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# Determination of royal jelly freshness by ELISA with a highly specific anti-apalbumin 1, major royal jelly protein 1 antibody<sup>\*</sup>

Li-rong SHEN<sup>†</sup>, Yi-ran WANG, Liang ZHAI, Wen-xiu ZHOU,

Liang-liang TAN, Mei-lu LI, Dan-dan LIU, Fa XIAO

(Zhejiang Key Laboratory for Agro-Food Processing, Fuli Institute of Food Science, Department of Food Science and Nutrition, Zhejiang University, Hangzhou 310058, China)

<sup>†</sup>E-mail: shenlirong@zju.edu.cn

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**Abstract:** Major royal jelly protein 1 (MRJP1), designated apalbumin 1, has been regarded as a freshness marker of royal jelly (RJ). A MRJP1-specific peptide (IKEALPHVPIFD) identified by bioinformatics analysis of homologous members of the major royal protein family was synthesized and used to raise polyclonal anti-MRJP1 antibody (anti-SP-MRJP1 antibody). Western blot analysis showed that anti-SP-MRJP1 antibody only reacted with MRJP1 in RJ. In contrast, the previously reported antibody against recombinant MRJP1 (anti-R-MRJP1 antibody) reacted with other members of MRJP family in RJ. Enzyme-linked immunosorbent assay (ELISA) using anti-SP-MRJP1 antibody demonstrated that MRJP1 content in RJ stored at 40 °C significantly degraded by 37.3%, 55.9%, 58.0%, 60.6%, 65.7%, 72.7%, and 73.1% at 7, 14, 21, 28, 35, 42, and 49 d, respectively, when compared with MRJP1 content in fresh RJ (0 d). Optical density analysis of MRJP bands from sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) profiles demonstrated that the degradation of MRJP1, MRJP2, MRJP3, and MRJP5 in RJ was strongly and positively correlated with the period of storage (*P*<0.0001). Our results indicated anti-SP-MRJP1 antibody was highly specific for MRJP1, and ELISA using the antibody is a sensitive and easy-to-use method to determine the freshness and authenticity of RJ.

**Key words:** Freshness, Royal jelly, Major royal jelly protein 1 (MRJP1), Enzyme-linked immunosorbent assay (ELISA), High specific antibody

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# 1 Introduction

Royal jelly (RJ) is secreted by nurse honeybees from the hypopharyngeal and mandibular glands. This key nutrient, food for all young larvae, is vital in caste determination, development, and reproduction of the queen (Salazar-Olivo and Paz-Gonzales, 2005). Previous studies have demonstrated a wide range of health benefits to humans and model organisms, including vasodilative and hypotensive activities (Tokunaga *et al.*, 2004), antitumor (Tamura *et al.*, 1987), anti-inflammatory (Fujii *et al.*, 1990; Kohno *et al.*, 2004), antibacterial and disinfectant properties (Fujiwara *et al.*, 1990; Shen *et al.*, 2010a), anti-fatigue activity (Kamakura *et al.*, 2001a), and increasing the average lifespan in mice (Inoue *et al.*, 2003) and nematodes (Honda *et al.*, 2011). Because of its great potential benefits to human health, RJ is widely used as a key component in many commercial products including medicine, dietary supplements, and cosmetics (Kamakura *et al.*, 2001c). Thus, an easy-to-use and reliable method for the qualitative

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ORCID: Li-rong SHEN, http://orcid.org/0000-0002-3197-9245

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and quantitative evaluation of commercially available RJ products is needed. RJ contains 12%-15% protein, of which 80% belongs to soluble RJ proteins, the major RJ protein (MRJP) family (Howe et al., 1985; Simúth, 2001). Nine members of MRJP (MRJP1-9) have been reported; among them five (MRJP1-5) make up 82% of the total RJ protein content (Schmitzová et al., 1998; Malecová et al., 2003; Albert and Klaudiny, 2004; Drapeau et al., 2006; Shen et al., 2010b). MRJP1, designated as apalbumin 1, constitutes 45% of water-soluble proteins and is the most abundant protein in RJ (Furusawa et al., 2008). It is a glycoprotein of 432 amino acid residues with a molecular weight of 55-57 kDa (Hanes and Simúth, 1992; Shen et al., 2010b). MRJP1, in the form of an oligomeric complex, has more biological functions than its nutritional role in the larval development of honeybees. It is expressed in the brain mushroom bodies and nervous system of the adult (Peixoto et al., 2009). Previously designated as royalactin, it was recently found to induce the differentiation of honeybee larvae into queens through an epidermal growth factor receptor (EGFR)-mediated signaling pathway. Royalactin is a special factor for increasing body size, promoting ovary growth and shortening developmental time in both honeybees and Drosophila melanogaster (Kamakura, 2011). Moreover, MRJP1 was also demonstrated to have various biological functions in different model organisms in addition to its anti-fatigue effect in mice (Kamakura et al., 2001a). MRJP1 stimulates rat hepatocyte DNA synthesis, prolongs the proliferation of hepatocytes, increases albumin production (Kamakura et al., 2001b), and regulates mouse macrophages to release tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ) (Majtan *et al.*, 2006). Peptides derived from MRJP1 by enzymatic hydrolysis in the gastrointestinal tract possess potent antihypertensive activity in rats (Matsui et al., 2002). Furthermore, the C-terminal of MRJP1 may be a precursor of the antimicrobial peptides known as Jelleines (Fontana et al., 2004).

MRJP1 was suggested as a marker for the freshness of RJ because the degradation of this protein was especially associated with its storage temperature and storage period (Kamakura *et al.*, 2001c). Currently, 10-hydroxy-2-decenoic acid (10-HDA), which is found only in RJ, has been used as the typical quality criterion of RJ. However, its content changes little during storage at 40 °C for one week (Kamakura *et al.*, 2001c; Sabatini *et al.*, 2009). As no correlation between 10-HDA content and storage duration was found whatever the storage temperature, new criteria to measure the freshness in the duration of RJ storage are needed (Antinelli *et al.*, 2003).

High performance liquid chromatography (HPLC) analysis of MRJP1 separated from RJ with column chromatography has been used primarily as the standard method to determine the content of MRJP1 in RJ (Kamakura et al., 2001c). Recently, enzymelinked immunosorbent assay (ELISA) using purified MRJP1 with polyclonal antibody against recombinant MRJP1 expressed in Escherichia coli (anti-R-MRJP1 antibody) has been developed to measure MRJP1 in honey (Bíliková and Simúth, 2010). However, the polyclonal antibody against the recombinant MRJP1 is unsuitable for determining the content of MRJP1 in RJ because the quantification reflects the total amount of MRJP family members, which share a common evolutionary ancestor of the yellow protein from Drosophilia melanogaster (Albert et al., 1999). Therefore, more recently, a new method using a polyclonal antibody with a synthesized peptide (C+SGEYDYKNNYPSDID) specific for MRJP1 for the determination of RJ freshness was developed (Yamaguchi et al., 2013).

In this study, we have identified a specific MRJP1 peptide via bioinformatics analysis of amino acid sequences of homologous members of the MRJP family, and generated a new MRJP1 antibody specific for MRJP1. Our results demonstrated that this new molecule, anti-SP-MRJP1 antibody, binds more specifically and sensitively to MRJP1 than previously reported anti-R-MRJP1 antibody. Using ELISA with this antibody, we accurately detected the degradation of MRJP1 in RJ during storage at high temperature. Our ELISA method with anti-SP-MRJP1 antibody will be more applicable for quality control and authenticity identification of RJ in production and commerce around the world.

### 2 Materials and methods

### 2.1 Reagents

RJ from the western honeybee (*Apis mellifera*) was harvested 72 h after seeding young (1 d old)

larvae from an apiary provided by the Hangzhou Biyutian Health Food Corporation, Ltd. (Tonglu County, Zhejiang, China) and then was stored at -20 °C (Yamaguchi *et al.*, 2013). Diaminobenzidine (DAB) and goat anti-rabbit IgG-linked horseradish peroxidase (IgG/HRP) were purchased from Boster Biological Technology, Ltd. (Wuhan, China). Protein molecular weight standard was purchased from TaKaRa (Dalian, China). Nitrocellulose membranes (NCs) were purchased from Millipore (Billerica, MA, USA). CNBr-activated Sepharose 4B was purchased from GE (NY, USA). Sulfosuccinimidyl-4-(*N*-maleimidomethyl) cyclohexane-1-carboxylate (Sulfo-SMCC), keyhole limpet hemocyanin (KLH), and SulfoLink Resin were from Pierce (Appleton, WI, USA).

# 2.2 Homology analysis of MRJP sequences and specific peptide synthesis

The complete amino acid sequences of nine members of the MRJP family in the honeybee were acquired from GenBank web (http://www.ncbi.nlm. nih.gov). These MRJP sequences were aligned using Genetyx-Win Version 5 (Software Development Co., Tokyo, Japan). The unique peptide regions with more than three amino acid residues specific for MRJP1 were selected as the highly specific peptide region. A specific peptide that corresponds to the selected region was synthesized using the AMS 422 robot (ABIMED GmbH, Langenfield, Germany) by China Peptides Co., Ltd.

# 2.3 Preparation of polyclonal antibodies against MRJP1

The polyclonal anti-R-MRJP1 antibody was previously obtained by immunization of male New Zealand White rabbits with a recombinant MRJP1 expressed in *E. coli* in our laboratory (Shen *et al.*, 2010b). The anti-R-MRJP1 antibody was purified to homogeneity by HPLC, and was then conjugated to the amino groups of KLH, a carrier protein through the thiol group of the terminal cysteine residue for immunization of male rabbits to generate anti-SP-MRJP1 antibody. Briefly, three intradermal injections of KLH-coupled regional peptide of MRJP1 (KLHpeptide) to male New Zealand white rabbits at three-week intervals were followed by two subcutaneous injections of the antigen dissolved in complete Freund's adjuvant (Majtan *et al.*, 2006). The same purified specific synthetic peptide was conjugated to the amino acid of bovine serum albumin (BSA), a carrier protein through the sulfhydryl group of the terminal cysteine residue for preparation of antigen (BSA-peptide) affinity column and detection of binding activity to anti-SP-MRJP1 antibody as the antigen.

# 2.4 Purification and binding activity assay of anti-SP-MRJP1 antibody

Anti-SP-MRJP1 antibody was purified with antigen affinity column filled with SulfoLink Resin conjugated to BSA-peptide. The column was washed with 10 volumes of phosphate-buffered saline (PBS). The rabbit serum was filled with a Millex-GV 0.45  $\mu$ m filter, loaded onto the column, and then washed with PBS. The purified antibody was eluted with buffer (7.5% (0.075 g/ml) of glycine, pH 2.7), then dialyzed with deionized water (ddH<sub>2</sub>O). The titer of anti-SP-MRJP1 antibody to react with BSA-peptide was measured with ELISA (see Section 2.8).

# 2.5 Preparation of antibody column and purification of MRJP1

CNBr-activated Sepharose 4B was dissolved in 1 mmol/L HCl solution before the purified anti-SP-MRJP1 antibody was added. The mixture was then added to an affinity column and incubated overnight at 4 °C. In order to purify MRJP1, 10 g of fresh RJ was dissolved in 10 ml of ddH<sub>2</sub>O and vortexed. Then, RJ was ultracentrifuged (CP70MX Ultracentrifuge, Hitachi, Japan) at 245000g (Simúth, 2001), and placed on anti-SP-MRJP1 antibody affinity column for 2 h. The column was washed with 3 volumes of normal saline (0.9% (9 g/L) of NaCl) to remove unbound proteins. The native MRJP1 was eluted with 2 ml elution buffer (pH 3, ddH<sub>2</sub>O), analyzed by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE), and freeze-dried and stored at -20 °C. The purified MRJP1 was then freeze-dried and stored at -20 °C.

#### 2.6 SDS-PAGE and Western blot assay

To measure the specificity of the antibodies, Western blot analysis was performed using MRJP1 purified from fresh RJ. Eight  $\mu$ g (15  $\mu$ l) of each protein sample was separated by 12% SDS-PAGE according to the standard method (Sambrook *et al.*, 1989). After electrophoresis, protein bands on SDS-PAGE gels were transferred to NC for Western blot analysis. The anti-R-MRJP1 antibody and anti-SP-MRJP1 antibody were used as the primary antibodies at a dilution from 3000- to 5000-fold, respectively. The goat anti-rabbit IgG/HRP was used as the secondary antibody. The treated NC was reacted with DAB substrate as described previously (Shen *et al.*, 2010b).

# 2.7 Degradation analysis of MRJPs in RJ by gel documentation system

The protein concentration in RJ was determined by a total nitrogen Kjeldahl method (AQSIQ, 2008). Fresh RJ samples were dispensed into sterile 50 ml glass bottles. These bottles were then incubated at 40 °C, and were sampled every 7 d (0, 7, 14, 21, 28, 35, and 42 d). All samples were stored at -40 °C after thermal treatment immediately for SDS-PAGE analysis as previously described (Kamakura et al., 2001c). A total of 15 µl protein samples were analyzed by SDS-PAGE. MRJP bands from SDS-PAGE were quantified using SensiAnsys Quantity Analysis program in the gel documentation system (Shanghai Peiqing Science & Technology Co., Ltd., China). The differences (relative value, %) of the optical density (OD) of the MRJP protein band in the object RJ samples (7, 14, 21, 28, 35, and 42 d) relative to that of the same protein in fresh RJ (0 d) control were displayed according to the method described by the manufacturer's manual. The OD of the MRJP protein band in fresh RJ (0 d) control was regarded as 100%. Each assay was repeated with at least three independent replicates.

# 2.8 Determination of MRJP1 in RJ by ELISA with different antibodies

Micro-titration plates were coated overnight at 4 °C with 100  $\mu$ l/well carbonate buffer solution (CBS: Na<sub>2</sub>CO<sub>3</sub> 1.59 g, NaHCO<sub>3</sub> 2.93 g, pH 9.6) with 0, 2, 4, 6, 8, and 10  $\mu$ g/ml of standard MRJP1, and then blocked with 1.5% (0.015 g/ml) dried milk in PBS (137 mmol/L NaCl, 2.7 mmol/L KCl, 100 mmol/L Na<sub>2</sub>HPO<sub>4</sub>, 2 mmol/L KH<sub>2</sub>PO<sub>4</sub>, pH 7.4). Subsequently, 100  $\mu$ l of anti-SP-MRJP1 antibody and anti-R-MRJP1 antibody diluted with 0.1% (1 g/L) dried milk in PBS were added, and incubated in the coated wells for 1.5 h at 37 °C, respectively. The binding (titer) of

anti-SP-MRJP1 antibody and anti-R-MRJP1 antibody with MRJP1 as antigen was visualized using 100  $\mu$ l of an anti-rabbit IgG conjugated to peroxidase for 1 h at 37 °C, respectively. The color reaction was initiated by adding 100  $\mu$ l/well of tetramethyl benzidine. The reaction was stopped after 15 min by adding 2 mol/L H<sub>2</sub>SO<sub>4</sub> to the wells. OD was measured at 450 nm with an ELISA plate reader. Calibration curves in the concentration range from 0 to 10  $\mu$ g/ml MRJP1 were generated.

The content of MRJP1 in total RJ protein was measured to quantify the RJ freshness. The calibration curves of MRJP1 in a broad concentration range from 2 to 20 g/ml against anti-SP-MRJP1 antibody and anti-R-MRJP1 antibody were prepared in a 96-well plate in PBS separately, corresponding to the MRJP1 content of diluted PBS, and no antigen (MRJP1) was used as the blank control. The ELISA detection for MRJP1 content in fresh RJ samples using anti-SP-MRJP1 antibody and anti-R-MRJP1 antibody was implemented separately. Then, ELISA assay was performed to determine the degradation of MRJP1 in RJ samples from different storage periods (0, 7, 14, 21, 28, 35, 42, and 49 d) under 40 °C with anti-R-MRJP1 and anti-SP-MRJP1 antibodies, respectively. The MRJP1 content of each well was calculated based on its OD value with reference to the calibration curve. Each assay was repeated with at least three independent replicates. For the comparison with the ELISA quantitation, the density values of the MRJP1 bands on SDS-PAGE gel of the RJ samples from different storage periods were generated using ImageJ program (http://rsb.info.nih.gov/ij/index.html).

### 2.9 Data processing and statistics

For ELISA, fitting curves were performed with Microsoft Office Excel 2007. The statistics analysis was carried out with the SPSS 16.0 software (SPSS, Inc., Chicago, IL, USA). One-way analysis of variance (ANOVA) with Duncan's post hoc tests was adopted to determine any significant difference among treatment groups. The mean and standard deviation were reported, and P value of <0.05 was considered statistically significant. A linear regression model was used to determine the relationship between the density value of MRJP1 band on SDS-PAGE gel and MRJP1 concentrations measured by ELISA in RJ.

## 3 Results

# 3.1 Identification of specific regional peptide of MRJP1

The alignment of amino acid sequences of the nine members of the MRJP family (Fig. 1) showed high homology among them. MRJP1–5 share at least 10 regions with 3–5 identical amino acid residue sequences; MRJP1–9 share at least 7 regions with 3–4 identical amino acid residue sequences. This suggests that the anti-MRJP1 antibody raised with full length MRJP1 might recognize all MRJP family members. Through multi-alignment analysis, we identified four peptides with more than three unique amino acid residues, which are quite different from the corresponding regions of MRJP2–9 members: QDAILSG

MRJP1.	1:MTR-LFMLV-CLGIVCQGTTGNILRGESLNK-SLPILHEWKFFDYDFGSDERRQDAILSGEYD	
MRJP2.	1:MTR-WLFMVACLGIACQGAIVRENSPR-N-LEK-SLNVIHEWKYFDYDFGSEERRQAAIQSGEYD	
MRJP3.	1:MTK-WLLLVVCLGIACQDVTSAAVNHQRKSANNLAH-SMKVIYEWKHIDFDFGSDERRDAAIKSGEFD	
MRJP4.	1:MTK-WLLLMVCLGIACQNIRGGVVRENSSGKNLT-N-TLNVIHKWKYLDYDFDNDERRQAAIQSGEYD	
MRJP5.	1:MT-T-WLLLVVCLGIACQGITSVTVRENSPRK-LANSMNVIHEWKYLDYDFGSDERRQAAMQSGEYD	
MRJP6.	1:MTN-WLLLIVCLSIACQDVTSAIHQRKSSK-N-LEH-SMNVIHEWKYIDYDFGSDEKRQAAIQSGEYD	
MRJP7.	1:MTR-WLFMVACLGIACQGAILRENSAR-N-LKN-SLKVMHEWKYIDYDFGSEEKRQAAIQSDEYD	
MRJP8.	1:MIRWLLLMYLG-ITCQGVTDIHSRNLTNSLKV-IYEWKYIDYDFGSDEKRQAAIQSGDYN	
MRJP9.	1: MSFN1WWL1L-YFS-1VCQAKAHYSLRDFKAN1FQVKYQWKYFDYNFGSDEKRQAA1QSGEYN	61
	*	
MRJP1.	61 : YKNNYPSD I DQWHDK I FVTMLRYNGVPSSLNV I SKKVGDGGPLLQPYPDWSFAKYDDCSG I VSASKLA I D	
MRJP2.	62: HTKNYPFDVDQWRDKTFVTILRYDGVPSTLNVISGKTGKGGRLLKPYPDWSFAEFKDCSKIVSAFKIAID	
MRJP3.	67: HTKNYPFDVDRWRDKTFVTIERNNGVPSSLNVVTNKKGKGGPLLRPYPDWSFAKYEDCSGIVSAFKIAVD	
MRJP4.	66 : RTKNYPLDVDQWHNKTFLAV I RYNGVPSSLNVVSDKTGNGGRLLQPYPDWSFAKYEDCSG I VSAHK I A I D	
MRJP5.	65: HTKNYPFDVDQWRGMTFVTVPRYKGVPSSLNVISEKIGNGGRLLQPYPDWSWANYKDCSGIVSAYKIAID	
MRJP6.	65 : YTKNYPFDVDQWHNKTFLAV I RYDGVPSSLNV I SEK I GNGGCLLQPYPDWSWANYKDCSG I VSAYK I A I D	
MRJP7.	62: HTKNYPFDVDQWRDKTFVTVLRYDGVPSSLNVISEKTGNGGRLLQPYPDWSWTKYKDCSGIVSAYSIAID	
MRJP8.	59: YTMNYLLDTDQWGDKTFVIIMKFNGVPSSLNVITNKTGNGGPLLAPYPDWTWAKNENCSGITSAYKIEID	
MRJP9.	62: YKNNVPIDVDRWNGKTFVTILRNDGVPSSLNVISNKIGNGGPLLEPYPNWSWAKNQNCSGITSVYRIAID	131
	**.*.*	
MRJP1.	131: KCDRLWVLDSGLVNNTQPMCSPKLLTFDLTTSQLLKQVE I PHDVAVNATTGKGRLSSLAVQSLDCNTNSD	
MRJP2.	132: KFDRLWVLDSGLVNRTVPVCAPKLHVFDLKTSNHLKQIEIPHDIAVNATTGKGGLVSLAVQAIDLAN	
MRJP3.	137: KFDRLWVLDSGLVNNNQPMCSPKLLTFDLKTSKLVKQVE I PHN I AVNATTGMGELVSLAVQA I DRTNTM-	
MRJP4.	136: EYERLWVLDSGLVNNTQPMCSPKLFAFDLNTSQLLKQVEIPHDVATT-GKGELVSLTVQAMDSTNTMV	
	135: KFDRLWILDSGIINNTQPMCSPKLHVFDLNTSHQLKQVVMPHDIAVNASTGNGGLVSLVVQAMDPVNTIV	
MRJP6.	135 : KFDRLWVLDSGL I NN I QLMCSPKLLAFDLNTSKLLKQ I E I PHN I AVNASTGMGGPVSL VVQAMDPMNTT-	
	132 : KFDRLWVLDSGLVNNTQPMCFPKLLVFDLNSSQL I KQVD I PHE I AVNTTTEQGRLKSLAVQA I SSVN	
	129 : MCDRLWVLDSGL I NNVRSVCPPQLLVFDLNTSQLLKQVK I PHD I AVNTTTEKGALVTLSVQLLSCEVNGS	
MRJP9.	132: EWDRLWVLDNG I SGETS-VCPSQ I VVFDLKNSKLLKQVK I PHD I A INSTTGKRNVVTP I VQSFDYNN-TW	199
	201 : TMVY I ADEKGEGL I VYHNSDDSFHRL TSNTFDYDPKFTKMT I DGESYTAODG I SGMALSPMTNNL YY	267
	199: TLVYMADHKGDAL IVYQNADDSFHRLTSNTFDYDFRYAKMT IDGESTTLKNG I CGMALSPYTNNLY	
	206 :VY I ADEKGEGL IMYONSDDSFHRLTSNTFDYDPRYTKLTVAGESFTVKNG I VG I ALSPVTNNLYY	
	200 :YINDERGEGE IMTGRISDDSFRRETSNTFDTDFRFTRETVAGESFTVRNG	
MRJP4. MRJP5.		
	205:TMADDRGDALITYGNSDESFHRLTSNTFDTDPKTTRMMDAGESFTAGUGI-FGMALSPMTNNLTT 204:VYTADDRGDALITYGNSDDSFHRLTSKTFDNDLRYSELAVAGESFTVHDGTFGMALSPVTNNLYY	
	199: TLVY I ADNKGDGL I VYQNSDDSFIRL TSNTFNYDPRYTKMTVEGESFTVQDG I YGMALSPMTNNLYY	
	199: TLVY I GDNEGFAL I I YNNSDNSFORL TSATF AND RYTTKIII VEGESI FVGDA	
	200:	
micor9.		204
	.**	
MRJP1.		337
	266 : SPLASHGLYYVNTAPFMKSQFGENNVQYQGSED1LNTQSLAKAVSKNGVLFVGLVGNSAVGCWNEHQSLQ	
MRJP3.		
MRJP4.		
MRJP5.	269 : SPLSSRSLYVVNTKPFMKSEYGANNVQYQGVQD I FNTES I AK I MSKNGVLFFGLMNNSA I GCWNEHQPLQ	
MRJP6.		
MRJP7.		
MRJP8.		
	265 : SAMSSHNLNYVNTKQFTQGKFQANDIQYQGASDILWTQASAKA I SETGALFFGLVSDTALGCWNENRPLK	

corresponding to region 51–57, HDKIFVTM to region 73–80, NIRTVAQ to region 340–346, and IKEALPHVPIFD to region 359–370 of MRJP1 (Fig. 1). The longest peptide M1, with 12 residues at region 359–370 (IKEALPHVPIFD; Fig. 1), was selected as the region unique to MRJP1 for synthesizing the specific peptide.

# 3.2 Specific properties of anti-SP-MRJP1 antibody and anti-R-MRJP1 antibody

SDS-PAGE of MRJP1 isolated from fresh RJ by ultracentrifugation and antibody affinity showed that one protein band of about 57 kDa (Fig. 2a) was identical to MRJP1 in size as previously reported (Simúth, 2001). In contrast, the fresh RJ showed more than four proteins including MRJP1, MRJP2, MRJP3,

	338 : RHN I RTVAQSDETLQM I ASMK IKEAL PHVP I FDRY INR-E-Y ILVLSN-	
	336 : RONLEMVAQNDRTLQMIAGMKIKEELPHFVGSNKPVKDEYMLVLSN-	
	341 : RESEDVVAQNEETLQMIVSMK IMENLPQSGRINDPEGNEYMLALSN-	
MRJP4.	336 : RON I DVVARNEDTLQMVVSMK I KONVPQSGRVNNTQRNEYLLALSD-	381
	339 : RENMOMVAQNEETLQTVVAMKMMHL-PQSNKMNRMHRMNRVNRVNRMDRMDRIDRMDRMDRMDTMDTMDR	
	339 : RONMDMVAQNEKTLQM11SVK11QNLAYSGRMNR1HKNE-YMLALSN-	
	336: RENTDMVAQNEETLQMIVGMKIKQLLPHIVIIDIDNIINDEYMLVLTN-	
	336 : RRN I EVVAKNKETLQA I TGLKVKRR	
MRJP9.	335 : RRN I E I VAKNNDTLQF I SG I K I I KQ	359
	* ** *	
MRJP1.	384 :KMQKMVNNDFNFDDVNFR IMNANVNEL ILNTRCENPDNDR	423
MRJP2.	382 :RMQK I VNDDFNFDDVNFR I LGANVKEL I RNTHCVNNNQNDN I	423
MRJP3.	387 : RMQK I I NNDFNFNDVNFR I LGAN VDDLMRNTRCGRYHNQNAGNQNADNQNADNQNADNQNAD	448
MRJP4.	382 :RNQNVLNNDLNLEHVNFQ ILGANVNDL IRNSRCANFDNQDNNHYNHNHN	430
MRJP5.	408: IDRMDR-MDR IDR IDR MHTMDTMDTMDRTDKMSSMDRMDRMDRVDRMDTMDRTDKMSSMDRMDRMDRVDT	476
MRJP6.	385 :RMQK I VNNDFNFDEVNFR I LGANVNNL I KNTRCAKSNNQNNN	426
MRJP7.	384 : RMQK I LNNDLNFND I NFR I LIGGVSDLLENTRCTNFN I	421
	361 :ISF ILVHGFPLEY	
MRJP9.	360 :I SSN I YERONNE	371
	424:TPFK1S1HL	
MRJP2.	424 :QNTNNQNDNNQKNNKKNANNQKNNNQN	450
MRJP3.	449 : NQNANKQNGNRONDNRQNDNKQNGNRQNDNKQNGNRQNDNKONGNRQNGNKQNDNKQN	506
MRJP4.	431 :QARHSSKSDNQNNNQHN	447
	477: MDTMDTMDRMDRMDRMDRMDRMDRMDRMDRMDRTDKMSRIDRMDKIDRMDRMDRTNRMDRMNRMNRQMNEYMM	
	427 :QNKYKNQAHLD	
	422 : QNDDSDENNDDS I R I T I DASFN	
MRJP8.	374:EYVL	377
MRJP9.	372:YIW	374
	433 :	433
MRJP2.	451 : DN	452
MRJP3.	507 : GNRQNDNKRNGNR-QNDNQNNQNDN-NRNDNQVHHSSKLH-	544
MRJP4.	448 :DQAHHSSK-SNNRHNNND	464
	547: ALSMKLQKFINNDYNFNEVNFRILGANVNDLIMNTRCANSDNQNNNQNKHNN	598
	438 :	438
	444 :	444
MRJP8.	378 : AVSNR I QK V I YG-FDFND VNFR I LIAN VNDLIKNTRC I SP	416
MRJP9.	375: IVSNKYQKIANGDLNFNEVNFRILNAPVNQLIRYTRCENPKTNFFSIFL	423

## Fig. 1 Homology analysis of amino acid sequences encoded by MRJP1-9 genes of Apis mellifera

Conserved amino acid residues of the MRJP family were marked with asterisks under the amino acid residues of MRJP9. The specific amino acid residues of MRJP1 selected by homology analysis of the MRJP family are in gray. The peptide M1 with 12 residues at regions 359–370 designed for raised antibody is indicated by underlining. The sequence numbers of MRJP1–9 in GenBank are NM\_001011579 (MRJP1), NM\_001011580 (MRJP2), NM\_001011601 (MRJP3), NM\_001011610 (MRJP4), NM\_001011599 (MRJP5), NM\_001011622 (MRJP6), NM\_0010114429 (MRJP7), NM\_001011564 (MRJP8), and NM\_001024697 (MRJP9), respectively

and MRJP5 with 25–87 kDa (Fig. 2a), which were reported previously (Simúth, 2001). More importantly, in Western blot assay, anti-SP-MRJP1 antibody recognized MRJP1 only in RJ (Fig. 2b), whereas anti-R-MRJP1 antibody bound to other members of the MRJP family (Fig. 3b) corresponding to the SDS-PAGE profile of MRJPs (Fig. 3a). This suggests that anti-R-MRJP1 antibody lacks specificity for MRJP1. This finding demonstrated that anti-SP-MRJP1 antibody is highly specific for MRJP1, and the purified natural MRJP1 proportion is suitable for use as the standard in RJ quality control.

#### 3.3 Binding activity of anti-SP-MRJP1 antibody

The results in Table 1 illustrate the binding activities of anti-SP-MRJP1 antibody. The OD values of the antibody at 80000-fold dilution were 10 times that of the unimmunized rabbit serum (OD at 1250-fold dilution). This suggests that the antibody possesses a very high binding activity to MRJP1. Our previous result showed that the binding activity of anti-R-

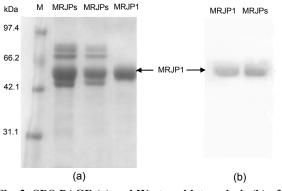


Fig. 2 SDS-PAGE (a) and Western blot analysis (b) of MRJPs probed with primary anti-SP-MRJP1 antibody Lanes: M, protein marker; MRJPs, soluble RJ proteins; MRJP1, purified natural MRJP1

MRJP1 antibody is 20000-fold dilution (Shen *et al.*, 2010b), which is only one fourth of the binding activity of anti-SP-MRJP1 antibody.

### 3.4 Thermal stability of MRJPs in RJ

Degradation of MRJPs in RJ samples stored at 40 °C from 7 to 42 d was measured using SDS-PAGE (Fig. 4). We found that overall the key MRJP proteins (i.e., MRJP1, MRJP2, MRJP3, and MRJP5) in RJ stored at 40 °C from 7 to 42 d, degraded consistently when compared with those of the fresh RJ (0 d). These findings suggest that not only MRJP1 but also other MRJP proteins are prone to degradation under thermal treatment.

#### 3.5 Standardization of ELISA with both antibodies

Based on ELISA with both antibodies, we generated two calibration curves: y=0.036x+0.264 ( $R^2=0.993$ ) for anti-SP-MRJP1 antibody (Fig. 5) and y=0.1100x+0.1239 ( $R^2=0.998$ ) for anti-R-MRJP1 antibody in a concentration ranging from 0 to 10 g/ml of MRJP1 in PBS, respectively.

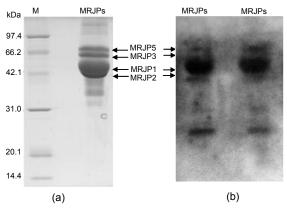
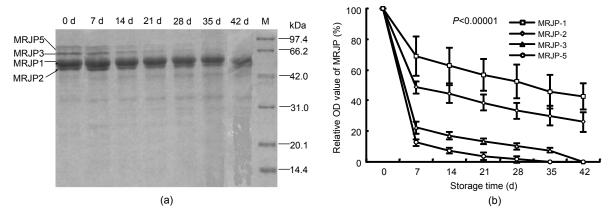


Fig. 3 SDS-PAGE (a) and Western blot analysis (b) of MRJPs probed with primary anti-R-MRJP1 antibody Lanes: M, protein marker; MRJPs, soluble RJ proteins

Table 1 Binding activity of anti-SP-MRJP1 antibody by ELISA

				OD	value at 450	) nm		
Sample	Dilution fold of anti-SP-MRJP1 antibody							CK at 1250-fold
-	1250	2500	5000	10000	20000	40 000	80000	dilution
1	2.376	2.375	2.375	2.368	2.071	1.233	0.637	0.071
2	2.375	2.375	2.375	2.376	2.375	2.184	1.611	0.068

CK: serum from normal rabbits





(a) SDS-PAGE analysis of RJ proteins stored at 40 °C from 0 to 42 d. Lanes of 0, 7, 14, 21, 28, 35, and 42 d refer to the RJ proteins stored at 40 °C from 0 to 49 d, respectively; Lane M refers to protein marker. (b) Degradation analysis of MRJPs in RJ. Relative OD value of MRJPs refers to the ratio of the OD of interest protein band in object RJ (7, 14, 21, 28, 35, and 42 d) relative to the OD of interest protein band in fresh RJ (0 d). The OD of interest protein band in fresh RJ was regarded as 100%. Data are expressed as mean $\pm$ SD (*n*=3)

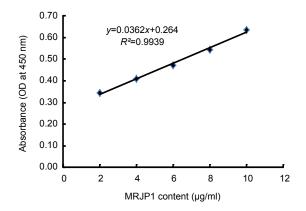


Fig. 5 Calibration curve of MRJP1 in PBS solution The calibration curve from 2 to 10  $\mu$ g/ml of MRJP1 in PBS was used for MRJP1 determination in 100-fold dilutions of RJ samples. The curves were prepared in triplicate for standard concentration

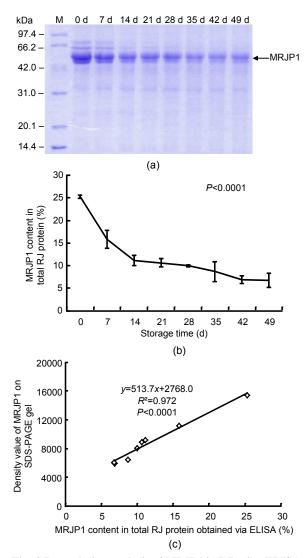
# 3.6 Comparisons of the specificity and sensitivity of both antibodies to MRJP1

ELISA results with anti-R-MRJP1 and anti-SP-MRJP1 antibodies showed different quantities of MRJP1 in the same RJ samples. The protein contents for RJ samples obtained with anti-R-MRJP1 antibody and anti-SP-MRJP1 antibody were  $(10.8\pm1.0)\%$  and  $(5.9\pm0.2)\%$ , respectively. The value  $(10.8\pm1.0)\%$  is consistent with the contents of MRJPs and MRJP1 for fresh RJ in previous reports, which showed fresh RJ contains 12%-15% crude proteins, of which about 80% are MRJPs (Furusawa *et al.*, 2008).

As demonstrated in Western blot analysis, anti-R-MRJP1 antibody bound to almost all members of MRJPs in RJ; this value (10.8%) measured by anti-R-MRJP1 antibody is an overestimate of MRJP1 protein in the fresh RJ. On the other hand, the value of 5.9% measured using ELISA with anti-SP-MRJP1 antibody is an accurate amount of MRJP1 in RJ, as MRJPs usually contains more than 45% MRJP1 (Furusawa *et al.*, 2008). Therefore, we consider that quantification of MRJP1 using ELISA with anti-SP-MRJP1 antibody is a reliable method.

### 3.7 Degradation of MRJP1 in RJ during storage

The concentration of total protein in fresh RJ was determined as (15.2±0.1)% using nitrogen Kjeldahl assay. Then, the content of MRJP1 in total protein of RJ samples stored at 40 °C from 0 to 49 d was estimated by measuring the density value of the MRJP1 band on SDS-PAGE (Fig. 6a), and quantitatively analyzed by ELISA using anti-SP-MRJP1 antibody (Fig. 6b). ELISA showed that the MRJP1 content of the total protein of RJ stored at 40 °C for 7, 14, 21, 28, 35, 42, and 49 d decreased by 37.3%, 55.9%, 58.0%, 60.6%, 65.7%, 72.7%, and 73.1% (n=3), respectively, relative to the MRJP1 content in the fresh RJ at 0 d (P<0.0001; Fig. 7). The relationship between the density value of MRJP1 and the MRJP1 content (measured by ELISA using anti-SP-MRJP1 antibody) in the total protein of RJ is given in Fig. 6c. The ELISA quantitation of MRJP1 in the total protein of RJ ranging from 5% to 30% was strongly and positively correlated with the density values of MRJP1 bands on the SDS-PAGE gel of RJ ( $R^2$ =0.972, P<0.0001). These findings suggest that the ELISA method measuring MRJP1 with anti-SP-MRJP antibody is a reliable tool for determining freshness and authenticity of RJ.



**Fig. 6 Degradation analysis of MRJP1 in RJ using ELISA** (a) SDS-PAGE analysis of RJ proteins stored at 40 °C from 0 to 49 d. Lanes of 0, 7, 14, 21, 28, 35, 42, and 49 d refer to RJ proteins stored at 40 °C from 0 d to 49 d, respectively; Lane M refers to protein marker. (b) Comparison of MRJP1 content in total protein (%) of RJ stored at 40 °C from 0 to 49 d using ELISA. Data are expressed as mean $\pm$ SD (*n*=3). (c) The correlation between the density value of the MRJP1 band on SDS-PAGE gel of RJ obtained via ImageJ program and the MRJP1 content in total protein of RJ samples measured by the ELISA method with anti-SP-MRJP1 antibody for RJ samples that were stored at 40 °C from 0 to 49 d

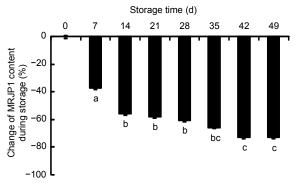


Fig. 7 Change of MRJP1 content in total protein of RJ during storage at 40 °C from 7 to 49 d

The change value (%) refers to the ratio of the MRJP1 content in total protein in object RJ sample (from 7 to 49 d) relative to the MRJP1 content of total protein in fresh RJ (0 d) control. The same letter marked at column stands for no significant difference (P>0.01). Data are expressed as mean±SD (n=3)

### 4 Discussion

In this study, we have successfully developed a new antibody, anti-SP-MRJP1, based on a unique peptide selected from the protein sequence of MRJP1. Western blot analysis demonstrated that this antibody binds specifically to MRJP1 only in RJ, in contrast to the previously reported anti-R-MRJP1 antibody that binds to almost all members of MRJP family (Li Y. et al., 2008). Thus, the ELISA method based on anti-R-MRJP1 antibody overestimated protein content of MRJP1 in RJ when compared with that based on ELISA with anti-SP-MRJP1 antibody. In addition, the anti-SP-MRJP1 antibody has a four-fold higher binding activity than anti-R-MRJP1 antibody. ELISA test of RJ using anti-SP-MRJP1 antibody can detect accurately the degradation of MRJP1 in RJ during the storage period of 49 d at 40 °C. SDS-PAGE analysis of those RJ samples demonstrated a similar phenomenon. Our results demonstrated that the anti-SP-MRJP1 antibody established in this study is highly specific for MRJP1.

RJ may spoil and lose its biological activities depending on storage duration and temperature. In order to preserve its biological properties, RJ has to be stored under freezing conditions (Zheng *et al.*, 2011). Therefore, it is essential to measure RJ freshness to ensure its quality for trade. Inappropriate storage of

RJ will be accompanied by acceleration of the Maillard browning reaction (Chen and Chen, 1995) and change of various physical and chemical properties including viscosity, titratable acidity, color, enzymatic activity, as well as the contents of MRJPs and free amino acids (Kamakura et al., 2001c). The compositional changes during storage at high temperature are not influenced by microorganisms and seem to be attributable to chemical and enzymatic activities. Interestingly, the degradation of MRJP1 in RJ was positively related to both storage temperature and storage period (Kamakura et al., 2001b; 2001c; Kamakura, 2011; Li J.K. et al., 2008). The results of native PAGE analysis for RJ stored at 4 and 40 °C showed that the two proteins in RJ, MRJP1 and MRJP2, were especially degraded when RJ was stored at 40 °C (Kamakura et al., 2001c). Furthermore, RJ stored at 40 °C for 7 d lost its activity for enhancing proliferation of rat hepatocytes, which was considered to be associated with degradation of MRJP1 (Kamakura et al., 2001b). According to two-dimensional PAGE, the amount of protein spots of MRJPs in fresh and -20 °C stored RJ was more than that in RJ samples stored at 4 °C and at room temperature for one year. The quantity of MRJP1 decreased significantly following the temperature trend in all images (Li J.K. et al., 2008). Therefore, MRJP1 is a typical marker for measuring freshness of RJ. Based on these findings, RJ heated at 40 °C for 40 d was used as the control to investigate the activity of MRJP1 for inducing the differentiation of honeybee larvae into queens (Kamakura, 2011). The presence of MRJP1 was also suggested as a promising parameter for evaluation of RJ authenticity in quality and standardization of RJ (Sabatini et al., 2009).

The anti-MRJP1 antibody generated with natural MRJP1 from RJ could recognize at least three MRJP members (MRJP1, MRJP3, and one 128-kDa MRJP member) in hypopharyngeal gland homogenates. In this investigation, the anti-MRJP1 antibody was generated with natural MRJPs as antigen, then separated by SDS-PAGE, and excised and electrotransferred on to an NC. Bioinformatics analyses on the analogous antigenic regions among MRJPs showed that MRJP1, MRJP2, and MRJP3 have more than 44% identity, and MRJP1–5 proteins share seven antigenic conserved regions corresponding to the epitopes for the anti-MRJP1 antibody (Peixoto *et al.*,

2009). The identities among MRJP sequences were suggested by hybridization assay and sequence alignment (Schmitzová et al., 1998; Albert et al., 1999; Drapeau et al., 2006; Peixoto et al., 2009). This antibody recognizes MRJP family members in the denatured protein. In addition, sequence analysis of all of the MRJP family members contains an N-terminal hydrophobic sequence that would function as a cleavable signal peptide as well as a putative N-linked glycosylation site (Drapeau et al., 2006; Peixoto et al., 2009; Klaudiny et al., 2010). Yamaguchi et al. (2013) were the first to determine MRJP1 levels in RJ samples from different companies with specific MRJP1 antibody. However, as a novel method, this article lacked essential information and details such as how to design peptide specific for MRJP1, purification of antibody, its difference from anti-R-antibody, and application in RJ freshness determination required by apiculture.

ELISA has been used as a rapid diagnostic tool in medicine, toxicology, as well as in quality-control in various industries around the world (Kajiura et al., 2008). Its accuracy is dependent upon the antigen used to create the polyclonal antibody. Special peptide immunogens are often used to generate polyclonal antibodies that target unique epitopes, especially for protein families of high homology (Shinnick et al., 1983; Lequin, 2005). Such antibodies are directed against a specific region of the protein chosen in advance by the investigator and so have a predetermined specificity (Huesca et al., 2000). ELISA with anti-R-MRJP1 antibody was used in quantification of MRJP1 in honeybees (Bíliková and Simúth, 2010). However, based on the observation in our study, the anti-R-MRJP1 antibody bound non-specifically to many members of the MRJP1 family because of common homology. Thus, ELISA with the anti-R-MRJP1 antibody may not be suitable for determining the freshness of RJ, but it may be used to measure the total content of the MRJP family

RJ adulteration is the most important problem affecting quality. Many studies on 10-10-HDA content (Kamakura *et al.*, 2001c), the ratios of the stable isotopes of the elements C and N, the fatty acid composition (Sabatini *et al.*, 2009) concerned with authenticity, qualification, and standardization of RJ have been carried out. Many indices to measure RJ freshness related to moisture, protein, total sugar, acidity levels and superoxide dismutase activity (Zheng et al., 2011), chromogenic reaction with HCl (Zheng et al., 2012), furosine (Messia et al., 2005), and changes in the content of amino acids (Wu et al., 2009) have been proposed. However, these indices either lack reliability as they are influenced by various factors or require complex sample preparation and expensive equipment, such HPLC for quantification. Thus, these indices are not widely accepted. MRJP1, the single specific protein and key factor of RJ associated with its biological activities as the unique marker of RJ freshness, has attracted much attention (Kamakura et al., 2001c; Sabatini et al., 2009; Yamaguchi et al., 2013). Moreover, compared with other methods described above, ELISA method developed by us has several advantages, including high reliability, lower cost, and ease-of-use.

In conclusion, we demonstrated that the anti-SP-MRJP1 antibody generated using a unique synthetic regional peptide corresponding to MRJP1 protein sequence as antigen is highly specific for MRJP1 in RJ. With this antibody, the quantification of MRJP1, the RJ freshness marker, is accurately determined. It is anticipated that such a method for measuring MRJP1 will be used as a reliable tool for determining the freshness and authenticity of RJ, and for detection of MRJP1 expression in honeybee tissues by using specific antibodies in the near future.

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### **Compliance with ethics guidelines**

Li-rong SHEN, Yi-ran WANG, Liang ZHAI, Wen-xiu ZHOU, Liang-liang TAN, Mei-lu LI, Dan-dan LIU, and Fa XIAO declare that they have no conflict of interest.

This article does not contain any studies with human or animal subjects performed by any of the authors.

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# <u>中文概要</u>

- 题 目:用高度特异性王浆主蛋白 1 多克隆抗体 ELISA 法检测蜂王浆新鲜度
- 目 的:为蜂王浆主蛋白 1 (MRJP1)的快速检测和鉴别 提供科学依据,为蜂王浆的质量控制提供技术 支持。
- 创新点: 首次比较了 MRJP1 特异性多克隆抗体与 MRJP1

重组表达蛋白多克隆抗体对王浆主蛋白家族的 免疫反应差异,验证了蜂王浆中 MRJP1 蛋白降 解与保温时间的相关性,建立了以 MRJP1 作为 蜂王浆新鲜度生物标志物的快速检测方法。

- 方 法:通过蜂王浆主蛋白家族蛋白的氨基酸序列同源性分析,筛选出 MRJP1 的特异性多肽区域,进行人工合成,免疫兔子后取血清制备成特异性多克隆抗体。用蛋白质印迹法(Western blot)检测了MRJP1 特异性多克隆抗体与 MRJP1 重组表达蛋白多克隆抗体对王浆主蛋白家族的免疫反应。以新鲜蜂王浆为对照品,用 MRJP1 特异性抗体酶联接免疫吸附剂测定(ELISA)法和变性电泳胶灰度扫描法分别测定保温(40℃)7~49天的蜂 王浆中 MRJP1 含量的变化,并进行了相关性分析。
- 结 论: MRJP1 的特异性抗体对 MRJP1 蛋白具有专一的 免疫识别特性,可特异性地检测代表蜂王浆新鲜 度的 MRJP1 含量变化,并鉴别蜂王浆的真伪。
- **关键词:**新鲜度;蜂王浆;王浆主蛋白 1; ELISA; 高度 特异性