

Doi:10.32604/phyton.2025.061099

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Comparative Karyotyping Reveals the Origin of Chinese Kale (*Brassica oleracea* var. *alboglabra*)

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Received: 17 November 2024; Accepted: 27 January 2025; Published: 06 March 2025

ABSTRACT: This study conducted karyotype analysis on 56 Chinese kale varieties from Guangdong and Fujian provinces using conventional chromosome analysis methods. The varieties were categorized into four groups based on their origin and flower color: white-flower Chinese kale originating from Guangdong (GW), yellow-flower Chinese kale originating from Guangdong (GY), white-flower Chinese kale originating from Fujian (FW), and yellow-flower Chinese kale originating from Fujian (FY). Karyotype differences among the four groups of Chinese kale were analyzed, and the evolutionary relationship between yellow-flower and white-flower Chinese kale from the two regions was inferred based on karyotype parameters. The results indicated that all Chinese kale varieties were diploid with 2n = 2x = 18, including a pair of satellites. The chromosome types included median-centromere (m) and sub-median-centromere (sm), and the karyotypes were 1A and 2A, a sper karyotype asymmetry index. The karyotype 1A of Chinese kale was identified for the first time. Partial least squares discriminant analysis (PLS-DA) identified nine karyotypic indicators that differentiated the four groups, and these differences were further visualized using heatmaps and box plots. Based on the evolution trends in the four groups and PLS-DA analysis, it was speculated that white-flower Chinese kale originated from Guangdong, yellow-flower Chinese kale originated from Fujian, and GY and FW were derived from the cross of GW and FY. This study provides a reference for understanding the genetic relationships between Chinese kale in Guangdong and Fujian, and offers a cytological basis for the evolution, hybridization, and phylogenetic relationships of Chinese kale.

KEYWORDS: Chinese kale; chromosome; karyotype; evolution

1 Introduction

Chinese kale (*Brassica oleracea* var. *alboglabra*), historically known as Gai Lan or Lan Cai, is a cruciferous brassica vegetable primarily consumed for its tender leaves and bolting stems. It is rich in glucosinolates, carotenoids, and ascorbic acid, and is considered to be beneficial to health [1–3].

Yellow-flower and white-flower Chinese kale, as two distinct types, exhibit variations in biological characteristics and nutritional properties [4]. For instance, white-flower Chinese kale had higher chlorophyll content but lower vitamin C content in its bolting stems compared to yellow-flower Chinese kale [5]. Additionally, white-flower Chinese kale possessed fewer branches and a sweeter, more tender quality, rendering it more popular among consumers [6]. In China, white-flower Chinese kale is widely cultivated



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and distributed, whereas yellow-flower Chinese kale is primarily grown in limited quantities, with most production concentrated in Fujian Province. Over the years, the cultivation of Chinese kale has expanded from southern to northern China, establishing it as an important *Brassica oleracea* vegetable for export [7,8].

Vegetables of *Brassica oleracea*, such as cauliflower (*Brassica oleracea* var. *botrytis*) and ornamental kale (*Brassica oleracea* var. *acephala*), are generally believed to have originated from the Mediterranean coast. However, the origin of Chinese kale remains a subject of debate, with two prevailing theories: one posits a Mediterranean coastal origin, while the other suggests a southern China origin. Zhang [9] conducted a comprehensive investigation of diverse graphic and textual materials, concluding that Chinese kale was introduced into China as early as the 5th and 6th centuries AD during the Southern and Northern Dynasties, it underwent mutation and long-term artificial selection in Guangdong, thereby supporting the theory of its southern China origin. Wang et al. [10] studied genetic diversity in 38 *Brassica olearacea* vegetable samples, including eight varieties of Chinese kale, and found that Chinese kale evolved from other *Brassica olearacea* vegetables via distinct routes, further supporting the southern China origin theory has gained consensus [11,12], research on the intraspecific origin of Chinese kale remains insufficient. To address this gap, this study collected 56 varieties of Chinese kale from Guangdong and Fujian and conducted karyotype analysis to provide a cytological basis for understanding the origin of Chinese kale.

Karyotype refers to the chromosome set's phenotype during mid-mitosis, summarizing chromosome number, size, and morphological features. Karyotype analysis is critical for cytogenetic classification and species kinship studies, particularly in plant research [13–16]. As stable carriers of genetic information, chromosomes are consistent in morphology, structure, and number, and they can be stained effectively by basic dyes. Karyotype analysis can be performed using non-banding or banding techniques [17]; this study utilized a non-banding technique. Mousavizadeh et al. [18] employed non-banding technology to analyze the karyotype and chromosome diversity of Iranian wild asparagus, providing insights into Asparagus evolution, while Wang et al. [19] investigated karyotypic differences among Chinese cherry (*Cerasus pseudocerasus* Lindl.) and four related Cerasus species to inform cross-breeding, phylogenetic relationships, and evolutionary dynamics.

In this study, 56 varieties of Chinese kale from Guangdong and Fujian were categorized into four groups: white-flower Chinese kale originating from Guangdong (GW), yellow-flower Chinese kale originating from Guangdong (GY), white-flower Chinese kale originating from Fujian (FW), and yellow-flower Chinese kale originating from Fujian (FY). Karyotype analysis was performed to elucidate karyotype variations and analyze evolutionary trends, aiming to provide cytological evidence for interspecific affinity and evolutionary origins of Chinese kale.

2 Materials and Methods

2.1 Plant Materials

This experiment utilized 56 varieties of Chinese kale from Guangdong and Fujian (which were preserved and propagated by our research team), abbreviated as GW (30 samples), GY (5 samples), FW (10 samples), and FY (11 samples) according to their flower color and origin. Each sample was as-signed a specific number, as detailed in Table 1.

No.	Variety	Origin	Color	No.	Variety	Origin	Color
GW1	Liuyetianjielan	Guangdong	White	GW29	Sijijielan	Guangdong	White
GW2	Baihuajielan	Guangdong	White	GW30	Dengfengzhongchi	Guangdong	White
GW3	Huaxianzi	Guangdong	White	GY1	Hongjiaojielan	Guangdong	Yellow
GW4	Guisunjielan	Guangdong	White	GY2	Zhengyuandasun	Guangdong	Yellow
GW5	Xianggujielan	Guangdong	White	GY3	Xianggangtiancui	Guangdong	Yellow
GW6	Taiwansiji	Guangdong	White	GY4	Huanghuajielankuai	Guangdong	Yellow
GW7	Mingfengxianggu	Guangdong	White	GY5	Cutiaojielankuai	Guangdong	Yellow
GW8	Kuaidasiji	Guangdong	White	FW1	JL-08	Fujian	White
GW9	Chenghaisiji	Guangdong	White	FW2	JL-14	Fujian	White
GW10	Cutiaojielan	Guangdong	White	FW3	JL-03-B	Fujian	White
GW11	Tezaosijidasun	Guangdong	White	FW4	JL-09	Fujian	White
GW12	Jielanwang	Guangdong	White	FW5	JL-05-B	Fujian	White
GW13	Minijielan	Guangdong	White	FW6	Bolicui	Fujian	White
GW14	Zhonghuajielan	Guangdong	White	FW7	JL-10-1B	Fujian	White
GW15	Tiancuijianye	Guangdong	White	FW8	JL-12-B	Fujian	White
GW16	Jienongjielan	Guangdong	White	FW9	Zhangzhou-B	Fujian	White
GW17	Hongjielantai	Guangdong	White	FW10	JL-07-B	Fujian	White
GW18	Jingxuancutiaoyusun	Guangdong	White	FY1	Huanghuajielan	Fujian	Yellow
GW19	Tiancuijielankuai	Guangdong	White	FY2	JL-03-H	Fujian	Yellow
GW20	Sijitianjielan	Guangdong	White	FY3	JL-10-1H	Fujian	Yellow
GW21	Nairecutiao	Guangdong	White	FY4	JL-12-1	Fujian	Yellow
GW22	Zhonghuasuinong	Guangdong	White	FY5	JL-12-H	Fujian	Yellow
GW23	Zhonghuajinma	Guangdong	White	FY6	Zhangzhou-H	Fujian	Yellow
GW24	Sijicutiao	Guangdong	White	FY7	Fuzhouhuanghua	Fujian	Yellow
GW25	Chenghaicutiaodarou	Guangdong	White	FY8	JL-1-2	Fujian	Yellow
GW26	Sijidazhong	Guangdong	White	FY9	JL-01-1	Fujian	Yellow
GW27	Darou-1	Guangdong	White	FY10	JL-07-H	Fujian	Yellow
GW28	Daroushuangcui	Guangdong	White	FY11	JL-10	Fujian	Yellow

Table 1: Experimental varieties of Chinese kale collected from different parts of China

2.2 Root Tip Culture

Selected full-grained seeds were soaked for 2 h at room temperature, placed in petri dishes lined with moist filter paper, and cultured in the dark at 25°C for 3 days. When the root tips reached a length of 10–15 mm, they were harvested at 9 am [20].

2.3 Chromosome Preparation

Root tips were harvested and pretreated in 2 mmol·L⁻¹ 8-hydroxyquinoline solution for 5 h at 4°C. The pretreated root tip materials were fixed in Carnoy solution (volume ratio of absolute ethanol to acetic acid was 3:1) at 4°C for 24 h, washed with 95%, 85%, and 75% ethanol gradient, and finally kept in 75% ethanol solution. The root tip was dissociated with 1 mol·L⁻¹ HCl at 60°C for 8 min, then washed with distilled water for 3 times, stained with Carbol-fuchsin solution for 5–10 min in dark, and squashed for viewing under 100 times oil lens and 10 times evepiece of OlympusCX21 microscope.

2.4 Karyotype Analysis

Thirty well-spread, morphologically clear metaphase cells were selected for chromosome counting and karyotype analysis, following the criteria of Li et al. [21]. Chromosome measurements and classifications were done based on the standards of Levan et al. [22]. The karyotype asymmetry coefficient was calculated

using the method of Arano [23], and karyotype classification was performed according to the regulations of Stebbins [24].

2.5 Data Analysis

Experimental data were processed using Excel 2019. Partial least squares discriminant analysis (PLS-DA) was conducted using SIMCA-P 14.1. Heatmaps were generated using MeV 4.9.0, and box plots were created using Origin 2021. Correlation analysis was performed with Excel 2019, and the results were visualized using Cytoscape 3.9.1 [25].

3 Results

In this study, metaphase chromosomes of 56 varieties of Chinese kale, categorized as GW, GY, FW, and FY, were observed (Fig. 1). Karyograms (Fig. 2) and karyotype diagrams (Fig. 3) were generated, and statistical analyses of these figures yielded the karyotype parameters (Table S1) and karyotypes (Table 2). The detailed analysis is as follows.

3.1 Chromosome Number and Characteristics

All 56 varieties of Chinese kale were diploid with 2n = 2x = 18, and no cells with abnormal chromosome numbers were detected (Fig. 2). Each variety possessed a pair of satellite chromosomes, with variations in length and location among different varieties. The satellites were predominantly located on chromosomes 7, 8, and 9, present in approximately 78.57% of all Chinese kale varieties (Fig. 3). Chromosome 8 had the highest frequency of satellites, observed in 17 varieties. Additionally, the satellite-bearing chromosomes were primarily of the sm type, representing 64.29% of all varieties, including 56.67% of GW, 100% of GY, 50% of FW, and 81.82% of FY.

3.2 Karyotype Analysis

As shown in Table 2, most Chinese kale varieties consisted of both median-centromere (m) and submedian-centromere (sm) chromosomes, while a few varieties contained only median-centromere (m) chromosomes. The specific karyotype formulas varied among different varieties of Chinese kale.

When analyzed together with Tables 2 and S1, the relative lengths of chromosomes ranged from 7.39% to 14.58% in GW, 8.54% to 14.33% in GY, 8.35% to 14.09 in FW, and 8.69% to 14.29% in FY. The centromere indices for GW, GY, FW, and FY ranged from 25.67–49.64, 27.61–48.7, 27.13–48.93, and 31.86–49.35, respectively. The ranges of relative chromosome lengths and centromere index of GW represented all the groups' ranges in this experiment. However, the range of ratios of the longest and shortest chromosomes for GW was observed at 1.31–1.87, encompassing those of GY (1.31–1.62) and FW (1.42–1.59), but not fully encompassing that of FY (1.21–1.61).



Figure 1: Metaphase chromosomes of different varieties of Chinese kale

GW1	A	2 N	88		**		24	44	A1	GY1 👗	8	A X	# 8	8 R			* 1	14	14
GW2	43		**		t #			4.6	••	GY2 🛔	8	1 R	X 2	25	# 8	ŝ.Ă.	• •	9 ð	• A
GW3	XX	3#		**	**	A R			66	GY3 🏮	X	XK	**		**	**	¢i,		# 6
GW4	11	11	13	6.2	58	\$1	¥ #	A 8		GY4 👗	K	X X	\$ 8	8 X	•	64		11	X n
GW5	14	14	**			**	à à			GY5 🍞	6	**	} X	ğe.	81	**	ńх	68	(*
GW6	* #	**		4.8	\$.2	4 X	**	18	48										
GW7	XX	A Y	**	ñ.#	::	λï		8.8											
GW8	58	11	AR	4.6			\$ ŝ	FA	44	FW1 🐇	K.	11	11	11				34	
GW9						• #		4.	۹.	FW2 🏅	ł	6.8	8.8	8.8	Å Å	2.2	25		
GW10		11	13			1.8	65		6 B	FW3 🛔	ă	8.8	4 X	8.6		-			
GW11	38							34		FW4 🝍	8	44	# 1	XX		# X.			
GW12	ХX	3.5	< X	£ \$	ā K	8.3) X	<u>j.</u> k	A 1	FW5 🍍	t			8 1					• 8
GW13				4 i	••	••	**	••	••	FW6 🚪	8	88	4.8			X 3	18		
GW14	XX		8.8	XX	43	8 ¥	2.2	4	8.A	FW7 🤰	X	8 R	X X	XX	6.6	6.1	8.8	x #	ź.ż
GW15	XX	88		<i>#</i> 1	4.8	4.8	25	14	8 A	FW8 👗	Ă	8 K	8.8	* *	6 X	á n'	4.6	**	
GW16	a 🕷	**	E K	# 6	* *		8 X	10	4 H	FW9 🍍	8	8 8	3.2	11	15	* *) x		z 1
GW17	81		13	2 2	32			11	• ¥	FW10 🚦	8	38	11	15	13	11	8.8		
GW18	1	81	-		XR	44			85										
GW19	33	13	11		*	**	23	\$3	4 #										
GW20	*					**	3 8	**	44	FY1 🌹	8			•	••			Å }	
GW21	48		8.8	X #		8 A	* 6			FY2 🍍	8			4 8			8.8	• •	6.6
GW22		8.8	8 X				48			FY3 🥇	7	8 X	XX	ññ	* *	8.8	XX	3 X	ńż.
GW23	11					A 8			88	FY4 🍍	Ă	8.8	* *	X X	8 R	8 R	Å Í	* *	4 A
GW24	* *	88	**	**		**	**		8.6	FY5 👗	×	8.8		X 3	**			* 4	6 %
GW25	5 3 3	••	81	8 8				66		FY6 🎁		X X	8 M	3.4	iı İ		=	61 AN	10 14
GW26	5 🛢 🛢				41			- 68		FY7 💄	•	4.8						Á 6	**
GW27	38	88					42			FY8 💧	£	á X	**	A A	¥ K	R 7	8.6	áà	展为
GW28	8 📕 📕	8 R			8 8				-	FY9 🝍	9		8.8	8.8	8 8	* *		4.8	8 Č
GW29	38	88	18	6 A	22		X 8	61	4.4	FY10 🌋	8		8.8						
GW30			88					ă Â	••	FY11 🄱		11			11		ž ž		8 8
																	10	μm	

Figure 2: Karyograms of different varieties of Chinese kale



Figure 3: (Continued)



Figure 3: Chromosome ideograms of different varieties of Chinese kale. In the *Y* axis, the upper part of 0 represents the short arm, and the lower part represents the long arm

No.	Karyotype formula	SAT		Lc/Sc	MAR	Karyotype	As. K (%)
		Number	Chromosome number				
GW1	2n = 2x = 18 = 12m + 6sm (2SAT)	2	9	1.57	1.59	2A	60.09
GW2	2n = 2x = 18 = 12m + 6sm (2SAT)	2	8	1.57	1.60	2A	60.68
GW3	2n = 2x = 18 = 14m + 4sm (2SAT)	2	9	1.47	1.41	2A	57.69
GW4	2n = 2x = 18 = 10m + 8sm (2SAT)	2	5	1.62	1.51	2A	59.88
GW5	2n = 2x = 18 = 18m (2SAT)	2	7	1.42	1.35	1A	57.25
GW6	2n = 2x = 18 = 10m + 8sm (2SAT)	2	9	1.48	1.59	2A	60.18
GW7	2n = 2x = 18 = 12m + 6sm (2SAT)	2	6	1.87	1.63	2A	60.91
GW8	2n = 2x = 18 = 16m (2SAT) + 2sm	2	7	1.68	1.48	1A	59.42
GW9	2n = 2x = 18 = 18m (2SAT)	2	7	1.43	1.24	1A	55.16
GW10	2n = 2x = 18 = 16m + 2sm (2SAT)	2	7	1.63	1.39	1A	57.95
GW11	2n = 2x = 18 = 18m (2SAT)	2	8	1.56	1.29	1A	56.24
GW12	2n = 2x = 18 = 10m + 8sm (2SAT)	2	8	1.59	1.55	2A	60.46
GW13	2n = 2x = 18 = 12m + 6sm (2SAT)	2	4	1.46	1.47	1A	59.23
GW14	2n = 2x = 18 = 12m (2SAT) + 6sm	2	8	1.44	1.66	2A	61.29
GW15	2n = 2x = 18 = 14m (2SAT) + 4sm	2	9	1.56	1.40	2A	58.01
GW16	2n = 2x = 18 = 18m (2SAT)	2	8	1.48	1.35	1A	57.29
GW17	2n = 2x = 18 = 12m + 6sm (2SAT)	2	8	1.64	1.50	2A	59.41
GW18	2n = 2x = 18 = 14m (2SAT) + 4sm	2	6	1.44	1.49	2A	59.46
GW19	2n = 2x = 18 = 10m + 8sm (2SAT)	2	5	1.60	1.62	2A	60.81
GW20	2n = 2x = 18 = 16m (2SAT) + 2sm	2	9	1.43	1.55	2A	59.48
GW21	2n = 2x = 18 = 16m (2SAT) + 2sm	2	7	1.44	1.49	2A	58.81
GW22	2n = 2x = 18 = 10m + 8sm (2SAT)	2	7	1.44	1.58	2A	60.41
GW23	2n = 2x = 18 = 16m + 2sm (2SAT)	2	6	1.64	1.52	2A	59.87
GW24	2n = 2x = 18 = 14m + 4sm (2SAT)	2	7	1.51	1.49	1A	59.43
GW25	2n = 2x = 18 = 16m (2SAT) + 2sm	2	8	1.40	1.41	1A	58.37
GW26	2n = 2x = 18 = 14m + 4sm (2SAT)	2	8	1.53	1.39	1A	57.63
GW27	2n = 2x = 18 = 16m (2SAT) + 2sm	2	7	1.45	1.30	1A	56.45
GW28	2n = 2x = 18 = 12m + 6sm (2SAT)	2	7	1.33	1.60	2A	60.83
GW29	2n = 2x = 18 = 16m + 2sm (2SAT)	2	8	1.31	1.33	2A	56.39
GW30	2n = 2x = 18 = 16m (2SAT) + 2sm	2	8	1.64	1.46	1A	58.65
GY1	2n = 2x = 18 = 14m + 4sm (2SAT)	2	8	1.60	1.48	1A	59.30
GY2	2n = 2x = 18 = 8m + 10sm (2SAT)	2	6	1.62	1.68	2A	61.39
GY3	2n = 2x = 18 = 14m + 4sm (2SAT)	2	7	1.62	1.55	2A	60.09
GY4	2n = 2x = 18 = 14m + 4sm (2SAT)	2	8	1.31	1.56	2A	59.51
GY5	2n = 2x = 18 = 14m + 4sm (2SAT)	2	7	1.33	1.49	2A	59.17
FW1	2n = 2x = 18 = 10m + 8sm (2SAT)	2	8	1.42	1.58	2A	59.90
FW2	2n = 2x = 18 = 10m + 8sm (2SAT)	2	5	1.49	1.62	2A	61.14
FW3	2n = 2x = 18 = 16m (2SAT) + 2sm	2	9	1.59	1.40	1A	58.45
FW4	2n = 2x = 18 = 18m (2SAT)	2	9	1.46	1.38	1A	57.67
FW5	2n = 2x = 18 = 14m + 4sm (2SAT)	2	7	1.48	1.43	2A	58.19
FW6	2n = 2x = 18 = 18m (2SAT)	2	8	1.55	1.31	1A	56.42
FW7	2n = 2x = 18 = 10m + 8sm (2SAT)	2	9	1.46	1.68	2A	61.93
FW8	2n = 2x = 18 = 12m + 6sm (2SAT)	2	6	1.57	1.64	2A	61.33
FW9	2n = 2x = 18 = 18m (2SAT)	2	7	1.42	1.38	1A	57.69
FW10	2n = 2x = 18 = 18m (2SAT)	2	6	1.54	1.36	1A	57.64
FY1	2n = 2x = 18 = 14m + 4sm (2SAT)	2	8	1.21	1.37	1A	57.44
FY2	2n = 2x = 18 = 12m + 6sm (2SAT)	2	9	1.47	1.52	1A	60.23
FY3	2n = 2x = 18 = 10m + 8sm (2SAT)	2	9	1.42	1.55	2A	59.83
FY4	2n = 2x = 18 = 14m + 4sm (2SAT)	2	7	1.59	1.53	2A	59.84
FY5	2n = 2x = 18 = 14m + 4sm (2SAT)	2	9	1.40	1.38	1A	57.11
FY6	2n = 2x = 18 = 12m + 6sm (2SAT)	2	5	1.46	1.52	1A	59.72

 Table 2: The karyotypes of different varieties of Chinese kale

(Continued)

Table 2 (continued)											
No.	Karyotype formula		SAT	Lc/Sc	MAR	Karyotype	As. K (%)				
		Number	Chromosome number								
FY7	2n = 2x = 18 = 16m + 2sm (2SAT)	2	8	1.39	1.44	1A	58.71				
FY8	2n = 2x = 18 = 16m (2SAT) + 2sm	2	8	1.61	1.41	1A	58.28				
FY9	2n = 2x = 18 = 12m + 6sm (2SAT)	2	9	1.39	1.61	2A	61.17				
FY10	2n = 2x = 18 = 18m (2SAT)	2	6	1.40	1.32	1A	56.66				
FY11	2n = 2x = 18 = 12m + 6sm (2SAT)	2	7	1.37	1.48	2A	59.14				

Note: Lc/Sc, ratio of the longest and shortest chromosome; MAR, mean of long-arm length and short-arm length ratio; As. K (%), index of the karyotypic asymmetry.

Among 56 samples, the highest mean arm ratio was 1.68, and the lowest was 1.24. In GW, the arm ratio varied from 1.01 to 2.90, with 8.89% of chromosomes having an arm ratio greater than 2 and a mean arm ratio of 1.47 for the entire group. In GY, the arm ratio ranged from 1.05 to 2.62, with 17.78% of chromosomes having an arm ratio greater than 2 and a mean arm ratio of 1.55 for the group. In FW, the arm ratio ranged from 1.04 to 2.69, with 10.00% of chromosomes having an arm ratio greater than 2 and a mean arm ratio of 1.45. In FY, the arm ratio varied from 1.03 to 2.14, with 45.45% of chromosomes having an arm ratio greater than 2 and a mean arm ratio of 1.47. Interestingly, although GW had the largest range in arm ratios among the four groups, it had the smallest proportion of chromosomes with an arm ratio greater than 2. In contrast, FY, which had the smallest range in arm ratios, had the largest proportion greater than 2. The karyotype asymmetry index was similar across the four groups, ranging from 55.16% to 61.29% for GW, 59.17% to 61.39% for GY, 56.42% to 61.93% for FW, and 56.66% to 61.17% for FY. Chinese kale exhibited two karyotype types: 1A, accounting for 44.64%, and 2A, accounting for 55.36% (Table 2).

3.3 Principal Component Analysis

To analyze the relationships among the four groups of Chinese kale based on karyotypic parameters, partial least squares discriminant analysis (PLS-DA) was conducted. As shown in Fig. 4, PLS-DA1 explained 12.8% of all parameter information, effectively differentiating between Chinese kale from Guangdong (GW and GY) and Fujian (FW and FY). PLS-DA2 explained 12.5% of all parameter information, distinguishing GY from FW. Overall, there was overlap among the groups, making complete clustering challenging. However, some regional differences were observed, providing a basis for further differential analysis.



Figure 4: PLS-DA analysis of karyotypes parameters of different varieties of Chinese kale

3.4 Karyotype Differences among Four Groups of Chinese Kale

To comprehensively study the karyotype differences among the four groups of Chinese kale, nine differential indicators were selected from VIP values greater than 1, based on PLS-DA. Heatmaps and box plots were generated to visually observe group differences (Fig. 5). As shown in Fig. 5A, the karyotype indicators were generally clustered into two major groups. One group included the relative length of chromosome 9, the relative length of the short arm of chromosome 7, and the minimum centromere index, while the other six indicators clustered into the second group.

From the nine box-plots in Fig. 5, although the ranges of GW indicators were broad, their median values were centered among all groups. The GY group showed higher median values than the other groups for the relative length of the short arm of chromosome 3, the relative length of chromosome 3, and the maximum arm ratio (Fig. 5B,C,G), but had significantly lower values for the relative length of the short arm of chromosome 7 and the minimum centromere index (Fig. 5I,J). The FW group had the lowest median value for the relative length of chromosome 2 among all groups, while the FY group had a significantly higher median value for the relative length of chromosome 9 compared to the other groups (Fig. 5D,H).



Figure 5: Heatmaps and box-plots of karyotypes parameters of different varieties of Chinese kale. (A) Heatmaps of 9 indexes of different Chinese kale varieties. (B) Relative length of the short arm of chromosome 3. (C) Relative length of chromosome 2. (E) Coefficient of variation in the average length of chromosome set. (F) Ratio of the longest and shortest chromosome. (G) Maximum arm ratio. (H) Relative length of chromosome 9. (I) Relative length of the short arm of chromosome 7. (J) Minimum centromere index

3.5 Correlation Analysis

To investigate the correlations among kale karyotype parameters, 41 relationships were identified (Fig. 6). The relative length of the long arm and the relative length of the short arm of chromosome 2 to 8 were negatively correlated. The chromosome type was positively correlated with the number of that chromosome type within the set. There was a strong negative correlation between the minimum centromere index and the maximum arm ratio, as well as between the maximum centromere index and the minimum arm ratio. Although karyotype classification is typically associated with chromosome length ratio, correlations were found between chromosome karyotype and maximum arm ratio, minimum centromere index, karyotype asymmetry, and the proportion of arm ratios greater than 2, but not with chromosome length ratio (Fig. 6).



Figure 6: Correlation between karyotypes parameters of different varieties of Chinese kale. The solid lines represent a positive correlation. The dashed lines represent a negative correlation. All correlations in the figure reflect Pearson correlation coefficient values above the threshold ($|\rho| > 0.65$)

3.6 Evolutionary Trend Analysis

To assess the evolutionary degree of Chinese kale, karyotypic evolutionary trends were analyzed using the average arm ratio and karyotypic asymmetry index as references. As observed in Fig. 7, the overall evolutionary level of GY was the highest, while the remaining three groups of Chinese kale varieties were evenly distributed across the plot. GW, the group with the most varieties, included the oldest Chinese kale, with GW9 notably distinct from the other varieties. Overall, GY was the most evolved among the four groups of Chinese kale.



Figure 7: Karyotypic evolutionary trend map of different varieties of Chinese kale

4 Discussion

4.1 Differences in Karyotypes between This Study and Previous Studies

In this study, the number of chromosomes for all Chinese kale was 2n = 2x = 18, each with two satellites, consistent with previous research [26–31]. However, the distribution of these satellites differed. Earlier studies typically localized most satellites on chromosome 7 [27,29,31]. In the study by Song et al. [26], satellites were found to be located on chromosomes 6 and 7, while Yuan et al. [30] reported their presence on chromosome 9. In contrast, our study, which examined multiple varieties, found satellites on chromosomes 4 to 9. Xia et al. [28] reported that the chromosome types of satellites were all sm, whereas this study revealed the presence of satellites on the m chromosome type in 35.71% of all varieties. Previous studies classified all karyotypes as "2A". However, we observed that many Chinese kale varieties had a "1A" karyotype, providing a reference for future studies on the karyotype of Chinese kale.

4.2 Karyotype Differences among Four Groups of Chinese Kale

The karyotype formulas of the four groups was primarily composed of multiple m chromosomes and a few sm chromosomes, with some groups having only m chromosomes. In GY, the predominant karyotype formula was 2n = 2x = 18 = 14m + 4sm (2SAT), with no exclusively m chromosomes. In both GY and FY, most chromosomes with satellites were of the sm type. This phenomenon was less pronounced in GW and FW, where only about half of the chromosomes with satellites were sm type. Further investigation is needed to determine if there is a relationship between satellite chromosome type and flower color in Chinese kale.

The average arm ratio of each group of Chinese kale was approximately 1.5, indicating genetic stability. The proportion of chromosomes greater than 2 differed among groups, with FY exhibiting a significantly higher proportion at 45.45% compared to the other three groups. Additionally, the relative length of chromosome 9 in FY was generally the highest among four groups, possibly contributing to its low chromosome length ratio (Fig. 5F,H). GY exhibited higher relative length of the short arm of chromosome 3 and maximum arm ratio compared to other groups (Fig. 5B,G), but significantly lower relative length of the short arm of chromosome 7 and the minimum centromere index (Fig. 5I,J). FW had the lowest relative length

of chromosome 2 among the four groups (Fig. 5D). These variations constitute key karyotypic differences among the different Chinese kale groups.

According to studies by Li et al. [32] and Zhang et al. [33], yellow flower color dominance over white flower color in Chinese kale followed single-gene inheritance as a qualitative trait. Li et al. [32] identified genes related to flower color inheritance on the long arm of chromosome 3. Although differences in short arm length and total length for chromosome 3 were observed among groups (Fig. 5), the long arm length of this chromosome did not differ significantly. Therefore, it can be concluded that flower color gene in Chinese kale was genetically stable and rarely mutate.

4.3 Evolution Analysis of Four Groups of Chinese Kale

The asymmetry index is widely recognized as a key indicator for assessing the evolutionary level of plants. In this study, the asymmetry index of the 56 varieties of Chinese kale was low, ranging from 55.16% to 61.93%, indicating that Chinese kale is an ancient plant.

In Fig. 6, there was no correlation between the karyotypes of Chinese kale and the chromosome length ratio, as the chromosome length ratios of all kale varieties were less than 2, consistent with previous studies [26–31]. It can be concluded that the karyotype of Chinese kale is determined by the proportion of arm ratios greater than 2. The stability of the chromosome length ratio in Chinese kale indicated the stability of its chromosomal inheritance.

As observed in Fig. 7, the scatter points representing the four groups of Chinese kale were evenly distributed on the karyotypic evolutionary trend map, with GY appearing to be the most evolved, likely due to subsequent hybridization. GW, however, included several primitive varieties, with GW9 identified as particularly primitive and distinct from the group. Geographically, genetic exchanges between Guangdong and Fujian have likely contributed to frequent genetic exchanges among Chinese kale varieties grown in these regions [34]. In practice, most varieties grown in Guangdong are white-flower Chinese kale, while those in Fujian are yellow-flower Chinese kale [35]. Combined with Fig. 4, GW and FY were located on the left and right sides of the distribution, respectively, while GY and FW were positioned between them. Therefore, it was hypothesized that white-flower Chinese kale originated in Guangdong, yellow-flower Chinese kale originated in Fujian, and GY and FW were hybrids of GW and FY. However, relevant molecular evidence is still lacking, and further research on the evolution of Chinese kale is needed.

5 Conclusions

In this study, 56 varieties of Chinese kale were karyotyped and categorized into four groups based on their origin and flower color. Karyotype differences among the four groups were analyzed, and the evolutionary history of Chinese kale was speculated. All varieties were diploid with 2n = 2x = 18. In addition to karyotype 2A, type 1A was also identified. This research provides a reference for studying the genetic relationship between Guangdong and Fujian Chinese kale and offers a cytological basis for understanding the evolution, hybridization, and phylogenetic relationships of Chinese kale.

Acknowledgement: None.

Funding Statement: This research was funded by National Natural Science Foundation of China (32372732, 32372683, 32460750, 32072586, 31500247), Sichuan Science and Technology Program (2023ZYD0090, 24NSFSC0404, 2025ZNS-FSC1112), Sichuan Innovation Team of National Modern Agricultural Industry Technology System (SCCXTD-2024-05), and, Key Laboratory of Storage of Agricultural Products, Ministry of Agriculture and Rural Aairs (kt202410), and Chengdu Science and Technology Program (2024-YF05-02375-SN).

Author Contributions: The authors confirm contribution to the paper as follows: study conception and design: Bo Sun, Fen Zhang; data collection: Xuena Yu, Kehao Liang, Victor Hugo Escalona; analysis and interpretation of results: Yi Tang, Zhi Huang, Huanxiu Li, Zhifeng Chen; draft manuscript preparation: Sha Luo, Shuang Wu, Junyan Song. All authors reviewed the results and approved the final version of the manuscript.

Availability of Data and Materials: The authors confirm that the data supporting the findings of this study are available within the article.

Ethics Approval: Not applicable.

Conflicts of Interest: The authors declare no conflicts of interest to report regarding the present study.

Supplementary Materials: The supplementary material is available online at https://doi.org/10.32604/phyton.2025. 061099.

Abbreviations

- S The relative length of short arm
- L The relative length of long arm
- L+S The relative length of chromosome
- m Median-centromere chromosome
- sm Submedian-centromere chromosome
- SAT Satellite
- Lc/Sc Ratio of the longest and shortest chromosome
- MAR Mean of long-arm length and short-arm length ratio
- As. K Index of the karyotypic asymmetry

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