

Article

The Comprehensive Evaluation of Safflowers in Different Producing Areas by Combined Analysis of Color, Chemical Compounds, and Biological Activity

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Abstract: In the present study, a new strategy including the combination of external appearance, chemical detection, and biological analysis was proposed for the comprehensive evaluation of safflowers in different producing areas. Firstly, 40 batches of safflower samples were classified into class I and II based on color measurements and K-means clustering analysis. Secondly, a rapid and sensitive analytical method was developed for simultaneous quantification of 16 chromaticity-related characteristic components (including characteristic components hydroxysafflor yellow A, anhydrosafflor yellow B, safflomin C, and another 13 flavonoid glycosides) in safflowers by ultra-performance liquid chromatography coupled with triple-quadrupole linear ion-trap tandem mass spectrometry (UPLC-QTRAP[®]/MS²). The results of the quantification indicate that hydroxysafflor yellow A, anhydrosafflor yellow B, kaempferol, guercetin, and safflomin C had significant differences between the two types of safflower, and class I of safflower had a higher content of hydroxysafflor yellow A, anhydrosafflor yellow B, and safflomin C as the main anti-thrombotic components in safflower. Thirdly, chemometrics methods were employed to illustrate the relationship in multivariate data of color measurements and chromaticity-related characteristic components. As a result, kaempferol-3-O-rutinoside and 6-hydroxykaempferol-3-O-β-D-glucoside were strongly associated with the color indicators. Finally, anti-thrombotic analysis was used to evaluate activity and verify the suitability of the classification basis of safflower based on the color measurements. It was shown that brighter, redder, yellower, more orange-yellow, and more vivid safflowers divided into class I had a higher content of characteristic components and better anti-thrombotic activity. In summary, the presented strategy has potential for quality evaluation of other flower medicinal materials.

Keywords: safflower; color; UPLC-MS²; chemometrics; quality evaluation



1. Introduction

Carthamus tinctorius L. (safflower), one of the oldest oil crops and major *Carthamus* species distributed in all of the world, has been applied in cut flowers, dyeing cosmetics, and coloring and flavoring foods [1,2], and safflower is a famous gynecological herbal medicine (Carthami Flos) in China, Korea, Japan, and other countries [3,4]. Modern pharmacological experiments show that safflower possesses a wide range of biological activities, such as dilating coronary arteries, improving myocardial ischemia, and regulating the immune system, as well as anticoagulation, anti-thrombosis, anti-oxidation properties [4–6]. In recent years, due to the medicinal value and health care value, more and more researchers have focused on investigating the active compounds, evaluating quality, and developing safflower products.

Recently, the safety and quality evaluation of safflower has been challenged, because it is often the target of artificial coloration and adulteration. Safflowers from different cultivars have different colors [7,8], such as white, yellow, orange, red, etc. The color of the total bloom stage turns from yellow to red gradually. Shades of orange, yellow, and red flowers are most common in early bloom, but post-bloom colors are darker. Additionally, different harvest times of the safflower cause different colors. The color is still yellow at the early harvest time, and the color turns purple and black at the late harvest time [9]. The light intensity can affect the color of the safflower and cause color changes as well [10]. The color of a safflower is usually related to the content of effective components, and it is an important factor to determine the quality of the safflower. Recent research has discovered that safflowers with different colors have great differences in the content of some chemical components. Xu et al. [11] discovered that when the content of hydroxysafflor yellow A (HSYA) is low, the safflower tends to be less red and darker. Tu et al. [12] indicated that the orange and white flower has a high content of HSYA and kaempferol-3-O-β-D-glucoside, respectively. Therefore, the color measurement is an important external appearance index of quality evaluation for raw and processed safflower, for determinations of conformity of safflower quality to certain specifications, or for the evaluation of quality changes as a result of safflower processing, storage, or other factors.

Additionally, in China, safflowers have been planted in Xinjiang, Yunnan, Gansu, Sichuan, and other places [13]. It is generally considered that plants grown in different areas have different contents and types of chemical markers, which are usually selected as important factors to study the quality evaluation and genuine producing areas of many functional foods and/or herbs, such as ginkgo seeds [14], *Ginkgo biloba* leaves [15], and *Ziziphus jujube* [16]. Currently, more than 200 compounds have been isolated and identified from safflower, including quinochalones, flavonoids, alkaloids, polyacetylene, and aromatic glucosides, among which quinochalcones and flavonoids are commonly considered as major bioactive and characteristic components associated with the therapeutic effects of safflower. Hence, the characteristic components (or chemical markers) might be a chemically important index to study the quality evaluation and genuine producing areas of safflower.

Nowadays, more and more studies indicate that quality evaluation and genuine producing areas of traditional Chinese medicines (TCM) or Chinese herbal medicine takes place via system engineering, which requires a lot of external appearances, and chemical and biological researches. However, current pharmacopoeias (such as Chinese Pharmacopoeia, U.S. Pharmacopoeia, and European Pharmacopoeia, etc.) are chemical markers-based, rather than bioactivity-oriented. Hence, to achieve the purpose of holistic quality evaluation of safflower, in this work, a new strategy based on a combination of the methods of external appearance, chemical detection, and biological analysis is proposed for the comprehensive analysis and evaluation of safflower samples from different producing areas (Figure 1). First, different safflower samples were classified into different types based on the results of external appearance of color measurements and K-means clustering analysis. Second, chromaticity-related characteristic components were comprehensive analyzed using UPLC-QTRAP[®]/MS² in different types

of safflower based on the above external appearance of color measurements. Third, chemometrics methods were employed to illustrate the relationship in multivariate data of color measurements and chromaticity-related characteristic components. Finally, anti-thrombotic analysis was used to evaluate the main biological activities and verify the suitability of the classification basis of different types of safflower based on the color measurements. By these efforts, the external, chemical, and biological evidence was obtained to unveil the quality of the different sources of safflower, which is beneficial to its holistic and more scientific quality control.



Figure 1. Technology roadmap of this study.

2. Results and Discussion

2.1. Color Measurements and Classification of Safflower

In the beginning, PCA of 40 samples was based on the content of characteristic components, according to the traditional classification method. It was found that the safflower samples from different regions could not be distinguished well. According to the previous literature, there was no direct evidence illustrating that safflower from different regions were different. According to the results, there was no significant difference in the content of characteristic components in safflowers from different regions. However, a significant difference in the content of characteristic components was presented in safflowers from the same region [17]. Additionally, from the previous reports, safflowers from different regions were also not distinguished based on the contents of the primary and secondary metabolites in safflowers and the in vitro bioactivity [18]. As we all know, the quality of safflowers was evaluated by their important appearance of color. The color measurement was selected as an important external appearance index to evaluate the quality of some vegetables and fruits such as tomatoes [19,20], blood oranges [21], blueberry [22], and bread [23]. Therefore, color measurement might be an effective and convenient method to evaluate the quality control of safflower, which has not previously been reported.

K-means clustering aims to partition *n* observations into *k* clusters, in which each observation belongs to the cluster with the nearest mean, serving as a prototype of the cluster [24]. The clustering algorithm refers to dividing a bunch of unlabeled data into different categories through an iterative process. This method ensures that the same type of data have similar features so that each cluster generated is compact inside, but the categories are independent of each other. Thus, safflowers with similar color characteristics would be classified into one category. In the previous analysis, the number of clusters was chosen to be 3, 4, or 5, but there was no significant difference between the groups in color indicators. At the same time, the color of the sample could be observed by the naked eye as orange, orange–red, red, and dark red. In order to ensure that the categories were independent of each other, the number of clusters was determined to be 2. The results of color measurements and K-means clustering analysis are shown in Table 1. Two types of safflower show significant differences in these five indicators. Ranges of L^{*}, a^{*}, b^{*}, C^{*}, and h_{ab} values in class I were 39.500~53.473,

20.403~34.543, 41.053~55.563, 50.422~64.749, and 55.055~69.345, respectively. Ranges of L*, a*, b*, C*, and h_{ab} values in class II were 30.930~49.697, 19.630~31.780, 25.860~40.527, 36.017~51.216, and 45.889~62.154, respectively. The values of L*, a*, b*, C*, and h_{ab} in class I were higher than class II (p < 0.01). This suggests that safflowers classified into class I were redder, yellower, brighter, more orange–yellow, and more vivid to the eye than class II. A comparison of class I and class II is shown in Supplementary Figure S2.

No	L*	a*	b*	Chroma	Hue Angle	Classification	No	L*	a*	b*	Chroma	Hue Angle	Classification
S1	47.3	33.3	47.7	58.2	55.1	Ι	S21	33.6	28.4	35.5	45.5	51.4	II
S2	38.3	29.0	35.1	45.5	50.4	II	S22	42.0	29.6	36.5	47.0	51.0	Π
S 3	46.7	33.0	43.3	54.5	52.7	Ι	S23	47.5	34.3	47.0	58.1	53.8	Ι
S4	46.8	31.8	43.8	54.1	54.0	Ι	S24	39.7	32.1	44.6	55.0	54.2	Ι
S 5	48.1	33.2	55.6	64.7	59.1	Ι	S25	47.3	32.4	46.5	56.7	55.1	Ι
S6	48.3	31.6	47.4	56.9	56.3	Ι	S26	37.0	29.4	40.5	50.1	54.1	II
S 7	43.2	31.9	42.3	53.0	53.0	Ι	S27	36.0	27.4	29.0	39.8	46.6	II
S 8	46.1	31.4	43.1	53.3	53.9	Ι	S28	49.2	34.5	48.5	59.5	54.5	Ι
S9	44.6	31.1	38.6	49.5	51.1	II	S29	36.3	27.8	32.0	42.4	49.1	Π
S10	38.5	29.3	38.2	48.1	52.5	II	S30	45.7	32.7	43.5	54.5	53.1	Ι
S11	48.9	33.3	48.9	59.2	55.7	Ι	S31	30.9	25.1	25.9	36.0	45.9	Π
S12	48.9	33.5	48.4	58.8	55.3	Ι	S32	39.5	32.5	42.7	53.7	52.7	Ι
S13	48.9	32.7	46.7	57.0	55.0	Ι	S33	53.5	20.4	54.1	57.8	69.3	Ι
S14	48.8	33.6	48.0	58.5	55.0	Ι	S34	42.5	31.8	40.2	51.2	51.6	II
S15	46.8	32.7	44.9	55.6	53.9	Ι	S35	42.7	26.7	36.3	45.1	53.6	II
S16	40.4	31.1	42.5	52.6	53.9	Ι	S36	43.4	22.4	37.2	43.5	58.9	II
S17	40.4	29.6	38.0	48.1	52.1	II	S37	49.7	19.6	37.2	42.0	62.2	Π
S18	50.9	26.3	49.3	55.9	61.9	Ι	S38	44.2	29.0	39.9	49.3	54.0	Π
S19	43.2	27.5	37.6	46.5	53.8	II	S39	43.0	29.9	41.1	50.8	53.9	Ι
S20	35.9	27.0	29.0	39.6	47.0	II	S40	45.1	28.8	41.4	50.4	55.1	Ι

 Table 1. Color measurements and classification of safflower.

2.2. Analysis of Chromaticity-Related Characteristic Components from Safflower

2.2.1. Selection of Chromaticity-Related Characteristic Components from Safflower

From the many previous reports, it has been confirmed that HSYA has a strong correlation with the color in the safflower [11,12]. Additionally, recent researches have shown that quinochalcones and flavonoids have a certain correlation with safflower color [12,25,26]. Therefore, three quinochalcones (HSYA, SC, and ASYB) and 13 flavonoids were selected, including their chromaticity-related characteristic components, for quantitative analysis in the present research.

2.2.2. Optimization of UPLC-QTRAP®/MS² Conditions

This study optimized chromatographic conditions such as column, mobile phase, flow rate, and column temperature. The Thermo Scientific Hypersil GOLD ($3 \times 100 \text{ mm}$, $1.9 \mu\text{m}$) column and the Acquity UPLC BEH C₁₈ ($2.1 \times 100 \text{ mm}$, $1.7 \mu\text{m}$) column were chosen to investigate the separation of the target compounds. The result indicates that the Acquity UPLC BEH C₁₈ ($2.1 \times 100 \text{ mm}$, $1.7 \mu\text{m}$) column could achieve better separation of 16 compounds. The mobile phase of the previous test, including methano–water, acetonitrile–water, methanol–water with 0.1% formic acid, and acetonitrile–water with 0.1% formic acid, were all tested. Acetonitrile had a stronger elution ability than methanol, which shortened the analysis time and obtained a better peak shape; and 0.1% formic acid kept the mobile phase at a suitable pH. The flow rate was set at 0.40 mL/min, and the column temperature was set at 35 °C, which were good for separating target compounds.

To optimize the QTRAP[®]/MS² conditions, Q1 full scans were conducted under both positive and negative electrospray ionization (ESI) modes [27]. The Analyst 1.6 software automatically collected and optimized the optimal ion fragmentation, declustering potential (DP), entrance potential (EP), collision energy (CE), collision cell exit potential (CXP), and other conditions. Three quinochalcone glucosides (HSYA, SC, and ASYB) were more sensitive in the negative ion mode. The MS/MS product ions at *m*/z 491.12 of HSYA resulted from the sugar moieties lost C₄H₈O₄. SC lost C₁₁H₁₀O₄ by cleavages of ring

A, then lost $C_4H_8O_4$ and formed the product ion at m/z 286.9. ASYB formed the product ion at m/z 449.11 (the base peak), together with a complementary product ion at m/z 593.15 in low abundance [28]. Flavonol $[M + H]^+$ product ions dehydrated to $[M + H - H_2O]^+$, followed by two sequential losses of CO: $[M + H - H_2O - CO]^+$ and $[M + H - H_2O - 2CO]^+$. These losses of carbon monoxide were also observed directly from the protonated flavonoid $[M + H - CO]^+$ and $[M + H - 2CO]^+$, and the loss of CO and dehydration occurred in the C-ring and proved the flavonol structure of the C-ring of an unidentified flavonoid [15]. Flavones $[M + H]^+$ ions exhibited only the $[M + H - H_2O]^+$ and $[M + H_2O - CO]^+$ fragments [19]. The optimized mass spectrometry conditions are shown in Table 2, and the chromatograms of the respective compounds are shown in Supplementary Figure S2.

	Compound	t _R (min)	Ion mode	MRM Transition	DP	EP	CE	СХР
1	hydroxysafflor yellow A	3.04	[M – H] [–]	611.06 > 490.8	-105	-10	-36	-53
2	safflomin C	12.20	[M – H] [–]	613.125 > 286.9	-40	-10	-45	-10
3	anhydrosafflor yellow B	8.80	[M – H] [–]	1043.122 > 448.9	-5	-10	-40	-10
4	kaempferol	12.99	[M – H] [–]	284.89 > 117	-155	-10	-56	-13
5	kaempferol-3-O-glucoside	8.86	[M – H] [–]	447.109 > 283.8	-120	-10	-32	-10
6	kaempferol-3-O-rutinoside	8.40	[M – H] [–]	593.003 > 284.9	-120	-10	-44	-29
7	kaempferol-3-O-β-sophoroside	5.78	$[M + H]^{+}$	611.8 > 286.9	56	10	31	20
8	6-hydroxykaempferol	12.31	$[M + H]^{+}$	303.933 > 168.8	186	10	47	14
9	6-hydroxykaempferol-3-O-β-D-glucoside	5.66	$[M + H]^{+}$	466.271 > 304.1	66	10	21	32
10	6-hydroxykaempferol-3,6-di-O-β-D-glucoside	4.68	[M + Na] ⁺	649.805 > 347	106	10	45	34
11	6-hydroxykaempferol-3,6,7-tri-O-β-D-glucoside	2.70	$[M + H]^{+}$	789.891 > 303.9	61	10	41	34
12	quercetin	12.56	[M – H] [–]	300.859 > 150.8	-90	-10	-28	-17
13	rutin	6.41	[M – H] [–]	609.048 > 300	-150	-10	-52	-33
14	luteoloside	7.22	[M – H] [–]	447.045 > 284.8	-80	-10	-36	-31
15	apigenin	12.95	$[M + H]^{+}$	270.989 > 153.1	171	10	49	14
16	quercetin-3-O-β-D-glucoside	6.88	$[M - H]^{-}$	461.98 3 > 298.9	-115	-10	-32	-31

Table 2. The ion mode and parameters for MRM of 16 compounds.

2.2.3. Method Validation

The established method was verified by measuring the linear, intraday and interday precision, stability, repeatability, and recovery. The calibration functions obtained by plotting the peak area versus the concentration of the compound were linear, and the determination coefficient higher than 0.9935 for all compounds. The limit of detections (LOD) and limit of quantitation (LOQ) ranges of 16 compounds were 1.182~176.130 ng/mL and 8.240~352.260 ng/mL, respectively. Calibration curves, correlation coefficients, linearity ranges, LODs, and LOQs of the 16 compounds are shown in Supplementary Table S1. The results of precision, repeatability, stability, and recovery are shown in Supplementary Table S2. The results of the experiment indicate that the developed method was accurate and reliable in this study.

2.2.4. Application to the Analysis of Real Samples

The results of quantitative determination are shown in Table 3. The contents of 16 compounds in two categories of safflower were compared, just as in Supplementary Figure S4. Safflower had a higher content of HSYA, SC, ASYB, kaempferol-3-O-rutinoside, 6-hydroxykaempferol-3-O- β -D-glucoside, and 6-hydroxykaempferol-3,6,7-tri-O- β -D-glucoside in class I. The content of HSYA and ASYB showed great differences between these two types of safflower (p < 0.01). HSYA (3.619~19.278 mg/g) and ASYB (2.196~14.124 mg/g) were the two compounds with the highest content. The color of HSYA and ASYB powders was orange–yellow, so the higher content of these two ingredients made the value of b* higher, which further caused the safflower to be more orange–yellow. Kaempferol, quercetin, and SC also had significant differences.

		Compound Number														
No	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
S 1	12.542	0.423	8.601	0.172	0.072	0.475	0.013	0.000	0.309	0.208	1.014	0.125	0.183	0.006	0.035	0.140
S2	7.279	0.225	2.975	0.166	0.070	0.437	0.019	0.000	0.384	0.219	1.095	0.049	0.065	0.009	0.037	0.044
S 3	14.583	0.277	4.351	0.138	0.061	0.501	0.013	0.098	0.332	0.159	0.701	0.034	0.074	0.006	0.034	0.041
S4	15.525	0.336	6.179	0.149	0.080	0.629	0.018	0.091	0.372	0.234	0.998	0.039	0.080	0.005	0.034	0.030
S5	11.456	0.265	4.989	0.142	0.055	0.628	0.020	0.101	0.335	0.135	0.669	0.029	0.100	0.005	0.000	0.039
S6	12.316	0.239	4.482	0.129	0.065	0.536	0.023	0.097	0.361	0.180	0.826	0.023	0.068	0.006	0.032	0.017
S 7	11.803	0.256	4.361	0.151	0.065	0.503	0.025	0.099	0.411	0.180	0.984	0.028	0.056	0.004	0.000	0.025
S 8	13.968	0.286	5.310	0.141	0.085	0.605	0.017	0.000	0.375	0.226	0.998	0.045	0.116	0.006	0.034	0.053
S9	13.611	0.240	5.812	0.165	0.259	0.580	0.023	0.096	0.576	0.444	1.378	0.059	0.074	0.006	0.000	0.112
S10	8.548	0.183	2.575	0.247	0.060	0.429	0.011	0.000	0.215	0.119	0.599	0.097	0.090	0.006	0.037	0.038
S11	15.204	0.188	5.437	0.147	0.061	0.450	0.004	0.000	0.263	0.128	0.504	0.019	0.080	0.004	0.035	0.048
S12	19.278	0.279	7.429	0.165	0.085	0.605	0.007	0.000	0.391	0.152	0.908	0.025	0.119	0.006	0.033	0.086
S13	17.615	0.245	7.250	0.131	0.073	0.527	0.008	0.000	0.382	0.169	0.756	0.012	0.089	0.004	0.033	0.042
S14	15.912	0.283	6.935	0.151	0.079	0.568	0.006	0.000	0.332	0.135	0.661	0.019	0.101	0.005	0.000	0.064
S15	17.714	0.363	7.098	0.222	0.202	0.454	0.010	0.000	0.629	0.315	1.152	0.048	0.060	0.007	0.000	0.136
S16	18.580	0.303	6.679	0.197	0.143	1.070	0.018	0.091	0.485	0.181	1.480	0.074	0.208	0.033	0.035	0.132
S17	11.124	0.225	3.561	0.185	0.090	0.536	0.015	0.096	0.310	0.220	0.939	0.095	0.143	0.013	0.033	0.080
S18	13.795	0.582	10.157	0.180	0.121	0.631	0.010	0.093	0.555	0.330	1.170	0.059	0.111	0.007	0.033	0.122
S19	8.689	0.200	3.439	0.142	0.061	0.472	0.015	0.095	0.270	0.137	0.672	0.031	0.059	0.002	0.033	0.033
S20	6.605	0.171	2.215	0.236	0.056	0.357	0.011	0.096	0.195	0.093	0.664	0.051	0.049	0.004	0.036	0.023
S21	8.745	0.329	3.913	0.198	0.074	0.562	0.014	0.097	0.323	0.111	0.759	0.046	0.059	0.004	0.039	0.023
S22	10.158	0.273	4.006	0.202	0.062	0.626	0.009	0.091	0.244	0.112	0.709	0.049	0.086	0.006	0.037	0.034
S23	15.730	0.343	7.145	0.190	0.098	0.617	0.010	0.093	0.562	0.241	1.025	0.052	0.102	0.005	0.033	0.062
S24	8.232	0.150	2.559	0.184	0.038	0.438	0.007	0.000	0.158	0.094	0.397	0.038	0.061	0.004	0.036	0.023
S25	17.862	0.377	8.845	0.171	0.101	0.643	0.009	0.099	0.493	0.264	1.176	0.040	0.114	0.005	0.000	0.094
S26	8.295	0.299	4.204	0.169	0.089	0.574	0.019	0.095	0.421	0.226	1.229	0.050	0.075	0.005	0.037	0.037
S27	7.625	0.159	2.332	0.232	0.056	0.353	0.012	0.000	0.198	0.111	0.655	0.051	0.051	0.005	0.037	0.032
S28	17.452	0.238	7.022	0.162	0.074	0.562	0.004	0.000	0.442	0.153	0.745	0.028	0.104	0.007	0.032	0.049
S29	5.806	0.178	2.333	0.252	0.051	0.430	0.009	0.000	0.218	0.109	0.592	0.067	0.051	0.004	0.044	0.031
S30	15.171	0.306	7.633	0.200	0.111	0.752	0.015	0.097	0.446	0.199	1.162	0.065	0.122	0.004	0.033	0.099
S31	7.495	0.304	2.746	0.292	0.088	0.360	0.013	0.000	0.267	0.117	0.755	0.109	0.061	0.004	0.041	0.045
S32	7.678	0.175	2.196	0.188	0.056	0.465	0.010	0.000	0.238	0.118	0.572	0.043	0.063	0.012	0.039	0.020
S33	13.284	0.757	14.124	0.227	0.126	0.707	0.008	0.096	0.738	0.314	1.351	0.055	0.083	0.003	0.032	0.090
S34	9.598	0.288	4.077	0.207	0.079	0.631	0.019	0.094	0.327	0.184	0.982	0.053	0.078	0.003	0.036	0.039
S35	9.160	0.308	4.245	0.185	0.083	0.568	0.015	0.095	0.352	0.227	0.789	0.083	0.161	0.010	0.039	0.121
S36	5.855	0.301	4.242	0.394	0.488	1.157	0.013	0.094	0.698	0.504	0.907	0.417	0.352	0.024	0.039	0.609
S37	3.619	0.186	3.642	0.331	0.318	0.906	0.007	0.094	0.390	0.285	0.681	0.307	0.301	0.012	0.041	0.296
S38	14.954	0.370	7.559	0.182	0.124	0.500	0.009	0.000	0.517	0.185	1.146	0.028	0.057	0.011	0.000	0.071
S39	16.308	0.359	8.346	0.182	0.104	0.567	0.011	0.095	0.504	0.164	1.045	0.021	0.073	0.007	0.000	0.054
S40	16.456	0.441	9.403	0.180	0.135	0.513	0.011	0.092	0.561	0.198	1.276	0.032	0.062	0.011	0.033	0.011

Table 3. The content of 16 compounds in 40 batches of safflower (mg/g).

2.3. Chemometrics Classification of Safflower Samples from Different Production Areas Based on the Chromaticity-Related Characteristic Components

During recent years, there has been increasing interest in illuminating the intrinsic relationship of multivariate data which are composed of numerous variables measured from many samples by validated analytical methods. Chemometrics methods have been widely used for depicting the intrinsic similarities and differences of samples by comparing particular chemical and biological parameters in different samples. The central idea of unsupervised PCA is to reduce the dimensionality of a data set consisting of a large number of interrelated variables. PCA is a very useful classification technique and widely used in the field of analytical chemistry. OPLS-DA is one of the most popular and common supervised methods. OPLS-DA is increasing its uses related to different chemical issues and could be considered as a popular tool in multivariate analysis. In this study, PCA was performed to classify safflower samples from different production areas based on the contents of the 16 chromaticity-related characteristic components. The supervised method of OPLS-DA was utilized to further separate clusters and validate the classify model. The scatter plots of PCA and OPLS-DA (Figure 2) show that the two types of safflower can be distinguished well. Compounds with VIP values larger than 1 were considered to be more important in the classification than other components. The VIP value represents the contribution of the variable to the model, If the value is larger, the contribution is greater. The VIP values of HSYA, ASYB, kaempferol, quercetin, and SC were more than 1. Thus, these five compounds were considered to be closely related to color characteristics.



Figure 2. PCA (**A**), OPLS-DA (**B**), permutation test (**C**), and VIP value (**D**) based on the content of the 16 compounds in two categories.

More and more researches have pointed out that the color is associated with bioactive compounds content [29]. Pearson correlation matrix analysis was used to discover the correlation between color indicators and compounds' contents [30]. The results of the correlation analysis are presented in Figure 3. The red and blue part in the figure indicate a strong correlation (r > 0.5 or r < -0.5). As shown in Figure 3, L* had a strong positive association with HSYA (r = 0.597), ASYB (r = 0.707), and 6-hydroxykaempferol-3-O- β -D-glucoside (r = 0.506). Additionally, a* had a strong positive association with HSYA (r = 0.609). The other parameter of b* had a strong positive association with HSYA (r = 0.706) and ASYB (r = 0.664). Chroma had a strong positive association with HSYA (r = 0.706) and ASYB (r = 0.576), and a strong negative association with HSYA (r = 0.706) and ASYB (r = 0.576), and a strong positive association with HSYA (r = 0.706) and ASYB (r = 0.576), and a strong positive association with HSYA (r = 0.706) and ASYB (r = 0.576), and a strong negative association with Kaempferol (r = -0.621). Hue angle had a strong positive association with ASYB (r = 0.651), 6-hydroxykaempferol-3-O- β -D-glucoside (r = 0.578), SC (r = 0.619),



and kaempferol-3-O-rutinoside (r = 0.521). It is reasonable to speculate that if the contents of HSYA and ASYB were higher, the color of safflower would turn lighter and more yellow.

Figure 3. Correlation between color indicators and compounds content.

The contents of HSYA, ASYB, and SC were higher in class I, but kaempferol and quercetin had a lower content. Guo et al. found that overexpression of CtCHS1 increased the quinochalcone glucoside accumulation and decreased flavonol aglycone and glycoside contents in safflower, and this study was based on the comparison of the yellow and white line safflower [31]. It indicates that accumulation of quinochalcone glucosides would make safflowers turn yellow and red. HSYA, ASYB, and SC belong to the quinochalcone *C*-glycosides, and the contents of the above three quinochalcone *C*-glycosides were higher in class I, while two flavonol aglycones (kaempferol and quercetin) presented a lower content in class I of safflower. Different expressions of CtCHS1 caused the different content in the five compounds, which might lead to differences in the color of the safflower.

2.4. Anti-Thrombotic Evaluation of Two Types of Safflower in Zebrafish

In this study, anti-thrombotic analysis was employed to evaluate the main biological activities and verify the suitability of the classification basis of different types of safflower based on the color measurements.

After being treated with 1.5 μ mol/L PHZ for 24 h, thrombus in the caudal vein was obviously increased, but the RBCs decreased significantly (p < 0.01), just as shown in Figures 4B, 5B, 6A. The thrombus in the caudal vein was decreased after treatment with aspirin for 24 h, and the RBCs increased significantly (p < 0.01), as shown in Figures 4C, 5C and 6A. The thrombus in the caudal vein was decreased after treatment with various concentrations of class I and II of safflower; 100 μ g/mL of class I of safflower could increase RBCs significantly (p < 0.01) and 200 μ g/mL of class II of safflower could also increase RBCs significantly (p < 0.05). The therapeutic efficacies of various concentrations of class I of safflower were

18.88%, 19.19%, 33.61%, 41.32%, 46.86%, and 46.80%, and the concentration of EC_{50} was 96.03 µg/mL. The therapeutic efficacies of various concentrations of class II of safflower were 11.94%, 15.14%, 15.26%, 26.92%, 35.48%, and 35.88%, and the concentration of EC_{50} was 181.8 µg/mL.



Figure 4. Thrombus in the caudal vein of vehicle control group (A), model group (B), aspirin group (C), the group treated with 200 μ g/mL class I of safflower (D), and the group treated with 200 μ g/mL class II of safflower (E).



Figure 5. The heart red blood cells (RBCs) of the control group (**A**), model group (**B**), aspirin group (**C**), various concentrations of class I of safflower (25–600 µg/mL) (**D–I**), and various concentrations of class II of safflower (25–600 µg/mL) (**J–O**). The blue circle indicates the staining of erythrocytes in the heart.



Figure 6. The heart RBCs, the therapeutic efficacies, and the concentration of EC₅₀ in class I of safflower (**A–C**) and class II of safflower (**D–F**). Data are expressed as mean \pm SEM of three separate experiments (n = 6). ^{##} p < 0.001 versus vehicle control group; ** p < 0.01 and *** p < 0.001 versus model group.

2.5. The Relationship of Color Measurements, Chemical Detection, and Anti-Thrombotic Analysis of Safflower

Safflowers with different colors showed different contents of target compounds. The contents of HSYA, ASYB, kaempferol, quercetin, and SC affected the color of the safflower. HSYA, ASYB, and SC made the safflower brighter, redder, yellower, and more orange–yellow. In addition, 6-hydroxykaempferol-3-O- β -D-glucoside and kaempferol-3-O-rutinoside made the safflower more orange–yellow too. To compare the activity of two types of safflower, class I of safflower had a better anti-thrombotic activity. The color of class I of safflower was brighter, redder, yellower, more orange–yellow, and more vivid to the eye. These color features of class I of safflower led to the higher content of characteristic components (HSYA, ASYB, and SC) and better anti-thrombotic activity.

Safflower is considered as a medicinal plant for the treatment of cardiovascular diseases, lowering blood cholesterol and relieving the pain. Recently, it was reported that chalcones possess the potential to treat cardiovascular disease [32]. HSYA and ASYB have been confirmed to be the main active chalcones in safflower [33–35]. Combined with the results of this study, SC might also be a main active ingredient. These three quinochalcone C-glucosides not only related to the color, but also had an association with anti-thrombotic activity.

Color not only reflected the chemical composition of safflower, but also reflected its pharmacological activity. Safflower which was brighter, redder, yellower, more orange–yellow, and more vivid had a higher content of characteristic quinochalcone C-glucoside components (HSYA, ASYB, and SC) and better anti-thrombotic activity. Therefore, the combination method of color features, chemical detection, and biological analysis is an effective, rapid, and repeatable strategy for the comprehensive analysis and evaluation of different safflower samples.

3. Material and Methods

3.1. Chemicals and Reagents

Sixteen chemical standards, including hydroxysafflor yellow A (1), safflomin C (2), anhydrosafflor yellow B (ASYB, 3), kaempferol (4), kaempferol 3-*O*-glucoside (5), kaempferol-3-*O*-rutinoside (6), kaempferol-3-*O*-β-sophoroside (7), 6-hydroxykaempferol (8), 6-hydroxykaempferol-3-*O*-β-D-glucoside (9), 6-hydroxykaempferol-3,6-di-*O*-β-D-glucoside (10), 6-hydroxykaempferol-3,6,7-tri-*O*-β-D-glucoside (11), quercetin (12), rutin (13), luteoloside (14), apigenin (15), and quercetin-3-*O*-β-D-glucoside (16), were used in this study. Compounds 3, 4, 7, 8, and 10 were purchased from Shanghai Yuanye

Biotechnology Company (Shanghai, China). Compounds **5**, **6**, and **13** were bought from Chengdu Chroma-Biotechnology Company (Chengdu, China). Compounds **1**, **12**, **14**, **15**, and **16** were acquired from Nanjing Liangwei Biotechnology Company (Nanjing, China). Compounds **2**, **9**, and **11** were isolated from safflower by the authors [36,37]. Dimethyl sulfoxide (DMSO), phenylhydrazine (PHZ), and acetylsalicylic acid (aspirin) were bought from Aladdin Company (Shanghai, China). *O*-dianisidine was obtained from J&K Scientific Ltd. (Beijing, China). Methanol, acetonitrile, and formic acid were HPLC grade and purchased from Merck (Darmstadt, Germany). Ultrapure water was prepared by the Milli-Q water purification system (Millipore, Bedford, MA, USA).

3.2. Plant Materials

Forty batches of samples from different regions were identified as *Carthamus tinctorius* L. by Dr. Hui Yan at the Department of Pharmacognosy, Nanjing University of Chinese Medicine, Nanjing, China. The voucher specimens were deposited at the herbarium in Nanjing University of Chinese Medicine, China. All the samples were pulverized into homogeneous powders (50 mesh), then stored under dry conditions at room temperature. The detailed information of the samples is shown in Table 4.

No	Location	No	Location
S1	Hefei City, Anhui Province	S21	Huocheng County, Xinjiang Uygur Autonomous Region
S2	Jiuquan City, Gansu Province	S22	Jimsar County, Xinjiang Uygur Autonomous Region
S 3	Jinhe Village, Jiuquan City, Gansu Province	S23	Jimsar County, Xinjiang Uygur Autonomous Region
S4	Jinhe Village, Jiuquan City, Gansu Province	S24	Jimsar County, Xinjiang Uygur Autonomous Region
S 5	Nanqu Village, Yumen City, Gansu Province	S25	Jimsar County, Xinjiang Uygur Autonomous Region
S6	Nanqu Village, Yumen City, Gansu Province	S26	Jimsar County, Xinjiang Uygur Autonomous Region
S 7	Nanqu Village, Yumen City, Gansu Province	S27	Emin County, Tacheng City, Xinjiang Uygur Autonomous Region
S 8	Dongqu Village, Yumen City, Gansu Province	S28	Emin County, Tacheng City, Xinjiang Uygur Autonomous Region
S9	Yumen county, Yumen City, Gansu Province	S29	Yumin County, Tacheng City, Xinjiang Uygur Autonomous Region
S10	Luoyang City, Henan Province	S30	Urumqi City, Xinjiang Uygur Autonomous Region
S11	Qixia District, Nanjing City, Jiangsu Province	S31	Huocheng County, Xinjiang Uygur Autonomous Region
S12	Qixia District, Nanjing City, Jiangsu Province	S32	Yining County, Xinjiang Uygur Autonomous Region
S13	Qixia District, Nanjing City, Jiangsu Province	S33	Yutian County, Xinjiang Uygur Autonomous Region
S14	Qixia District, Nanjing City, Jiangsu Province	S34	Emin County, Tacheng City, Xinjiang Uygur Autonomous Region
S15	Yongxin Couty, Jian City, Jiangxi Province	S35	Lijiang City, Yunnan Province
S16	Jianyang City, Sichuan Province	S36	Yongsheng County, Yunnan Province
S17	Xichang City, Sichuan Province	S37	Yongsheng County, Yunnan Province
S18	Chabuchar County, Xinjiang Uygur Autonomous Region	S38	Suzhou District, Jiuquan City, Gansu Province
S19	Yutian County, Xinjiang Uygur Autonomous Region	S39	Suzhou District, Jiuquan City, Gansu Province
S20	Huocheng County, Xinjiang Uygur Autonomous Region	S40	Suzhou District, Jiuquan City, Gansu Province

Table 4. Forty batches of safflower samples from different regions.

3.3. Color Measurements

The color of the sample powder was measured by Konica Minolta Spectrophotometer CM-5 and each sample was analyzed in triplicates. L* value indicated lightness, from dark (0) to white (100), a lower value indicated darker color and a higher value indicated lighter color. The other CIE parameter of a* value indicated from green ($-a^*$) to red ($+a^*$), while b* value indicated from blue ($-b^*$) to yellow ($+b^*$) [38]. Chroma (C^*) was the quantitative colorfulness attribute, as it determined the difference degree in comparison to a grey color with the same lightness for each hue [21]. Hue angle (h_{ab}) was a parameter that defined the colors traditionally as pinkish, yellowish, and greenish [21]. High values of hue angle represented more reddish–orange color and low values indicated more reddish–blue color. CIE LAB color space is shown in Supplementary Figure S1. The equations for calculating C*_{ab} and hue angle (h_{ab}) are Equations (1)–(3) [38]:

Chroma
$$(C^*_{ab}) = \sqrt{(a^{*2} + b^{*2})}$$
 (1)

Hue angle
$$(h^{\circ}_{ab}) = \tan^{-1}(b^*/a^*)$$
 when $a^* \ge 0$ and $b^* \ge 0$ (2)

Hue angle
$$(h^{\circ}_{ab}) = 180 + \tan^{-1}(b^*/a^*)$$
 when $a^* < 0$ (3)

3.4. Quantitative Determination of 16 Compounds

3.4.1. Preparation of Sample Solution

Firstly, 0.5 g of each dry sample powder was weighed accurately into a 50 mL conical flask with stopper, and 20 mL of 50% methanol was added accurately to each conical flask. Then, after weighing the total weight, ultrasonic extraction (40 kHz) for 40 min at 30 °C was conducted, using the same solvent to replenish the loss weight during extraction. After centrifugation (13,000× *g*, 10 min), the supernatants were stored at 4 °C and filtered through 0.22 µm cellulose membrane filters prior to injection.

3.4.2. Preparation of Standard Solution

A mixed standard stock solution containing the above target compounds **1–16** was dissolved by 70% methanol. The initial concentration of the compounds **1–16** were 875.625 μ g/mL, 54.375 μ g/mL, 64.375 μ g/mL, 32.5 μ g/mL, 40.234 μ g/mL, 38.734 μ g/mL, 81.875 μ g/mL, 58.750 μ g/mL, 63.750 μ g/mL, 60 μ g/mL, 33.437 μ g/mL, 62.5 μ g/mL, 63.125 μ g/mL, 9.609 μ g/mL, 21.562 μ g/mL, and 45.089 μ g/mL, respectively. Then, the mixed standard stock solution was diluted with 70% methanol to a series of appropriate concentrations [39]. All the standard solutions were stored at 4 °C until use, and filtered through a 0.22 μ m cellulose membrane before injection.

3.4.3. UPLC-QTRAP[®]/MS² Conditions

Chromatographic separation was performed on a Waters ACQUITY UPLC system (Waters, Milford, MA, USA). An UPLC BEH C₁₈ column ($2.1 \times 100 \text{ mm}$, $1.7 \mu\text{m}$) was used for analyzing all the samples. The mobile phase included A (0.1% formic acid aqueous solution) and B (acetonitrile) (v/v). The gradient elution procedure was as follows: 10-20% B at 0-2 min, 20-30% B at 2-7 min, 30-80% B at 7–12 min, 80-100% B at 12–13 min, and 100-10% B at 13–14 min. The mobile phase was set at a flow rate of 0.40 mL/min, and the injection volume was 2 μ L. The column temperature was set at 35 °C and the sample temperature was 4 °C.

Mass spectrometry detection was performed with an AB SCIEX Triple Quad 6500 plus (AB SCIEX Corp., Massachusetts, USA) equipped with an electrospray ionization source (ESI). The ESI–MS were acquired in both positive and negative ion multiple reaction monitoring (MRM) modes, with the capillary voltage of 5 kV, the desolvation gas flow rate set to 1000 L/h at a temperature of 550 °C, the cone gas flow rate set at 50 L/h, and the source temperature at 150 °C. The cone voltage (CV) and collision energy (CE) [40] were set to match the MRM of each compound. The dwell time was automatically set by the MultiQuant software 3.0.2 (AB SCIEX Corp., Massachusetts, USA).

3.4.4. Validation of UPLC-QTRAP®/MS² Method

A series of concentrations of standard solutions was prepared for obtaining the calibration curve. The limit of detections (LODs) and the limit of quantitations (LOQs) of 16 compounds were acquired while the signal-to-noise ratios (S/N) were 3 and 10, respectively. The peak height divided by the background noise value was calculated as the S/N.

The 16 standard solutions with six replicates were investigated during a single day and three consecutive days, respectively, for the intra-day and inter-day precision, and the results were displayed as the RSDs of the peak area for each standard compounds. The repeatability was confirmed by six different sample solutions, which were prepared from the same sample **S26**. Then, the variations were expressed by RSDs. Meanwhile, the sample solutions mentioned above were stored at 4 °C, injected, and analyzed at 0, 2, 4, 6, 8, and 16 h for the evaluation of the stability.

The purpose of the recovery test was to evaluate the accuracy of the developed method. It was performed by adding the 16 standards at low (80% of known amounts), medium (same as known amounts), and high (120% of known amounts) levels into the representative sample **S26** solution.

The spiked samples were then extracted, processed, and quantified in accordance with the methods mentioned above. The spike recoveries were calculated using Equation (4):

Recovery = [(measured amount – original amount)/spiked amount]
$$\times 100\%$$
 (4)

3.5. Anti-Thrombotic of Two Types of Safflower in Zebrafish

Firstly, 1 mL of the sample solution was accurately prepared according to the above steps. Then, the solvent was evaporated and residue was dissolved in 25 mL of 0.1% DMSO medium [41]. The final concentration of the solution was 1000 μ g/mL, then diluted with water as required. **S28** and **S29** were selected as representative samples, which both showed different color and represented two types of safflower.

Zebrafish at 48 HPF (hours post fertilization) were bought from Nanjing EzeRinka Biotechnology Co., Ltd (Nanjing, China). Firstly, 20 zebrafish larvae were placed in a 24-well microplate and treated with 1.5 μ mol/L PHZ for 24 h [42]. Secondly, PHZ was washed out, and then different groups of zebrafish were respectively treated with the positive drug of aspirin (5.625 μ g/mL) and two types of safflower at different dosing concentrations (25, 50, 100, 200, 400, and 600 μ g/mL). These different doses were consistent with the experimental concentrations, and the concentration of 50% maximal effect (EC₅₀) was calculated at the same time. Larvae placed in 0.1% DMSO were considered to be the vehicle control. The morphology of zebrafish was evaluated after incubation for 24 h at 28 °C. Thirdly, 10 zebrafish of each group were stained with 1.0 mg/mL *O*-dianisidine dye liquor for 15 min and washed with DMSO three times [43]. Lastly, the specimens were placed on glass slides and observed by a fluorescent inverted microscope (Leica, Germany). The heart red blood cells (RBCs) were quantitatively analyzed by Image-Pro Plus 6.0, based on the staining intensity (SI) of erythrocytes in the heart. The anti-thrombotic effects of safflower and aspirin were evaluated calculated by the following formula [42] Equation (5).

Therapeutic efficacy (%) =
$$\frac{SI(drug) - SI(model)}{SI(control) - SI(model)} \times 100\%$$
 (5)

3.6. Statistical Analysis

Data of color measurements were imported to SPSS 24.0 software (SPSS, Chicago, USA) for K-Means clustering analysis. The data were expressed as mean \pm SEM, and two-tailed *t*-test and analysis of variance (ANOVA) were performed by GraphPad Prism 7 (San Diego, CA, USA). A value of *p* < 0.05 was regarded as a significant difference, and a value of *p* < 0.01 was regarded as a very significant difference. Principal components analysis (PCA) and orthogonal partial least squares discriminant analysis (OPLS-DA) was performed by SIMCA-P 14.0 software (Umetrics, Umea, Sweden). Correlation analysis and heat-map were performed by Origin 2017 software (OriginLab, Massachusetts, USA).

4. Conclusions

HSYA had the highest content in safflower, and it was the main active ingredient [44,45]. Studies have also pointed out that HSYA content was related to the color of safflower. This research indicates that the content of ASYB and SC can also affect the color and its anti-thrombotic activity. The higher content of HSYA, ASYB, and SC makes safflower redder, yellower, brighter, more orange–yellow, and more vivid, due to the strong correlation between the color and compounds content. Furthermore, the high content of these compounds leads to better anti-thrombotic activity. Therefore, safflowers with different colors might have different contents of compounds and different pharmacological activities. Moreover, it is also feasible to evaluate the quality of other flower medicinal materials based on this strategy.

Supplementary Materials: The following are available online, Table S1: Calibration curves, correlation coefficients, linearity ranges, LODs, and LOQs of the 16 compounds, Table S2: Precision, repeatability, stability, and recovery of 16 compounds, Figure S1: Comparison of colour indicators between class I and II of safflower, Figure S2: Representative chromatogram of 16 compounds in safflower, Figure S3: Quantitative determination of 16 compounds from class I and II of safflower, Figure S4: CIE LAB colour space.

Author Contributions: Y.-P.T., S.-J.Y., S.-L.S. and J.-A.D. designed the experiments; Z.-J.P., X.-Q.S., Z.-H.Z., Y.-Y.C., S.-L.H., G.-P.P., performed the experiments; Z.-J.P., G.-S.Z., H.Y., J.-Q.B., X.-P.W., analyzed the data; Z.-J.P., S.-J.Y. and Y.-P.T., G.-S.Z. wrote the paper. All authors participated in the preparation of the manuscript and approved the final manuscript.

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Sample Availability: Samples of the compounds hydroxysafflor yellow A and anhydrosafflor yellow B are available from the authors.



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