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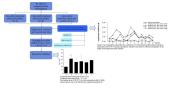
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An integral evaluation method for the synchrony of drug release based on the mathematics set was developed for guiding the preparation of multi-component TCM.

1	An integral evaluation method for the synchrony of drug release based on the
2	mathematics set in guiding the preparation of a multi-component traditional
3	Chinese medicine
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21	Running title: An integral evaluation method for the synchrony of drug release

1 Abstract

2	The quantitative characterization and evaluation of the synchrony of
3	multi-component release behavior are bottlenecks urgently needed to solve in the
4	studies on multi-component release kinetics and the quality control of
5	multi-component traditional Chinese medicine (MCTCM). MCTCM is from the
6	original prescription and preparation. At present, the evaluation of MCTCM is
7	generally full use of pharmacodynamics evaluation. But it is lack of convenience and
8	can not clearly reveal the correlation between the whole dynamic process of drug
9	release and <i>in vivo</i> absorption. It is also very limited for the control of preparation.
10	In addition, there are various types of effective ingredients in MCTCM, which exist
11	distinct differences in physicochemical properties. These differences may lead to
12	asynchrony of drug release between MCTCM and original preparation. So that they
13	thereby directly affect the efficacy. However, in this issue through the
14	pharmacodynamics evaluation we can't find the core reason. The papers reported
15	were confined to studying the release characteristics of one or a few components in
16	MCTCM and the drug release evaluation method reported was too macroscopic to
17	specifically identify the components which caused the integral asynchrony.
18	In this paper, in order to reveal the MCTCM release synchronous characteristics,
19	Fuzhenghuayu Capsules selected as the model, an original preparation, the integral
20	release evaluation method based on the mathematics set was established in guiding
21	the preparation of MCTCM. This method can not only be used in evaluating release
22	characterization of MCTCM by the parameter of asynchronous coefficient, but also

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1	be used in adjusting the dose of release unit by the relative error parameter. The
2	results demonstrated that this evaluation method was feasible, stable and
3	reproducible. And the Fuzhenghuayu Pellets guided by this method were showed
4	release synchrony and similar pharmacodynamics with the original capsules, and the
5	drug release mechanism was mainly frame erosion. Through this study, we could
6	then evaluate the quantifiable release characteristics of MCTCM and it could apply
7	an integral synchronous evaluation method for MCTCM.
8	
9	Keywords: drug release set; evaluation method of the synchrony; drug release;
10	chromatographic fingerprint; Multi-Component Traditional Chinese Medicine

11

1 1. Introduction

2	Currently, multi-component traditional Chinese medicine (MCTCM) is from an
3	effective and classical formulation, which has clear active substances in the clinic and
4	confirmed by pharmacodynamics. ¹ According to the characteristics of the different
5	effective components in the formulation, the MCTCM is usually comprised of two or
6	more types of release units, such as pellets, granules, microcapsules etc. Then
7	release units were recomposed into the capsules. Moreover, through optimizing
8	different release unit composition, MCTCM can produce similar or superior effect to
9	the original preparation. Presently, optimizing release unit composition of MCTCM
10	depended on an pharmacodynamics assessment, such as pharmacological effect
11	intensity, the onset time and the continuing role of time, et al. It is available for
12	ensuring the rationality of the preparation as an terminal evaluation. However, it can
13	not describe the whole release behavior of MCTCM. Furthermore, it can not find the
14	reason of asynchronous release behaviour due to the method of preparation, the
15	different physicochemical properties of ingredients, et al. These factors may also
16	directly affect the curative effect of the drug. Therefore, attention should be paid to
17	the quantification of synchrony of multi-component release research in MCTCM.
18	At present, there are mainly following three aspects of release evaluation of
19	MCTCM. ²⁻⁴ The first one is the evaluation method based on the chemical composition
20	which is with one or a few indicator components quantitative analysis. ⁵⁻¹¹ This
21	method is simple, fast and strong representative, but it is only suitable for single
22	component preparation. The second one is the evaluation method based on

1	pharmacological effect and biological effect, a biological potency measurement
2	method, which is used for the terminal evaluation. $^{12-14}$ The third one is the fuzzy
3	evaluation method based on component theory, including Quantified fingerprint
4	method ¹⁵⁻¹⁷ and Kalman filter method ¹⁸⁻²² which was used in the evaluation of the
5	synchrony of Yinqiaojiedu tablets ¹⁹ , Tongxuanlifei pills ¹⁶ , Dachuanxiongfang pellets,
6	Shaoyaogancaofang pellets and Yuchangning pellets ¹ . Although Kalman filtering
7	method was reported, because of its Ultraviolet spectrophotometry method with
8	detecting all absorption in only one wavelength as "component score", this method
9	was too general to be able to specify which components caused integral asynchrony.
10	As a result, it failed to identify which component accounted for the overall
11	asynchrony and also failed to provide an effective reference for the design of
12	MCTCM.
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1	shown in Supplementary Data 1.
2	The change of material composition would lead to change the coexistence
3	environment of different polar substances. It may lead to the asynchrony release
4	between the original preparation and recomposed release unit. Therefore, three
5	FZHY Pellets (FZHY-P) with different release behaviors were prepared using modern
6	technology according to the physical and chemical properties of the effective
7	components. And the integral evaluation method for the synchrony of drug release
8	based on the mathematics set was established. Moreover, relative error ($arepsilon$),
9	asynchronous coefficient (k) and other parameters were established for achieving the
10	synchrony of the drug release behavior compared with the original preparation and
11	recomposed preparation, screening the drug release unit type and dose. Through the
12	above research, it provided an effective evaluation tool of drug release for the
13	quantitative characterization and the synchrony of MCTCM.
14	2. Materials and methods
15	2.1 Chemicals and materials
16	FZHY-C was purchased from Shanghai Huanghai pharmaceutical Co., LTD
17	(Shanghai, China). FZHY-P were self-made, with fat-soluble immediate release unit
18	(pellets) of 2.5% drug loading for fat soluble active components (including Cordyceps
19	fat soluble components, Gynostemma pentaphyllum saponins and Amygdalin) and
20	water-soluble immediate release unit (pellets) or water-soluble sustained release
21	unit (pellets) of 33% drug loading for water soluble active components (including
22	Cordyceps polysaccharide, Salvianolic acid and Gynostemma pentaphyllum

1	polysaccharides). The preparation of fat-soluble immediate release unit (type: pellets,
2	No: A), water-soluble sustained release unit (type: pellets, No: B) and water-soluble
3	immediate release unit (type: pellets, No: C) were showed in Supplementary Data2.
4	All these pellets with different unit type and dose ratio were put into empty capsule
5	shells.
6	Reference standards of Protocatechuic aldehyde (>98%) and salvianolic acid B
7	(>98%) were purchased from the National Institute for the Control of Pharmaceutical
8	and Biological Products (Beijing, China). 50% total salvianolic acid extract (TSA) and
9	80% total gypenoside extract (TG) were obtained from Shanghai Huanghai
10	Pharmaceutical Co., Ltd. (Shanghai, China). Dimethylnitrosamine (DMN) were
11	provided by Tokyo Chemical Industry Co., Ltd (Tokyo, Japan). 4-Dimethylamino
12	benzaldehyde, Chloramines-T, Anhydrous ethanol, Xylene, formaldehyde,
13	Pentobarbital Sodium, Isopropanol (Analytic grade), Concentrated hydrochloric acid
14	were all obtained commercially by China Sinopharm Co., Ltd of Shanghai chemical
15	reagent Company (Shanghai,China). Sodium Chloride Injection (0.9%) was bought by
16	Anhui Shuanghe Pharmaceutical Co., Ltd (Anhui, China). Hydroxyproline (Hyp) was
17	bought by Sigma-Aldrich Chemie GmbH (American). Perchloric acid was bought by
18	Shanghai taopu Chemical plant (Shanghai, China). Phosphoric acid was excellent
19	pure-grade and all other chemicals were analytical grade. Acetonitrile and methanol,
20	both HPLC grade, were purchased from Merck (Germany). Pure water system (for
21	HPLC/UP) was bought by Labconco Company (American).
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22 **2.2** Dissolution test and preparation of reference solution and sample solution

1	A DT800-Ls intelligent dissolution instrument was purchased from Tianjin
2	university wireless power plants. The dissolution test of FZHY-C and FZHY-P was
3	prepared in accordance with the China Pharmacopeia. ²⁹
4	For each batch of reference solution, 20 FZHY-C were accurately weighted,
5	equivalent to 2.00 times of single capsule weight (\pm 0.1 mg), was "fully" dissolved in
6	250 mL of distilled water by dissolving for 4 hours. The rotational speed was set at 75
7	revolutions per minute , and the temperature was set at 37 \pm 0.5°C. The solution (5
8	mL) was filtered through a PTFE syringe filter (Millipore, pore size 0.22 μ m, diameter
9	13 mm) as the calibration sample with a gross concentration of 200%, which
10	represented 2.00 capsule was "fully" dissolved in 250mL of dissolution medium, then
11	dried on the water bath at 70 °C, dissolved in the mobile phase (1 mL), filtered
12	through a 0.45 μm membrane, and analyzed by high performance liquid
13	chromatography (HPLC). The 200% calibration sample was serially diluted with
14	distilled water to generate the other calibration samples, whose gross concentrations
15	were 25%, 50%, 75%, 100%, 125% and 150%.
16	For each batch of sample solution, 20 FZHY-P were accurately weighted,
17	equivalent to 2.00 times of FZHY-P weight (\pm 0.1 mg) . FZHY-P sample solutions were
18	collected at 10, 30, 60, 120 and 240 min throughout the experiment, and all were
19	prepared and analyzed as described above. Following sample removal, the
20	appropriate media (5 mL) was immediately replenished.
21	2.3 Instruments and chromatographic conditions
22	An Agilent Technologies 1200 HPLC system equipped with Agilent G1329A

1	automatic temperature control autosampler and Agilent G1314B variable wavelength
2	detector were used. Samples were separated using a Kromasil C18 column (250 mm
3	\times 4.6 mm, 5 μm). The mobile phase was composed of methanol (A), acetonitrile (B)
4	and water containing 0.2% H_3PO_4 (C) and was delivered at a flow rate of 0.8 mL \cdot min ⁻¹ .
5	A gradient program was used as follows: the initial elution condition at 0 min was
6	A:B:C (0.2% : 0.3% : 99.5% , v/v) , then linearly changed to A:B:C (1.2% : 1.8% :
7	97.0% , v/v) at 30 min, A:B:C (2.0% : 3.0% : 95% , v/v) at 31 min, A:B:C (4.0% : 6.0% :
8	90% , v/v) at 55 min, A:B:C (12.0% : 18.0% : 70.0% , v/v) at 90 min, A:B:C (20% : 30% :
9	50% , v/v) at 140 min, A:B:C (0.2% : 0.3% : 99.5% , v/v) at 150 min. The detection
10	wavelength was 280 nm and the column temperature was set at 30 °C.
11	Validation was performed by establishing precision, stability and repeatability
12	for all characteristic peaks. Chromatographic profiles were obtained for mixed
13	standards of protocatechuic aldehyde, salvianolic acid B, TSA, TG and the reference
14	solutions using the chromatographic conditions described above.
15	Intra-day precision tests were performed by analyzing FZHY-C reference
16	solutions during a single day and inter-day precision tests during 3 days. For
17	precision tests, reference solutions were analyzed six times continuously. For
18	repeatability tests, six independent reference solutions were prepared as described
19	above. For stability tests, six independent reference solutions were prepared and
20	analyzed at 0 h, 3 h, 6 h, 9 h, 12 h and 24 h following storage at −4° C.
21	2.4 Establishment of the evaluation method for the synchrony of drug release
22	based on the mathematic sets

1	A drug release set is defined as the assembly of chemical compounds dissolved
2	in a medium from the TCM preparation. ¹⁸ It can also be considered as the significant
3	part of the basis for pharmacodynamics.
4	The standard chromatographic fingerprint is defined as the characteristic
5	fingerprint of a drug release set for full drug release of an original preparation at the
6	specified dose. When the reference and sample preparations display similar release
7	behavior, it is defined as synchrony. Conversely, dissimilar release behavior is defined
8	as asynchrony.
9	When FZHY-C was fully released , the drug release set was comprised of several
10	components (corresponding subscript symbols as "n"), so the peak area of drug
11	release set could be described as {A _{s,1} , A _{s,2} ,, A _{s,(n-1)} , A _{s, n} }. Supposing the number
12	of FZHY-P which consists of different drug release units was defined as "p" and the
13	sampling time point of its in vitro release was defined as "t", then the peak area of
14	each component obtained from the "p" th sample can be expressed as $A_{p, n, t}$.
15	2.4.1 Establishment of standard curve of reference preparation
16	The reference preparation was "fully" dissolved as the calibration sample and it
17	was diluted to a series of concentrations, then the standard curve was constructed
18	by plotting peak area ($A_{s,n}$) against time on the x and y axes respectively. The
19	concentration of each component of sample preparation was obtained as followed.
20	Firstly, the $A_{\text{p,n,t}}\;$ of sample preparation were inputted into the standard curve
21	regression equation, then the relative release amount (C_t) of the drug release set
22	was be obtained according to the dilution ratio.

1	2.4.2 Accumulative release of the drug release set
2	After obtaining the relative release amount (C_t) of each characteristic peak in the
3	drug release set, the cumulative release degree (Q_t) could be calculated in
4	accordance with the formula below, and the curve for drug release could be fitted.
5	Finally, the release mechanism could be speculated.
6	$Q_t = \{ [C_t \times V_0 + \sum (C_{t-1} \times V_i)] / M \} \times 100\%$
7	In the equation, "t" represented sampling time point(h), " Q_t " represented
8	cumulative release degree (%) at sampling time point, " C_t " represented the
9	concentration of each characteristic peak (i.e. release quantity), " V_0 " represented
10	total volume of dissolution medium, "Vi" represented sampling volume, and "M"
11	represented release quantity of reference preparation (i.e., the product of the
12	reference " C_t " and " V_0 ").
12 13	reference " C_t " and " V_0 "). 2.4.3 The synchronous characterization of the drug release set
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1	$Lim\sum(Q_{p,n,i} - Q_{s,n}) = 0$. Hence, the absolute value of the difference was defined as
2	"Error" and was shown as " Q_{ε} ". The error of each component obtained from the "p" th
3	sample could be expressed as $Q_{\varepsilon(p,n,t)}$ t. $Q_{\varepsilon(p,n,t)}$ reflected the error between reference
4	preparation and samples. In order to reduce the calculation error, and thus the
5	synchrony of each compound in the drug release set could be evaluated by "relative
6	error" ($\varepsilon_{p,n,t}$), i.e. $\varepsilon_{p,n,t} = Q_{\varepsilon(p,n,t)} / Q_{s,n}$.
7	The error curve was obtained by plotting sample points "t" on the x-axis
8	against the relative error " ε " on the y-axis. The parameter $k_{p,n}$ (the slope of the error
9	curve) reflected the change of curve and was defined as the asynchronous coefficient
10	"k". The "k" for the reference and sample preparations was tested by a rank sum test
11	and the difference of the overall location and shape distribution was estimated
12	respectively. When no difference was observed, we could conclude that the
13	complete release process was similar or equal between the sample and reference
14	preparations.
15	2.5 Animals, Drug Administration, Sampling
16	Eighty-five white Sprague-Dawley (SD) rats were obtained commercially from
17	Chinese Academy of Shanghai Experimental Animal Center [Shanghai, China,
18	certification NO: SCXK (Shanghai): 2007-0005]. They were kept in an environmentally
19	controlled breeding room in Experimental Animal Center of Shanghai Traditional
20	Chinese Medicine University (Shanghai, China) for one week before starting the
21	experiments and fed with standard laboratory food and water ad libitum. The animal
22	facilities and protocols were approved by the Institutional Animal Care and Use

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1	Committee, Shanghai University of TCM. All procedures were conducted in
2	accordance with the Guide for the Care and Use of Laboratory Animal (The National
3	Academies Press, revised edition 2010).
4	Rats were divided into six groups as followed : normal group (n=10), model
5	group (n=19), the original preparation FZHY-C group (n=14), the high dose of FZHY-P
6	group (n=14), the medium dose of FZHY-P group (n=14) and the low dose of FZHY-P
7	group (n=14). The rats of model group were intraperitoneal injected with 2 mL·kg $^{-1}$
8	(weight) DMN (0.5%). The period of injection was lasted in 4 weeks. The first
9	injection was taken in 2/3 of the full dose. The injection was taken once a day. It had
10	the rest of 4 days after the continuous injection of 3 days. This period of injection
11	was total 3 weeks. In the 4^{th} week, the injection was taken 1/2 the amount of the full
12	dose in 1 $^{\rm st}$ day and 3 $^{\rm th}$ day of 4 $^{\rm th}$ week and was stopped 1 time during the period.
13	Then after 3 days, the injection was also taken the full dose one time. The normal
14	group rats were also intraperitoneal injected with the same amount of sodium
15	chloride injection as model group rats at the same time of model group.
16	Each group were orally given with the corresponding dose of an aqueous
17	solution with 10 mL·kg ⁻¹ (weight). The dose of FZHY-C group was 6.318 g·kg ⁻¹ (weight).
18	The dose of the FZHY-P was shown in Table 1S. Each group was given once a day and
19	lasted 4 weeks. The normal group and the model group were given the same dose of
20	drinking water.
21	After the end of the experiment, the rats were fasted, but not limited water, and

22 they were weighed after 24 hours. Then rats were sacrificed, and the livers and

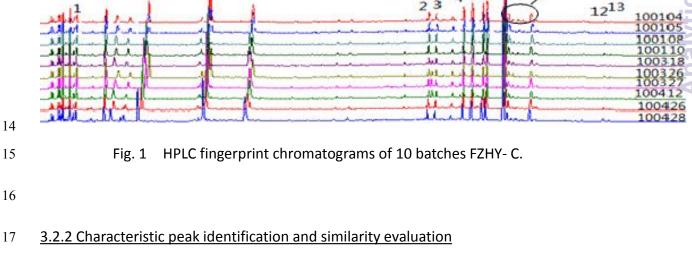
1	spleens were removed to weigh. And then about two 1.0 cm $\times 0.8$ cm $\times 0.3$ cm size
2	liver tissue was taken and fixed in neutral buffered formalin (10%).
3	2.6 Biomedical study
4	Hydroxyproline(Hyp) content of the liver tissue was determined by Jamall
5	methods. ³⁰ Microfuge lite was purchased from Beckman Company. Vortex mixer
6	(XW-80A) was obtained commercially by the Haimen Lindberg Instrument
7	Manufacturing Co., Ltd. M5 Multifunctional microplate reader was bought by
8	American Molecular Devices Company.
9	Approximately 100 mg of liver tissue was weighed, dried water out with a filter
10	paper, then it was placed in a homogenate tube, added 1.5mL amount of saline in
11	homogenizer, homogenized and transferred to ampoules. Homogenate tube was
12	flushed by 1 mL saline. The ampoule was added 2.5 mL HCl before sealed . Then the
13	sealed ampoule was put into the oven at 105 $^{ m o}$ C to hydrolyze for 20 hours. Then the
14	solution of hydrolyzate was filtered, 100 μL amount of which was plused into a
15	new homogenate tube. Each specimen was duplicated twice and each one was dried
16	in the oven at 40 °C.
17	0.2 mL amount of the chloramine-T working fluid and 1.2mL of the 50%
18	isopropanol were added to the homogenate tube. Then it was stayed at 25 °C room
19	temperature for 10 min. 1 mL of the ER working fluid was next added to it, i.e. the
20	Euclidean liquid [Ehrlich's reagent solution, 25% (w / v) dimethylamino benzaldehyde
21	and 27.3% (v / v) perchloric acid solution in isopropanol]. After mixed with shaking, it

22 was set at 50 °C warm bath for 90 min. OD value was measured at the wavelength of

23 558 nm after distilled water was used to zero. The OD value of each tube and its

1	duplicated tube were taken the mean, Hyp content was calculated based on the
2	standard curve, and corrected by precise liver tissue wet weight .
3	Standard curve was drawn as followed Table 2S. OD value was measured at the
4	wavelength of 558 nm after distilled water was used to zero. Standard curve was
5	drawn according to the standard concentration.
6	2.7 Statistical analysis
7	Standard chromatographic fingerprint was generated by using the software
8	"Similarity evaluation system for chromatographic fingerprint of TCM" (2004, A
9	Chinese Pharmacopoeia Commission). Differences between parameters for each
10	group of drug release were tested by rank sum test by using SPSS 18.0 software for
11	Windows (MapInfo Corporation, Troy, NY, USA).
12	The data was expressed as means \pm SD in pharmacodynamic evaluation. One-way
13	analysis of variance (ANOVA) and further LSD-t multiple comparisons were also
14	employed by SPSS 18.0. A <i>P</i> -value below 0.05 or 0.01 was taken to indicate significant
15	difference between data means.
16	3. Results and discussion
17	3.1 Validation of chromatography method
18	The results showed that linear relationship of each characteristic peak in the
19	standard fingerprint had a good linear relationship. Regression equations and
20	correlation coefficients of characteristic peaks are shown in Table 1. Table 1 also
21	listed the results of precision, repeatability and stability tests. For intra-day precision
22	analysis, the data showed that the relative standard deviations (RSDs) of retention

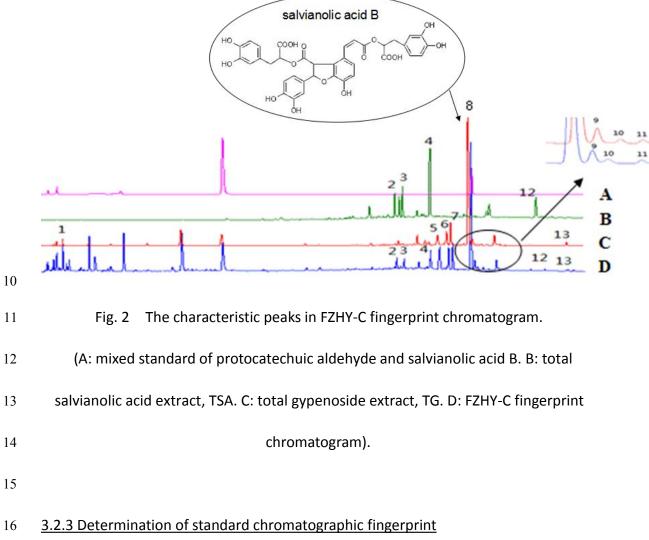
1	time and the peak area were less than 0.01% and 3.67% respectively. Meanwhile
2	those for inter-day variation were less than 0.04% and 3.80%, respectively. The
3	results for repeatability were less than 0.03% with respect to retention time and
4	5.93% for peak area. For the stability tests, the <i>RSDs</i> for retention time and peak area
5	were less than 0.03% and 4.15% respectively, indicating that the sample was stable
6	over 24 h. The validation data indicated that the analytical method was specific,
7	sensitive and stable. 3.2 Standard chromatographic fingerprint of the drug release set 3.2.1 Acquisition of standard chromatographic fingerprint Characteristic fingerprint chromatograms of 10 batches FZHY-C were shown in Fig.1. This data was generated from the reference standard fingerprint chromatogram by using the software, a similarity evaluation system for chromatographic fingerprint of TCM.
8	3.2 Standard chromatographic fingerprint of the drug release set
9	3.2.1 Acquisition of standard chromatographic fingerprint
10	Characteristic fingerprint chromatograms of 10 batches FZHY-C were shown in
11	Fig.1. This data was generated from the reference standard fingerprint
12	chromatogram by using the software, a similarity evaluation system for
13	chromatographic fingerprint of TCM.



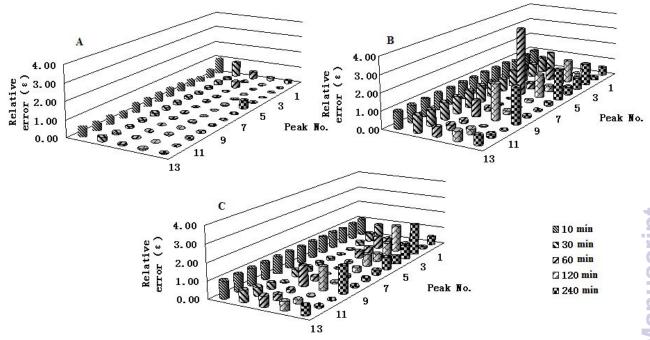
18 13 Characteristic peaks were identified, in which the fourth peak (94.33 min) was

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1	the reference peak (4 th s). Others had the following relative retention times: 0.07 min
2	(1 st), 0.96 min (2 nd), 0.98 min (3 rd), 1.00 min (4 th s), 1.02 min (5 th), 1.04 min (6 th), 1.06
3	min (7 th), 1.09 min (8 th), 1.10 min (9 th), 1.12 min (10 th), 1.14 min (11 th), 1.16 min (12 th)
4	and 1.23 min (13 th). From the Fig.2, based on comparisons of the mixed standards of
5	protocatechuic aldehyde and salvianolic acid B, total salvianolic acid extract (TSA),
6	total gypenoside extract (TG) and FZHY-C reference preparation, peak numbers 1, 5,
7	6, 7, 8, 9, 10, 11 and 13 were from TSA, peak numbers 2, 3, 4, 12 were from total
8	gypenoside, and peak 8 was salvianolic acid B. The average similarities of the 10
9	randomly-selected samples were >0.96.



1	The reference fingerprint "R" generated by the similarity evaluation system for
2	chromatographic fingerprint of TCM software was regarded as the standard
3	fingerprint chromatography of the evaluation method. The drug release set for the
4	standard chromatogram was formed by characteristic peak areas and could be
5	expressed as: $\{A_{s,1}, A_{s,2}, A_{s,3}, A_{s,4}, A_{s,5}, A_{s,6}, A_{s,7}, A_{s,8}, A_{s,9}, A_{s,10}, A_{s,11}, A_{s,12}, A_{s,13}\} = \{79.2, 10, 10, 10, 10, 10, 10, 10, 10, 10, 10$
6	181.6, 20.7, 87.8, 168.8, 351.1, 433.8, 3949.9, 154.6, 96.4, 50.1, 31.1, 81.3}.
7	3.3 Optimization of release unit type of FZHY-P by using the synchronous
8	evaluation method based on the mathematics set
9	For optimizing the release unit type of FZHY-P, the release unit composition was
10	designed. Sample 1 was contained of fat-soluble immediate release unit (A, 0.015 g)
11	and water-soluble sustained release unit (B, 0.285 g). Sample 2 was contained of A
12	(0.015 g) and water-soluble immediate release unit (C, 0.285 g). The peak area of the
13	drug release set in both standard and sample fingerprint chromatograms were
14	recorded in the release experiment described in sections 2.2. The chromatographic
15	conditions described in section 2.3. Each experiment was repeated three times. The
16	parameter (s) of the reference preparation and samples are shown in Table 2, relative
17	errors are shown in Fig. 3 and the statistical results are shown in Table 3.



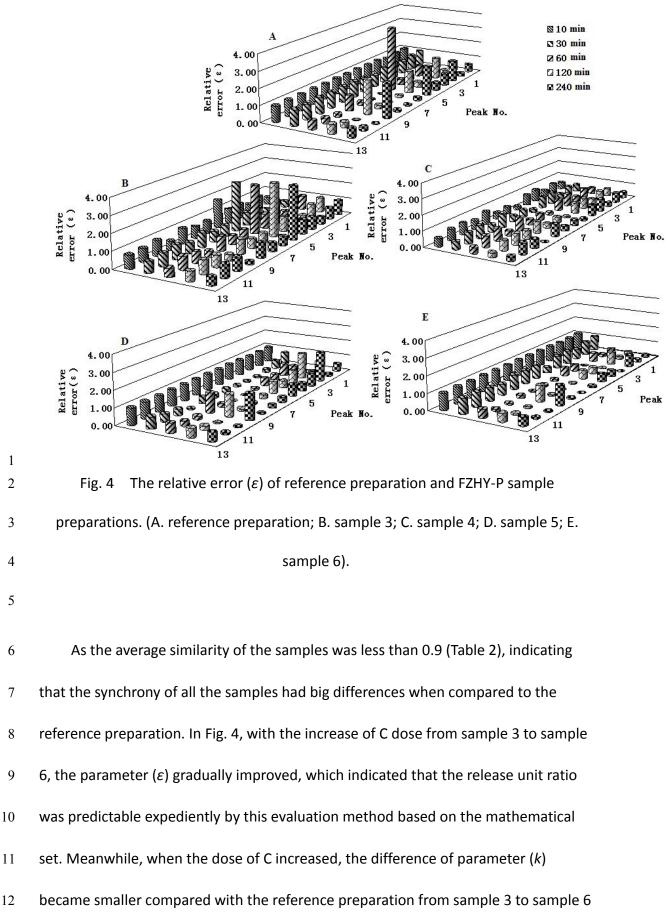
2 Fig. 3 The relative error (ϵ) of reference preparation and FZHY-P sample preparations. (A. reference preparation; B. sample 1; C. sample 2).

3

4

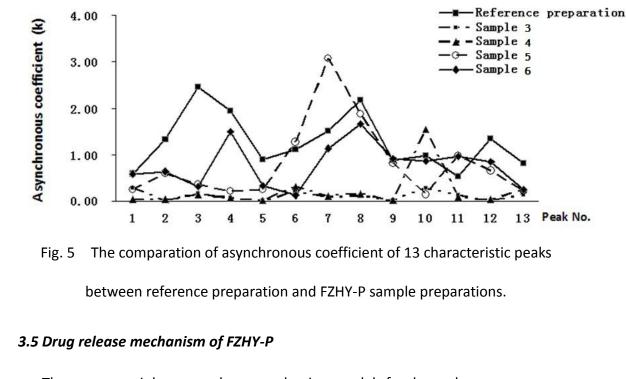
The average similarity (s) of both samples 1 and 2 were less than 0.9, as shown in 5 6 Table 2, indicating a big difference between the two samples and the reference preparation, especially within 60 minutes. One explanation could be the sustained 7 8 release effect which was caused by the sustained release pellets in sample 1. After 60 9 minutes, the similarity had become much closer to 1, which indicated that the release degree of samples was becoming more synchronous to the reference 10 11 preparation. We further observed that the peak numbers 3, 6 and 10 in sample 1 and 12 2 showed the biggest change in relative error (ε) by comparison with the reference 13 preparation (Fig. 3). It explained that the asynchrony was from the above three peaks 14 and next we should pay attention to adjusting the release unit ratio to improve synchrony. From Table 3, there were significant differences (P < 0.05) between 15

- 1 sample 1 and the reference preparation, but not sample 2.So the sample 2 was more
- 2 similar than sample 1 to the reference preparation.
- 3 In summary, this data showed that it was suitable that FZHY-P was composed of A
- 4 and C.
- 5 **3.4** Optimization of dose ratio of FZHY-P by using the synchronous evaluation
- 6 *method based on the mathematics set*
- 7 While some components such as peak number 6 and 10 were found to affect the
- 8 synchrony of the overall sample. Since these components were mostly derived from
- 9 TSA, TSA was one of water soluble active component, so we adjusted the
- 10 water-soluble immediate pellets dose to ensure that the pellets was similar to the
- 11 reference preparation in release character.
- 12 For optimizing the ratio dose of FZHY-P, all the A were 0.015 g. The C of Sample 3,
- 13 Sample 4, Sample 5, Sample 6 were 0.200 g, 0.250 g, 0.285 g, 0.400 g, respectively.
- 14 The similarity of the reference preparation and samples are shown in Table 2, relative
- 15 errors are shown in Fig. 4 and the statistical results are shown in Table 3.

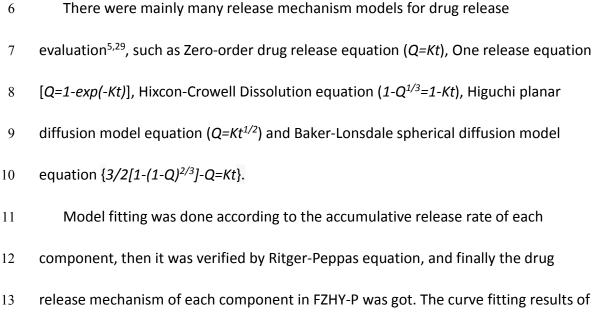


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1	(Table 3). Through the analysis of variance, sample 3 had significant difference ($P <$
2	0.05), and sample 6 was the most similar ($P = 0.42$). With the exception of sample 6,
3	the parameter (k) of the other samples all had a significant difference ($P < 0.05$, $P <$
4	0.01). So sample 6 was more similar than other samples to the reference
5	preparation.
6	In summary, this data showed that the dose ratio of the C and A were 0.4 : 0.015,
7	suggesting that the release character of FZHY-P was similar to the reference
8	preparation.
9	At the same time, we observed that statistical method could affect the accuracy
10	of the evaluation method. Initially we used the t-test ²² to assess the parameter(k)
11	between each sample and reference preparation. The results showed that samples 1,
12	2, and 3 had no distinct difference ($P > 0.01$). In contrast, by the rank sum test ³¹⁻³⁴ ,
13	samples 3, 4 and 5 were significantly different when compared to the reference
14	preparation ($P < 0.01$). We observed that the variation in the statistical methods was
15	due to the data normality and homogeneity of the variance test. It was assessed that
16	the data of the reference preparation and samples did not meet the requirements of
17	the t-test for data normality, therefore two kinds of statistical methods produced
18	different results. Unlike the t-test, the rank sum test was more suited to
19	non-normally distributed data and was not restricted by the overall distribution. Due
20	to the consideration of the overall position and peak shape, the Kolmogorov-
21	Smirnov z test was more creditable than Wilcoxon test used commonly ³⁵⁻³⁹ . Fig. 5
22	could demonstrated intuitively that the asynchronous coefficients of sample 6 was



1 closer to reference preparation in the shape and position than other samples.



- 14 sample 6 are shown in Table 4. The results showed that the model fitting of the
- 15 Hixcon-Crowell dissolution equation was the best for the vast majority of
- 16 components in FZHY-P except the third peak, which indicated that the drug release
- 17 mechanism of FZHY-P was mainly in corrosion process.

2 3

4

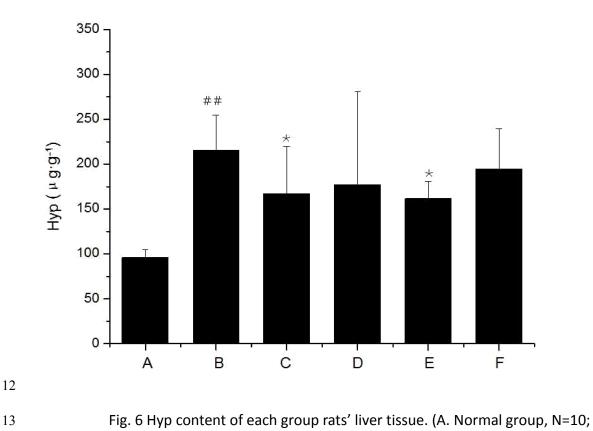
5

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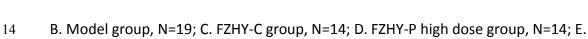
1	According to Ritger-Peppas ⁴⁰ , for ball type preparation, the meaning of exponent
2	of time 't' is as followed. Drug release is mainly in the Fickian diffusion when $n \leq 0.43$.
3	When 0.43 < n < 0.85, drug release is for non-Fickian diffusion (drug diffusion with
4	skeleton solutional phase coordination). When $n \ge 0.85$, drug release mode belongs
5	to mainly frame erosion. In this paper, the evaluation method based on the
6	mathematics set was used to represent the synchronicity of drug release on FZHY-P.
7	Through the validation of this model equations, it was shown that the third peak's n
8	value was greater than 0.43 and less than 0.85, which release mechanism belonged
9	to the diffusion dissolution synergy, meanwhile the remaining 12 characteristic peaks
10	of the release mechanism were mainly frame erosion.
11	3.6 Evaluation of efficacy against liver fibrosis
12	The fresh spleen of each group was observed (including the color, texture, surface
13	smoothness). The liver of normal group was red, tender, soft and had smooth
14	surface and the edge was sharp, and the spleen was dark red and medium texture.
15	However, the liver volume of model group was significantly reduced and the liver
16	was dark red, hard texture, rough surface and the edge of it was blunt. The spleen
17	volume of model group was also largely increased. Compared with the model group,
18	the livers of other drug groups (except the low dose of FZHY-P group) were better in
19	color and texture than model's, and spleens were also narrowed in some degree.
20	Compared with the normal group, the body weight, liver weight and liver
21	weight / weight of the model group were significantly lower ($P < 0.01$), spleen weight
22	and spleen weight /weight of model group increased significantly ($P < 0.01$).
23	Compared with the model group, the weight of original preparation FZHY-C group,

1	the high dose, medium dose of FZHY-P groups were higher than the model group
2	and the medium dose of FZHY-P group had significant higher($P < 0.05$). Compared
3	with the model group, the body weight and liver weight of the low dose of FZHY-P
4	group decreased, and its of medium dose of FZHY-P had statistically decreased
5	($P < 0.01$). The results were showed in Table 3S.
6	The Hydroxyproline content of each groups were also tested. Compared with
7	the same period of normal group, Hyp content of model group liver tissue was
8	significantly higher ($P < 0.01$). Compared with model group, Hyp content of the
9	original preparation FZHY-C group and medium dose of FZHY-P group liver tissues
10	had decreased significantly ($P < 0.05$). The results were showed in Table 3S and
11	Fig .6.









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1	FZHY-P medium dose group, N=14; F. FZHY-P lose dose group, N=14)
2	Compared with normal group, $^{^{\#\#}}P{<}0.01$; Compared with model group, $^{*P}{<}0.05$.
3	
4	The above results had shown that the medium dose of FZHY-P in the
5	intervention effect of DMN rat liver fibrosis was more effective than original
6	preparation. In this study, the content of Hyp in liver tissue, as the main
7	pharmacodynamic indicator could evaluated the reasonableness of the FZHY-P.
8	The results demonstrated the same effect as the original preparation, indicating that
9	the evaluation method for the synchrony of drug release based on the mathematics
10	set was reasonable and convenient in guiding its preparation .
11	4. Conclusions
12	Using the original preparation FZHY-C as the model, an integral evaluation
13	method for the synchrony of drug release based on the mathematics set for MCTCM
14	has been established. This method can not only be used in evaluating release
15	characterization of MCTCM by the parameter of asynchronous coefficient, but also
16	be used in adjusting the dose of release unit by the relative error parameter.
17	The results demonstrated that this evaluation method was feasible, stable and
18	reproducible. And the FZHY-P guided by this method showed release synchrony and
19	similar pharmacodynamics with the original preparation, and the drug release
20	mechanism was mainly frame erosion. Through this study, we could then evaluate
21	the release characteristic of MCTCM and it could apply an integral synchronous
22	evaluation method for multi-component dissolution / release.
23	Acknowledgments

1	The work was supported by grants from the National Natural Science
2	Foundation of China (Grant No.30801548), the Shanghai Municipal Education
3	Committee (Grant No. 12ZZ124), the Shanghai Science and Technology Committee
4	(Grant No. 08DZ1971100 and 12401900402) and the Shanghai Education
5	Commission Leading Academic Discipline Project (Grant No. J50302). Thanks to Xu
6	Lieming researcher for the contribution on anti-hepatic fibrosis pharmacodynamics
7	of FZHY-P.
8	
9	

inter-day and intra-day precision, repeatability and stability. 12

	6	Lieming r	esearcher fo	or the cont	ribution o	n anti-hep	oatic fibro	sis pharma	codynamic	S	·ipt
	7	of FZHY-P	».								SCL
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	,										D
	10	Table leg	ends								ote
	11	Table 1	Regression e	equations,	related co	oefficients	of each o	characterist	ic peak,		Cel
	12	intor day	and intra-da	av procisio	n ronosta	hility and	ctability				AC
	12	liitei-uay		ay precisio	n, repeata	Dinty and	Stability.				
Peak	Retention	Regressio		Retenti	ion time (% R	SD)			Peak Area	a(% <i>RSD</i>)	Ö
No.	time	n	Related	Intra-day	Inter-day	Repeata-	Stability	Intra-day	Inter-day	Repeata-	Stability
		equation	coefficient	precision	precision	bility		precision	precision	bility	Ť.
			(<i>r</i>)								G
1	6.445	Y =0.0855	0.978 0	0.00	0.01	0.00	0.33	1.89	3.58	5.93	4.15
		X -0.6864									ğ
2	91.002	Y =0.0332	0.999 3		0.00						C D
	91.002		0.5555	0.00	0.00	0.00	0.00	1.55	1.64	0.00	1.82
	91.002	X +0.0175		0.00		0.00	0.00	1.55		0.00	1.82
3	92.809	X +0.0175 Y =0.2501	0.984 7	0.00	0.00	0.00	0.00	1.55 2.28	1.64 2.66	0.00	Z
		X +0.0175 Y =0.2501 X +0.2804	0.984 7		0.04				2.66		1.82 2.05
3		X +0.0175 Y =0.2501 X +0.2804 Y =0.0640									Z
4	92.809	X +0.0175 Y =0.2501 X +0.2804 Y =0.0640 X -0.1081	0.984 7 0.999 6	0.00	0.04	0.00	0.02	2.28	2.66 2.50	1.60	2.05
	92.809	X +0.0175 Y =0.2501 X +0.2804 Y =0.0640 X -0.1081 Y =0.0264	0.984 7	0.00	0.04	0.00	0.02	2.28	2.66	1.60	2.05
4	92.809 94.811	X +0.0175 Y =0.2501 X +0.2804 Y =0.0640 X -0.1081 Y =0.0264 X +0.0184	0.984 7 0.999 6 1.000 0	0.00 0.00	0.04 0.04 0.02	0.00 0.00	0.02	2.28 2.36	2.66 2.50 0.61	1.60 4.39	2.05 00 0.71
4	92.809 94.811	X +0.0175 Y =0.2501 X +0.2804 Y =0.0640 X -0.1081 Y =0.0264 X +0.0184 Y =0.0962	0.984 7 0.999 6	0.00 0.00	0.04	0.00 0.00	0.02	2.28 2.36	2.66 2.50	1.60 4.39	2.05 00 0.71
4 5 6	92.809 94.811 96.798	X +0.0175 Y =0.2501 X +0.2804 Y =0.0640 X -0.1081 Y =0.0264 X +0.0184 Y =0.0962 X -0.3394	0.984 7 0.999 6 1.000 0 0.999 2	0.00 0.00 0.00	0.04 0.04 0.02 0.00	0.00 0.00 0.00	0.02 0.02 0.04	2.28 2.36 0.44	2.66 2.50 0.61 3.29	1.60 4.39 2.72	2.05 00 0.71 1.41
4	92.809 94.811 96.798	X +0.0175 Y =0.2501 X +0.2804 Y =0.0640 X -0.1081 Y =0.0264 X +0.0184 Y =0.0962 X -0.3394 Y =0.0129	0.984 7 0.999 6 1.000 0	0.00 0.00 0.00	0.04 0.04 0.02	0.00 0.00 0.00	0.02 0.02 0.04	2.28 2.36 0.44	2.66 2.50 0.61	1.60 4.39 2.72	2.05 00 0.71 1.41
4 5 6 7	92.809 94.811 96.798 98.659	X +0.0175 Y =0.2501 X +0.2804 Y =0.0640 X -0.1081 Y =0.0264 X +0.0184 Y =0.0962 X -0.3394 Y =0.0129 X +0.0627	0.984 7 0.999 6 1.000 0 0.999 2 1.000 0	0.00 0.00 0.00 0.00	0.04 0.04 0.02 0.00 0.00	0.00 0.00 0.00 0.00	0.02 0.02 0.04 0.03	2.28 2.36 0.44 3.11	2.66 2.50 0.61 3.29 0.70	1.60 4.39 2.72 2.49	2.05 000 0.71 1.41 0.54
4 5 6	92.809 94.811 96.798 98.659	X +0.0175 Y =0.2501 X +0.2804 Y =0.0640 X -0.1081 Y =0.0264 X +0.0184 Y =0.0962 X -0.3394 Y =0.0129	0.984 7 0.999 6 1.000 0 0.999 2	0.00 0.00 0.00 0.00	0.04 0.04 0.02 0.00	0.00 0.00 0.00 0.00	0.02 0.02 0.04 0.03	2.28 2.36 0.44 3.11	2.66 2.50 0.61 3.29	1.60 4.39 2.72 2.49	2.05 000 0.71 1.41 0.54

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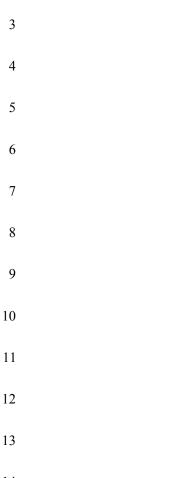
9	104.761	Y =0.0360	1.000 0	0.00	0.00	0.00	0.03	0.36	0.34	2.33	2.22
	104.701	X +0.0038		0.00		0.00	0.03	0.50		2.55	2.22
10	106.601	Y =0.0674	0.997 4	0.01	0.04	0.03	0.02	0.74	0.89	3.48	2.52
	100.001	X -0.3836		0.01		0.05	0.02	0.74		5.40	2.52
11	108.017	Y =0.1186	1.000 0	0.00	0.02	0.00	0.04	1.24	1.58	2.13	1.82
		X +0.0020		0.00		0.00	0.04	1.24		2.15	1.02
12	110.016	Y =0.2075	0.999 7	0.00	0.02	0.00	0.04	3.15	3.68	0.96	0.76
	110.010	X -0.0666				0.00	0.04	5.15		0.90	0.70
13	116.826	Y =0.0731	0.999 8	0.00	0.03	0.00	0.05	3.67	3.80	1.66	3.90
	110.820	X +0.0845		0.00		0.00	0.05	5.07		1.00	5.90

1

Sample No.				S		
-	10min	30min	60min	120min	240min	AVERAGE
Reference	0.872	0.926	0.951	0.974	0.976	0.940
Sample 1	0.105	0.402	0.936	0.953	0.943	0.668
Sample 2	0.000	0.906	0.937	0.979	0.978	0.760
Sample 3	0.504	0.697	0.783	0.851	0.900	0.747 *
Sample 4	0.650	0.794	0.887	0.937	0.952	0.844
Sample 5	0.000	0.906	0.937	0.979	0.978	0.760
Sample 6	0.000	0.916	0.965	0.972	0.986	0.768

1 Table 2 Similarity (s) of release of samples and reference preparation.

2 Compare to the reference preparation, *p < 0.05.



15

16

- 1 Table 3 The statistical results of asynchronous coefficient (*k*) of FZHY-P and reference
- 2 preparation.

				k			
No.	Reference preparation	Sample 1	Sample 2	Sample 3	Sample 4	Sample 5	Sample 6
1	0.579	0.248	0.232	0.269	0.032	0.232	0.581
2	1.330	0.564	0.594	0.021	0.033	0.594	0.627
3	2.462	0.173	0.365	0.148	0.139	0.365	0.303
4	1.934	0.134	0.1984	0.051	0.065	0.198	1.499
5	0.900	0.301	0.239	0.013	0.010	0.239	0.318
6	1.117	0.247	1.262	0.183	0.307	1.262	0.126
7	1.508	1.097	3.078	0.083	0.107	3.078	1.126
8	2.184	1.074	1.876	0.101	0.156	1.876	1.661
9	0.879	0.680	0.800	0.020	0.024	0.800	0.917
10	0.982	0.275	0.1382	0.269	1.539	0.138	0.864
11	0.534	1.397	0.972	0.119	0.081	0.972	0.965
12	1.333	0.519	0.660	0.012	0.032	0.660	0.833
13	0.800	0.211	0.210	0.140	0.232	0.210	0.239
K-S Z		1.569*	1.177	2.550**	2.353**	1.177*	0.981

3 Compare to the reference preparation, **P* < 0.05

4 Compare to the reference preparation, ***P* <0.01

- 5
- 6
- 7
- 8
- 9
- 10
- 11
- 12
- 13

Peak NO.	Model	Fitting equation	Correlation coefficient(<i>r)</i>
1	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-0.6395 <i>t</i>	0.977 1
2	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-0.4945 <i>t</i>	0.989 6
3	Higuchi Diffusion equation	$Q=0.1039t^{1/2}$	0.955 3
4	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-0.2306 <i>t</i>	0.984 2
5	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-4.6473 <i>t</i>	0.941 1
6	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-1.4185 <i>t</i>	0.948 1
7	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-0.1969 <i>t</i>	0.967 6
8	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-0.8478 <i>t</i>	0.970 8
9	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-0.5575 <i>t</i>	0.980 3
10	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-1.5170 <i>t</i>	0.996 4
11	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-0.6989 <i>t</i>	0.990 2
12	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-0.8494 <i>t</i>	0.974 0
13	Hixcon-Crowell Dissolution equation	(1-Q) ^{1/3} =1-0.4785 <i>t</i>	0.994 3

1	Table 4	The drug release	curve fitting res	ults of sample 6
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