, we used SERS to further research the mode of interaction between gigantol and the AR gene. Experimental studies in recent years showed that SERS allows the Raman scattering signal of the tested material enhancement 1014–1015 times, and  $10^{-7}$ M of DNA can also be detected (Yang et al., 2016; Kim et al., 2016; Li et al., 2016; Vaverkova et al., 2014). In this experiment, the SERS peaks of AR gene were greatly enhanced with the increasing amounts gigantol. The mechanism of enhancement is ascribed to the appearance of resonance between AR gene and gigantol, based on the fact that the resonance enhancement relative to the electromagnetic enhancement alone (Zhou et al., 2015; Chakraborty et al., 2016).

#### 5. Conclusions

In summary, in this research we provide new insights on the interaction of AR gene with gigantol extracted from *Dendrobium*  *chrysotoxum* Lindl. Our preliminary data revealed that gigantol interacts with AR gene through intercalate into the DNA base pairs and does not groove Binding. This is the first report on gigantol–AR gene interactions, and the biological data acquired in the present research should be valuable for the rational drug design and greater clinical efficacy.

#### **Conflict of interest**

The authors have declared no conflict of interest.

#### Authors' contribution

XYW(jidewowxy2@qq.com) was responsible for the design of the experiments; JW(445741136@qq.com), XL(xiaoxuesnowflak@ 163.com), WFM(550517704@163.com) and DC(1048456990@ qq.com) were contribution to the most of the experimental operation; WCW(wanwencheng@gzucm.edu) and QHY(297010172@qq.com) gave the guidance on experimental operation and the data statistic; JMH(hujiangmiao@mail.kib.ac.cn) was responsible for the extraction of Gigantol. JW and XL finished the writer of the article; OC (Oliver.Chen@tufts.edu) gave guidance to the modification of the article.

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#### J. Wu et al.

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