

## 1 **Repository**

## 2 **Methods**

### 3 ***E1. FAHF-2 formula and product***

4 Dried aqueous extract of FAHF-2, produced in a good processing practice (GPP) certified  
5 facility (Xiyuan Chinese Medicine Research Pharmaceutical Manufacturer, Beijing, China), was  
6 obtained from Beijing Shen Hua Shi Di Medical Technology, Beijing China, and stored at room  
7 temperature. The quality of the raw herbs was established and described in details in previous  
8 publications.<sup>(1;2)</sup> In brief, based on organoleptic and microscopic examination, the raw herbal  
9 materials used in FAHF-2 were identified as the fruits of *Prunus mume*, the skin of the fruits of  
10 *Zanthoxylum schinifolium*, roots of *Angelica sinensis*, rhizome of *Zingiber officinalis*, twigs of  
11 *Cinnamomum cassia*, bark of *Phellodendron chinense*, rhizome of *Coptis chinensis*, roots of  
12 *Panax ginseng*, the fruiting body of *Ganoderma lucidum*. Product quality was monitored by  
13 HPLC fingerprinting according to the FDA's Guidance for Industry Botanical Drug Products<sup>(3)</sup>  
14 and as described in a previous publication.<sup>(2)</sup> The results of safety testing, including heavy metals,  
15 pesticide residue and microbes all met the standards for botanical products.<sup>(4-6)</sup> FAHF-2 and PN  
16 preparations were tested for endotoxin using the Pyrogen Plus assay kit (Lonza, MA). They  
17 were both undetectable with low limit of detection of <0.03 Endotoxin Unit/ml (EU/ml).<sup>(2)</sup>

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### 19 ***E2. Extraction and Isolation of Fractions and Compounds from FAHF-2***

20 Powered FAHF-2 extract (5 g) was dissolved into distilled water (125 ml). The solution was  
21 sonicated for 10 minutes followed by vortexing. This procedure was repeated for three times.  
22 The solution was transferred to a separatory funnel, and extracted with equal volume of butanol  
23 (125 ml, Fisher Scientific, Pittsburgh, PA, USA). After the separation, the organic layer was

24 removed and saved. The aqueous layer was re-extracted with butanol for three more times. All  
25 separated butanol extracts were combined and washed with distilled water at ratio 3:1. The  
26 butanol extracts were then evaporated using Rotorvaper R-210 (Buchi Corporation, New Castle,  
27 DE, USA) under reduced pressure. The dried extract butanol FAHF-2 () was collected and stored  
28 at 4°C.

29 Sub-fractions of B-FAHF-2 were collected using Preparative-HPLC (pre-HPLC) system coupled  
30 with UV detector (Waters, Milford, MA). Briefly, methanol dissolved B-FAHF-2 was loaded  
31 onto prep-HPLC with Xbridge™ C18 reverse-phase column 150mm×19mm×5µm (particle size)  
32 from Waters. The mobile phase used were 0.1% formic acid (mobile phase A) and acetonitrile  
33 (mobile phase B). The separation gradient started at 97% of A to 75% of A for 10min and to  
34 55% of A in another 20 min at a flow rate of 20mL/min. Four sub-fractions were collected based  
35 on their polarities.

36 Liquid chromatography mass spectrometry (LC-MS) system and commercially available  
37 standards were used to analyze the constituents in sub-fraction 2 of B-FAHF-2. Berberine and  
38 palmatine were purchased from Sigma-Aldrich (St Louis, MO). Jatrorrhizine was purchased  
39 from National Institute for the control of Pharmaceutical and Biological product of China  
40 (Beijing, P. R. China). Exact mass of each constituent was measured by high-resolution  
41 micromass LCT premier time-of-flight mass spectrometer (Waters Corporation, Milford, MA,  
42 USA) coupled to Waters Alliance 2695 HPLC system. The mass spectra range was set to be  
43 mass-to-charge ration (m/z) 50 to 1000. The mass spectrometer was set in positive mode with  
44 source conditions as follow: capillary voltage of 3200v, cone voltage of 25v, desolvation gas  
45 flow of 500L/h, cone gas flow of 40L/h, desolvation temperature of 350 °C , source temperature

46 of 110 °C. All spectra were obtained with an internal calibrant (lock mass with Leucine  
47 Enkephalin as reference).

### 48 ***E3. Assessment of systemic anaphylactic symptoms***

49 Anaphylactic symptoms were evaluated approximately 30 minutes after the challenge dose  
50 utilizing a previously described scoring system<sup>(7)</sup>: 0 - no signs; 1 - scratching and rubbing around  
51 the snout and head; 2 - puffiness around the eyes and snout, redness around snout, diarrhea, pilar  
52 erecti, reduced activity, and/or decreased activity with increased respiratory rate; 3 - wheezing,  
53 labored respiration, cyanosis around the mouth and the tail; 4 - no activity after prodding, or  
54 tremor and convulsions; 5 - death. Rectal temperatures were measured approximately 30 minutes  
55 following challenge using a thermal probe (Harvard Apparatus, Newark, NJ).

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### 57 ***E4. Histological assessment of mucosal and cutaneous mast cells degranulation***

58 For assessment of mucosal mast cell degranulation, sham treated and FAHF-2 treated mice were  
59 sacrificed forty hours after the final challenge at wk 18. The proximal jejunum from each mouse  
60 was snap frozen in OCT and 8µm transverse sections each were fixed in Mota's fixative  
61 containing lead acetate and stained with acidic toluidine blue. Mast cells were counted in 400x.  
62 Cutaneous mast cell degranulation during systemic anaphylaxis was assessed by examination of  
63 skin samples collected after challenge as described previously.<sup>(7)</sup>

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### 65 ***E5. Flow cytometry measurement of peripheral blood basophil numbers using FcεRI***

66 ***<sup>+</sup>CD49b<sup>+</sup> as marker to identify mouse basophils***

67 In addition to the previous method using the FcεRI<sup>+</sup>CCR3<sup>-</sup>Gr<sup>-</sup>,<sup>(8-10)</sup> other methods such as  
68 FcεRI<sup>+</sup>CD49b<sup>+</sup> have also been used recently as markers to identify mouse basophils.<sup>(11-13)</sup> To  
69 further validate the study on peripheral blood basophils, and to determine prolonged inhibitory  
70 effect of FAHF-2 on basophils, we conducted additional sets of experiments, in which mice were  
71 sensitized with peanut and CT as previously described.<sup>(2;14)</sup> FAHF-2 treatment began at wk 16  
72 (32 mg in 0.5 ml water , twice a day i.g) for 10 wks. Approximately 100μl blood samples were  
73 collected from each mouse 23 wk post FAHF-2 therapy. PBLs were processed and Fcγ  
74 receptors were blocked with anti-mouse CD16/32 mAb (2.4G2). PE-conjugated anti-mouse CD3,  
75 PE-conjugated anti-mouse B220, FITC-conjugated anti-mouse FcεRI, and APC-conjugated anti-  
76 mouse CD49b were added to the cell suspension. CD49b<sup>+</sup> FcεRI<sup>+</sup> CD3<sup>-</sup>B220<sup>-</sup> cells were  
77 defined as mouse blood basophils.<sup>(13)</sup> Data were analyzed using Flowjo software (Tree Star, Inc.  
78 Ashland, OR).

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#### 80 ***E6. Quantifying FcεRI subunit mRNA levels by real-time PCR (qPCR)***

81 Real-time reverse transcription quantitative PCR (qPCR) was performed using a SYBR green  
82 protocol and an ABI7900 HT machine (Applied Biosystems, Foster City, CA, USA) as  
83 previously described.<sup>(15)</sup> Briefly, 1x10<sup>6</sup> MC/9 cells were cultured in the presence or absence of  
84 FAHF-2 (20μg/ml) for 24 hr. Cells were then exposed to 2 μg/ml of mouse anti-DNP IgE for 5h.  
85 Cells were harvested and RNA was extracted by using Trizol reagent ( Invitrogen, Carlsbad, CA)  
86 according to the manufacture's instruction. The RNA was then deoxyribonuclease (DNase)  
87 treated (Invitrogen). cDNA was prepared from these samples using oligo-dT primers (Invitrogen)  
88 and Omniscript reverse transcript (QIAGEN, Valencia,CA).The cDNA was used to quantify the

89 mRNA levels of FcεRI α, β and γ subunits by using real-time PCR. The primers used were as  
90 follows: FcεRIα (sense primer 5'-TGGGAACAATCACCTTCAAA-3 and anti-sense  
91 5'CAGCCAATCTT GCGTTACAT-3' ); FcεRIβ(sense 5'-  
92 CATTAAAGGTCCAGACTCCA-3' and anti-sense 5'-CGTCCCATATCAATTACCCA-3');  
93 FcεRIγ (5'-CGCAGCTCTGCTATATCCTG-3' and antisense 5'-  
94 ACAGCATCTGCTTTCTCACG-3'). The primers were designed based on National Center for  
95 Biotechnology Information accession numbers NM01084 (α subunit), AB033617 (β-subunit)  
96 and NM10185 (γ subunit).

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#### 98 ***E7. Western blot***

99 Determination of early FcεRI-mediated signal events was performed by Western blot analysis  
100 using anti-Syk and anti-phospholated Syk antibodies (Cell Signaling, Danvers, MA) as  
101 described previously.<sup>(16)</sup> Briefly, IgE-sensitized RBL-2H3 cells ( $7 \times 10^6$ ) were cultured with or  
102 without B+P+J for 24 hours and stimulated with DNP (150μg/ml) for 5 minutes. The cells were  
103 then lysed following manufacturer's instruction for Nuclear/Cytoplasmic Extract Kit (Active  
104 Motif, Carlsbad, CA) and 50 μg of whole cell proteins was separated by SDS-PAGE and blotted  
105 onto a nitrocellulose membrane. After overnight blocking with 5% fat-free milk in PBST, the  
106 membranes were incubated with anti-Syk/phospho antibody for 2 hours. After washing with  
107 phosphate buffered saline Tween-20 (PBST), the membranes were incubated with peroxidase-  
108 conjugated anti-rabbit IgG. Immunoreactive bands were visualized by using the ECL Western  
109 detection system (Amersham Pharmacia Biotech, Piscataway, NJ).

#### 110 ***E8. IgE binding test by flow cytometry***

111 IgE binding test was performed as described previously.<sup>(17)</sup> RBL-2H3 Fc  $\gamma$  receptors were first  
112 blocked with anti-Fc $\gamma$ RII/III(2.4G2). 20 $\mu$ g/ml of berberine, palmatine or jatrorrhizine and then  
113 1 $\mu$ g/ml mouse IgE antibody were incubated with RBL-2H3 (1X10<sup>6</sup>/ml) at 4°C for 1 hour. Cells  
114 were washed twice with PBS, then FITC-anti-mouse IgE was added to detect binding of IgE on  
115 RBL-2H3 cells (4°C, 30 minutes) and then analyzed. Cells were incubated without IgE in the  
116 presence or absence of berberine, palmatine or jatrorrhizine ,and then stained with FITC-anti-  
117 mouse IgE for use as negative controls.

## 118 **Results**

### 119 **E1. Reduction of basophil number and anaphylactic reactions**

120 In additional set of experiments, using the CD49b<sup>+</sup>Fc $\epsilon$ RI<sup>+</sup> basophil criteria, numbers of basophil  
121 in FAHF-2 treated mice remained significantly lower than sham treated mice 23 wks post  
122 treatment (Figure E1A,B, p<0.01). Consistent with these findings, FAHF-2 treated mice were  
123 protected following peanut challenge at this time point (Figure E1C,D,E, p<0.05 for all).

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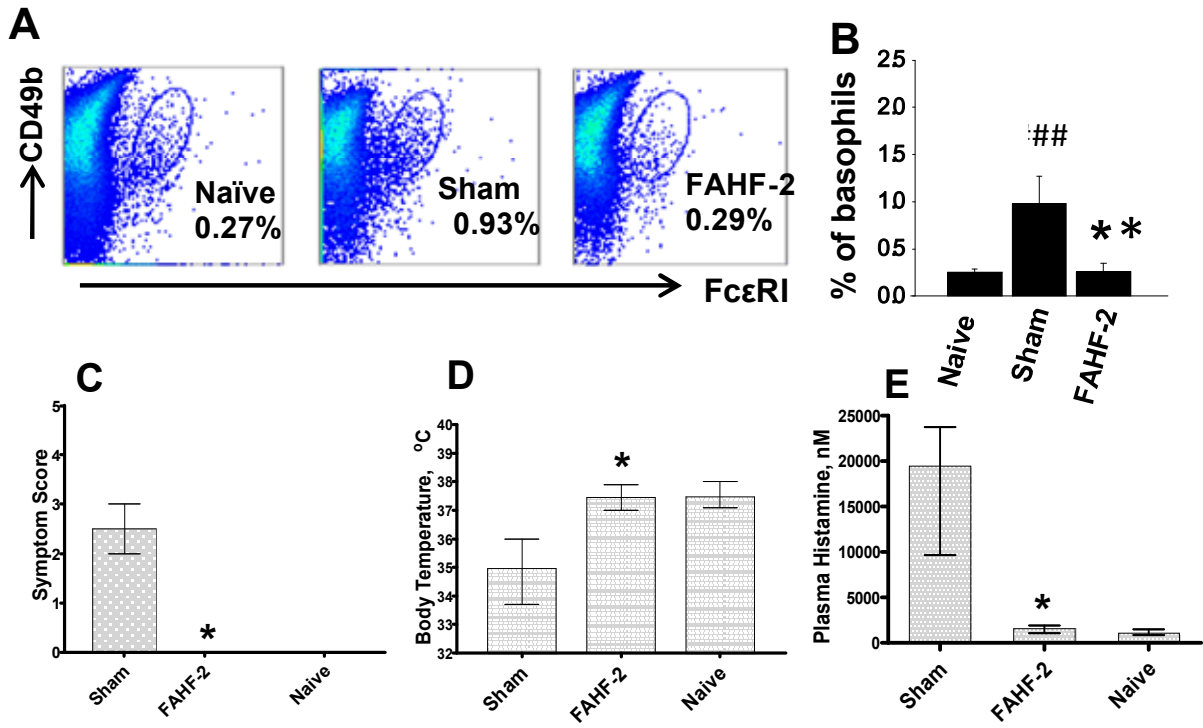
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130 E. Figure

131 Figure E1



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135 **E. Figure legend**

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137 **E1: FAHF-2 induced prolonged inhibition of peripheral basophils and protection against**

138 **peanut anaphylaxis. A.** Dot plots are representative of individual mouse blood basophils from

139 each group 23 wks post therapy. **B.** Bar graph shows Mean  $\pm$  SEM of 3 individual mouse from

140 each group. Data were analyzed by One-way ANOVA followed by Bonferroni test. \*\* $p < 0.01$ ,

141 Sham vs. FAHF-2; ##  $p < 0.01$ , Sham vs. Naïve. **C-E.** Clinical reactions 30 mins following oral

142 peanut challenge. **C.** Anaphylactic symptom score. **D.** Rectal temperature. **E.** Plasma histamine

143 levels. Data were analyzed by Kruskal-Wallis One Way Analysis of Variance on Ranks followed

144 by Student-Newman-Keuls Method. Bars show median; error bars show range.  $n=4$ . \* $p < 0.05$ ,

145 Sham vs. FAHF-2.

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