

Cocktail of Astragalus Membranaceus and Radix Trichosanthis Suppresses Melanoma Tumor Growth and Cell Migration Through Regulation of Akt-Related Signaling Pathway

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Edited by:

Jiang-Jiang Qin, Institute of Cancer and Basic Medicine (CAS), China

Reviewed by:

Jun Li, Huazhong University of Science and Technology, China Man-Tat Lau, Children's Medical Research Institute, Australia Sung-Gook Cho, Korea National University of Transportation, South Korea

*Correspondence:

Shengli Wei wsl7491@126.com Xiuhua Hu xiuhuahu@126.com

[†]These authors have contributed equally to this work and share first authorship

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¹School of Life Sciences, Beijing University of Chinese Medicine, Beijing, China, ²Dongfang Hospital, Beijing University of Chinese Medicine, Beijing, China, ³School of Traditional Chinese Medicine, Beijing University of Chinese Medicine, Beijing, China, ⁴School of Chinese Materia Medica, Beijing University of Chinese Medicine, Beijing, China

Background: Malignant melanoma has high morbidity and mortality and limited treatment options. Traditional Chinese medicine has great potential in the clinical therapy of cancer, and the theory of compatibility is one core content of Chinese medical theory. *Astragalus Membranaceus* and *Radix Trichosanthis* are clinically effective for the treatment of various cancers.

Methods: We verified the effects of AMD, RTD, and their "cocktail" on melanoma model *in vitro* and *in vivo* and the mechanism of its effect on the Akt-related signaling pathway by network pharmacology, MTT, flow cytometry, LDH, SOD, MDA assay, and Western blot.

Results: The network pharmacology analysis indicated that the PI3K-Akt pathway plays a crucial role in the treatment of malignant melanoma with these two herbs. In addition, AMD, RTD, and their "cocktail" could inhibit the proliferation of A375 cells by reducing the survival rate in a concentration-dependent manner and by regulating the cell cycle, and the compatibility of two herbs also could inhibit melanoma growth. They could, respectively, induce apoptosis and inhibit migration by affecting the expression of Bcl-2, Bax, p53, snail, E-cadherin, and N-cadherin. Furthermore, LDH activity was decreased, while SOD increased and MDA reduced. The factors of the Akt-related signaling pathway, Akt and p-Akt, were decreased.

Conclusion: This study showed that AMD, RTD, and their "cocktail" could regulate cell proliferation, apoptosis, and metastasis in A375 cells through the suppression of the Akt-related signaling pathway, and the "cocktail" groups had detoxification and additive effects. The best compatibility of the two herbs also can inhibit tumor growth and metastasis *in vivo*.

Keywords: Astragalus Membranaceus, Radix Trichosanthis, herbal "cocktail", Akt-related singnaling pathway, proliferation, migration

INTRODUCTION

Malignant melanoma (MM) is a malignant cancer characterized by increasing incidence and high metastatic capacity and which is originated from the dysplasia of melanocytes. In America, MM ranks third as the most common cancer in men and fifth in women (Miller et al., 20192019). Although the overall cancer incidence rate is reportedly in decline, the incidence of MM continues to increase, with an estimated annual increase of approximately 3%. In addition, the patients with MM are mostly younger, with 47% of the patients ≤65 years old and $33\% \leq 50$ years (Miller et al., 20192019). Surgery is the main treatment for early-stage MM, but it has a frequent postoperative recurrence due to early malignant invasion. Chemotherapy and radiotherapy are employed in patients unsuitable for surgery and patients refusing surgery, but they are prone to side effects and poor prognosis (Davis et al., 2019). Therefore, there is an urgent need to seek new treatment methods for MM. Moreover, the potential values of traditional Chinese medicine (TCM) in the treatment of melanoma have gradually been revealed.

Astragalus Membranaceus (AM) and Radix Trichosanthis (RT) are two herbs widely used in TCM to treat different cancers, which have been proven to inhibit tumor growth and metastasis (Zhu et al., 2021; Bian et al., 2022). Chinese herbal compound formula is used most commonly in the clinic, and it is more effective when certain specific herbs and certain ratios are used together. In ancient TCM books, such as "Yixue Zhongzhong Canxilu" written by Zhang Xichun in the Qing dynasty and "Orthodox Manual of External Medicine" written by Chen Shigong in the Ming dynasty, there are also cases of compatibility of AM and RT for disease, especially diabetes. The compatibility of TCM is like mixing a cocktail, so we called it "cocktail" here. Therefore, it has the positive significance of clinical treatment to research the effect of AM, RT, and their "cocktail" on MM cells. However, the specific mechanism of AM and RT in MM treatment is still unclear. In this research, we performed a network pharmacological analysis and evaluated the influence on the tumor growth and proliferation, apoptosis, and migration of MM cells with different concentrations of AM, RT, and their "cocktail," which might provide a reference to clinical applications. We also researched the influence on factors related to apoptosis and migration; the change of LDH activity, SOD activity, and MDA content; and the effect on the Akt-related signaling pathway, which might reflect the mechanisms of the treatment of AM, RT, and their "cocktail."

MATERIALS AND METHODS

Herb Preparation

Every 100 g herb (AM and RT, from Beijing Tongrentang) was soaked in deionized water for 1 h and decocted twice, separately. The decoction was poured into a beaker for 50 ml concentration, transferred, and centrifuged at 1000 g for 20 min. The supernatants, respectively, were the *Astragalus Membranaceus* decoction (AMD) and *Radix Trichosanthis* decoction (RTD).

LC-ESI-MS/MS

The LC-ESI-MS/MS analysis system was equipped with an ultrahigh-performance liquid chromatograph (UHPLC) and an MS system. A Waters ACQUITY UPLC HSS T3 C18 column (2.1 mm × 100 mm, 1.8 μ m) was used at 30°C and 5 ul of injection volume. Gradient elution conditions were set as given in **Supplementary Table S1**. MS parameters were a sheath gas flow of 40 arb, an auxiliary gas flow rate of 15 arb, a capillary temperature of 320°C, an auxiliary gas heater temperature of 350°C, and a positive spray voltage of 3.2 kv. The resolution of MS and MS/MS was 70,000 and 17,500, respectively.

Network Pharmacology

The research process is shown in Supplementary Figure S1. The active chemical compositions of AM and RT were retrieved from the Traditional Chinese Medicine System Pharmacology (TCMSP) database and screened, which conformed to the oral bioavailability $(OB) \ge 30\%$ and drug-likeness $(D)L \ge 0.18$. The potential targets of herbs were further screened in target relative databases and completely reserved. The known potential melanoma-related targets were found with "Homo sapiens" in disease target databases and merged and deleted replicas of all targets. We used the STRING database to establish a protein-protein interaction (PPI) network and visualized the PPI by Cytoscape software. Subsequently, KEGG analysis was performed by linking targets to the Database for Annotation, Visualization, and Integrated Discovery database (DAVID). The drug-target-pathway-disease network was established via Cytoscape software.

MTT Assay

Human malignant melanoma A375 cells and human L-02 liver cells (purchased from Cell Resource Center, Peking Union Medical College) were plated onto 96-well plates with 3×10^3 cells/well. A control group (medium), AMD groups (medium with concentrations 2, 4, 8, 10, and 20 mg/ml of AMD), and RTD groups (medium with concentrations 0.16, 0.32, 0.64, 1.28, and 2.56 mg/ml of RTD) were set and incubated for 48 h or 72 h. After MTT solution and DMSO were used successively, the OD value was estimated at an absorbance of 490 nm. IC₀, IC₂₅, and IC₅₀ (the cell inhibitory rates were 0, 25, and 50%, respectively) were calculated. Afterward, the control group, different concentrations of AMD, RTD, and "cocktail" groups were set up for further tests.

Animal Experiments

Male and female C57BL/6J mice aged 6–8 weeks (purchased from Huafukang Biotechnology) were randomly divided into six groups of six with the abdomen depilated. Following the experimental progress (**Figure 4A**), 0.1 ml PBS was subcutaneously injected at the abdomen of the mice of the normal control group (NCG), while suspension (prepared by 0.1 ml of PBS) with 5×10^5 B16-F10 cells were injected into others. This day was marked as d0. On d2, small black spots greater than 2 mm diameter appeared at the injection site. At d3, the NCG and model control group (MCG) were given ddH₂O intragastric administration, while the three compatible groups were given AMD and RTD solution at a ratio of 5:3 (L 3.12 g/kg, M 6.24 g/kg, and H 12.48 g/kg, respectively), and the positive control group (PCG) was given compound cyclophosphamide solution (19.5 mg/kg) once a day

TABLE 1 Information of compounds of compatibility of AMD and RTD with a ratio of 5 to 3 detected by LC-ESI-MS/MS in different Ion modes.

No.	RT [min]	lon mode	Name	Formula
1	1.292	[M-H] ⁻¹	3-[(Carboxycarbonyl)amino]-L-alanine	C ₅ H ₈ N ₂ O ₅
2	1.551	[M-H] ⁻¹	∟-Glutamic acid	C ₅ H ₉ N O ₄
3	1.555	$[M + H]^{+1}$	∟-Threonine	C ₄ H ₉ N O ₃
4	1.685	$[M + H]^{+1}$	Trigonelline HCl	C ₇ H ₇ N O ₂
5	1.7	$[M + H]^{+1}$	2-Pyrrolidinecarboxylic acid	$C_5 H_9 N O_2$
6	1.718	$[M + H]^{+1}$	Betaine	C ₅ H ₁₁ N O ₂
7	2.428	$[M + H]^{+1}$	∟-Valine	C ₅ H ₁₁ N O ₂
8	2.913	$[M + H]^{+1}$	Adenine	C ₅ H ₅ N ₅
9	2.952	$[M + H]^{+1}$	Nicotinic acid	$C_6 H_5 N O_2$
10	3.574	[M-H] ⁻¹	Citric acid	C ₆ H ₈ O ₇
11	4.03	[M-H] ⁻	Maleic acid	$C_4 H_4 O_4$
12	5.367	$[M + H]^{+1}$	L-lyrosine	$C_9 H_{11} N O_3$
13	5.404	$[M + H]^{-1}$	p-Coumaric acid	$C_9 H_8 O_3$
14	17.872		Asperulosidic acid	$C_{18} H_{24} O_{12}$
15	19.059		4-Hydroxybenzoic acid	$C_7 H_6 O_3$
10	20.156	$[VI + \Pi]$	Asaryiaideniyde	$C_{10} H_{12} O_4$
10	20.228		Sidiricose AS	$C_{22} H_{30} U_{14}$
10	20.257	$[VI + \Gamma]$ [M + \Box] ⁺¹		$C_9 H_{11} N O_2$
19	20.337	$[1 \vee 1 + 1]$	Discusic aciu	
20	20.712			
21	20.094	$[N_{+}]^{+1}$	7-Methoxy.coumarin	$C_{31} \Gamma_{38} C_{17}$
22	20.342	$[M + H]^{+1}$	2-Hydroxy-4-methoxybenzaldebyde	
20	21.047	$[M + H]^{+1}$		Care Hare Our
25	21.773	[M-H] ⁻¹	L-Tryptophan	$C_{22} = 1_{22} = 0_{10}$
26	22.015	$[M + H]^{+1}$	Liquiritigenin	
27	22.010	[M-H] ⁻¹	Liquiritin	C_{01} H ₂ C_{04}
28	22.076	$[M + H]^{+1}$	3 5-Dimethoxy-4-hydroxybenzaldehyde	$C_2 + C_2 = C_3$
29	22.468	$[M + H]^{+1}$	Genistin	C_{21} H ₂₀ O_{10}
30	22.726	$[M + H]^{+1}$	Tectoridin	$C_{22} H_{22} O_{11}$
31	22.878	$[M + H]^{+1}$	Artemisinin	C ₁₅ H ₂₂ O ₅
32	23.187	[M-H] ⁻¹	Phloridzin	$C_{21} H_{24} O_{10}$
33	23.194	$[M + H]^{+1}$	Isosakuranetin	C ₁₆ H ₁₄ O ₅
34	23.522	[M-H] ⁻¹	Azelaic acid	$C_9 H_{16} O_4$
35	23.541	$[M + H]^{+1}$	Jatrorrhizine	C ₂₀ H ₁₉ N O ₄
36	23.66	$[M + H]^{+1}$	Epiberberine	C ₂₀ H ₁₇ N O ₄
37	23.898	[M-H] ⁻¹	Salicylic acid	C ₇ H ₆ O ₃
38	24.055	$[M + H]^{+1}$	6"-O-Acetylglycitin	C ₂₄ H ₂₄ O ₁₁
39	24.219	$[M + H]^{+1}$	Baicalin	C ₂₁ H ₁₈ O ₁₁
40	24.243	$[M + H]^{+1}$	Ononin	C ₂₂ H ₂₂ O ₉
41	24.45	[M-H] ⁻¹	Hydroxygenkwanin	C ₁₆ H ₁₂ O ₆
42	24.534	$[M + H]^{+1}$	Ligustilide	C ₁₂ H ₁₄ O ₂
43	24.98	$[M + H]^{+1}$	Palmatine	C ₂₁ H ₂₁ N O ₄
44	25.005	$[M + H]^{+1}$	Methylnissolin-3-O-glucoside	C ₂₃ H ₂₆ O ₁₀
45	25.011	$[M + H]^{+1}$	4',7-Di-O-methylnaringenin	C ₁₇ H ₁₆ O ₅
46	25.013	$[M + H]^{+1}$	o-Veratraldehyde	C ₉ H ₁₀ O ₃
47	25.239	$[M + H]^{+1}$	Berberine	C ₂₀ H ₁₇ N O ₄
48	25.383	[M-H] ⁻¹	Isomucronulatol 7-O-glucoside	C ₂₃ H ₂₈ O ₁₀
49	25.385	$[M + H]^{+1}$	Benzoic acid	C ₇ H ₆ O ₂
50	25.933	$[M + H]^{+1}$	Calycosin	C ₁₆ H ₁₂ O ₅
51	26.178	$[M + H]^{+1}$	Wogonoside	C ₂₂ H ₂₀ O ₁₁
52	26.329	$[M + H]^{+1}$	Pectolinarigenin	C ₁₇ H ₁₄ O ₆
53	26.448	$[M + H]^{+1}$	Glycitin	C ₂₂ H ₂₂ O ₁₀
54	27.973	$[M + H]^{+1}$	Artemisinic acid	G ₁₅ H ₂₂ O ₂
55	28.624	$[VI + H]^{+1}$	Resibutogenin	$C_{24} H_{32} O_4$
57	20.030		Astragologido IV	
59	23.407	[IVI-F1] [NA , LII+1	Astragaluside IV	
50	3U.U3Z	[IVI + □] [N/I → □]+1	Medicarpin	
60	30.202 21 12/	[N1 ± H] ⁺¹	Wilforlide A	
61	31.595	[M + H] ⁺¹	Roburic acid	
62	31 661	[N] + H] ⁺¹	5-Hydroxy-6.7-dimethoxylflayone	$O_{30} \cap_{48} O_2$
63	32.78	$[M + H]^{+1}$	Chrysosplenetin B	$O_{17} \cap_{14} O_5$ $O_{10} \cap_{14} O_5$
64	32.913	[M-H] ⁻¹	Astragaloside II	C40 H-20 O4-
		f	(Co	ontinued on following page)

No.	RT [min]	lon mode	Name	Formula
65	5 33.378 [M-H] ⁻¹		Cucurbitacin B	C ₃₂ H ₄₆ O ₈
66	35.27	[M+2H] ⁺²	Isoastragaloside I	C ₄₅ H ₇₂ O ₁₆
67	35.283	[M-H] ⁻¹	Dioscin	C ₄₅ H ₇₂ O ₁₆



TABLE 1 | (Continued) Information of compounds of compatibility of AMD and RTD with a ratio of 5 to 3 detected by LC-ESI-MS/MS in different Ion modes.

FIGURE 1 | Mass spectra of AMD, RTD, and compatibility of AMD and RTD in negative and positive ion modes. (A) Mass spectra of AMD in the negative ion mode. (B) Mass spectra of AMD in the positive ion mode. (C) Mass spectra of AMD in the negative ion mode. (D) Mass spectra of RTD solutions in the positive ion mode. (E) Mass spectra of the compatibility of AMD and RTD with a ratio of 5:3 in the negative ion mode. (F) Mass spectra of the compatibility of AMD and RTD with a ratio of 5:3 in the positive ion mode.



FIGURE 2 | Outcomes of network pharmacology analysis predicted. (A) Venn diagram of AM, RT, and MM. (B) PPI network of common targets. The layout of the three circles was in a clockwise direction according to the nodes from large to small and red to purple. (C) KEGG signaling pathways. The larger the area, the more the counts, and the bluer the color, smaller the *p*-value. (D) Drug-target-pathway-disease network. The yellow nodes represent herbs, red of the top 20 targets, orange of 34 pathways, and green of MM.

for 14 consecutive days. The mice weight, tumor length, and transverse diameter were recorded every second day, and the tumors of MCG were photographed every 4 days. Finally, the mice were anesthetized, and the blood, tumor, and liver were collected. The pathological features were determined by H&E staining. All these experiments were in accordance with the guidelines of the Animal Care and Use Committee.

Flow Cytometry Assay and Hoechst Staining

After treating A375 cells with AMD, RTD, and their "cocktail" groups, the cell cycle and apoptosis (processed by an Annexin-FITC Apoptosis Detection Kit) were observed by flow cytometry. In addition, we stained the cells with Hoechst 33342 and photographed them under a confocal microscope (400x image) to observe the apoptotic morphology.

Scratch Wound Healing Assay

After A375 cells were cultured for 24 h, a cross was scraped at the center of wells by using a 10- μ l micropipette tip and photographed to record. Then, the cells were cultured after they were treated with different concentrations of AMD, RTD, and their "cocktail." After 48-h or 72-h treatment, images were captured again.

LDH, SOD, and MDA Assay

The treatment of A375 cells was consistent as mentioned before; then, the cell supernatant (treated for 48 h) and mice serum were



TABLE 2		10	and	10	of		and	RTD	in	oach	action	timo	
IADLE Z	$1 \cup_0$	IC25,	anu	1050	0I	AIVID	anu	שוח	11.1	each	action	ume.	

Drug	Action time (h)	IC ₀ (mg/ml)	IC ₂₅ (mg/ml)	IC ₅₀ (mg/ml)	
AMD	48	1.1	3.6	14	
	72	1.2	4.5	12	
RTD	48	0.137	0.863	2.113	
	72	0.081	0.769	2.020	

collected (treated for 14 days). As per the guidelines of the corresponding kit manufacturer's instructions, the value of LDH, SOD, and MDA was measured. Finally, the results were calculated by using the formula given in the instructions.

Western Blot Assay

The protein in tissues and cells were extracted to determine protein concentration by using a BCA protein detection kit. Western blot experiments for β -actin, Bcl-2, Bax, p53, E-cadherin, N-cadherin, snail, Akt, and p-Akt were performed and photographed by using the Gel imaging instrument. ImageJ software was used to analyze the bands.

Statistical Analysis

The experiments were repeated at least three times, and differences between all data were evaluated using ANOVA, and p < 0.05 was considered statistically significant.

RESULTS

Analysis of Chemical Components in AMD, RTD, and Their Cocktail

The Compound Discoverer 3.1.0.305, mzCloud, and mzVault databases were used to identify unknown compounds. The chemical components are listed in Supplementary Table S2, S3 and Table 1. There were 25 and 49 chemical components of AMD and 14 and 16 components of RTD in the negative and positive ion modes, respectively (Figure 1), while 20 and 47 components were observed in their cocktail at a ratio of 5:3 in different ion modes (Table 1). The AMD compounds include flavonoids, organic acids, and amino acids. Terpenoids, organic acids, and amino acids are the main compounds in RTD. Compounds of the cocktail of AMD and RTD are also flavonoids, organic acids, and amino acids.Interestingly, cocktail contains some unique compounds like berberine and diosmetin-7-O-β-Dglucopyranoside.

Network Pharmacology Analysis Predicted Potential Signaling Pathways of Melanoma Treated by AM and RT

Overall, 20 active chemical compositions of AM and 2 of RT were obtained via the TCMSP database (**Supplementary Table S4**). According to aforementioned databases, there are a total of 691 component targets of herbs, and 901 known potential melanoma-related targets were obtained. After summarizing these targets, 107 common targets were obtained, as shown in the Venn diagram in **Figure 2A**. According to the degree of alignment, the key proteins included SRC, PIK3CA, PIK3R1, Akt1, MAPK1, HRAS, RAC1, MAPK3, HSP90AA1, and PTPN11 (**Figure 2B**). According to KEGG analysis, 34 signaling pathways were obtained (**Figure 2C**). The action pathways mainly include PI3K-AKT, FOXO, and VEGF signaling pathways. The drug-target-pathway-disease network was established to reveal the relationship between them (**Figure 2D**).

AMD, RTD, and Their Compatibility Inhibit the A375 Cell Proliferation

A375 cell proliferation was significantly inhibited by AMD and RTD in a dose-dependent manner (**Figure 3A**), while we found there was a low inhibitory effect on L-02 cells (**Figure 3B**). The corresponding IC₀, IC₂₅, and IC₅₀ of each period were calculated (**Table 2**). They were used to further assess the different effects on A375 cell proliferation with AMD, RTD, and their "cocktail." The proliferation inhibition rate of "cocktail" groups was higher than those of the corresponding AMD and RTD groups (p < 0.05) (**Figure 3C**). Among all the "cocktail" groups, AMD IC₂₅ + RTD IC₅₀ (compatibility in a ratio of 5:3) had a stronger inhibition than others on the A375 cell proliferation (**Figure 3C**).

Compatibility of AMD and RTD Could Inhibit Melanoma *In Vivo*

We had already confirmed that the compatibility group has better effects than the single drug, and the compatibility has an additive effect. Therefore, in the animal experiment, we directly selected the best compatibility ratio (5:3) to detect its effect on mice with malignant melanoma. After B16-F10 cells were injected 2 days later, the tumor began to form—2 mm in diameter (**Figure 4B**). Then the mice body weight in most groups showed an increasing trend, while the PCG obviously decreased compared with NCG (**Figure 4C**), and the spirit and activity of the mice had no obvious change. Compared with MCG, the tumor weight of medication administration groups was decreased significantly (p < 0.05) (**Figure 4D**). The H&E staining results indicated a large area of tumor tissues had disappeared in administration groups (**Figure 4E**), compared with the MCG. Interestingly, there was no significant difference in the pathological sections of liver tissues in all groups.

AMD, RTD, and Their Compatibility Affect the A375 Cell Cycle and Apoptosis

To investigate the mechanism of tumor growth and cell proliferation inhibition, we further analyzed the cell cycle and apoptosis *in vitro*. AMD, RTD, and their "cocktail" could play an inhibitory role in cell proliferation through the induction of cell cycle arrest (**Figure 5**). Furthermore, AMD, RTD, and their "cocktail" could promote A375 cell apoptosis in a dose-dependent manner, and the numbers of early and total apoptotic cells increased significantly in the "cocktail" groups, which showed an additive effect of compatibility (**Figure 6A**).



versus MCG. ${}^{\#}p < 0.05$, compared with NCG.

Hoechst staining results showed that the most nuclear membranes in groups RTD IC_{50} (AMD $IC_0+RTD IC_{50}$) and (AMD $IC_{25} + RTD IC_{50}$) were obviously wrinkled, and the chromatin in nuclei was condensed (**Figure 6B**). Meanwhile, the herbs could reduce Bcl-2 expression and enhance the Bax protein level, with a ratio of Bax/Bcl-2 significantly increased in a dose-dependent manner (**Figures 6C,D**), and the "cocktail"

groups had better effects, especially (AMD IC_0+RTD IC_{50}) and (AMD $IC_{25} + RTD IC_{50}$). In addition, we found that the herbs could also increase p53 protein expression, and group (AMD $IC_{25} + RTD IC_{50}$) had the best effect (**Figure 6E**). In vivo, the compatibility treated in the melanoma model could also promote p53 and Bax protein expression (**Figure 6F**).



resulst. n = 3RTD group.

Cell Migration Could Be Inhibited in A375 Cells Treated With AMD, RTD, and Their Compatibility

The wound healing assay showed that the control group suffered a closing tendency of a scratch wound and the cell number increased, while the others led to slower closing of a scratch wound with statistical difference (**Figure 7A**). Thus, AMD, RTD, and their "cocktail" could inhibit the A375 cell migration in a dose-dependent manner. The

"cocktail" groups had a stronger cell migration inhibition, especially groups (AMD IC_0+RTD IC_{50}) and (AMD IC_{25} + RTD IC_{50}).

Furthermore, in treatment groups of A375 cells, especially "cocktail" groups (AMD IC₀ + RTD IC₅₀) and (AMD IC₂₅ + RTD IC₅₀), E-cadherin increased and N-cadherin decreased, resulting in an increased ratio of E-cadherin/N-cadherin (**Figures 7B,C**). In addition, snail expression of A375 cells decreased (**Figure 7D**). *In vivo*, E-cadherin increased and N-cadherin decreased in tumor tissues treated by the compatibility (**Figure 7E**).







AMD, RTD, and Their Compatibility Affect the LDH, SOD, and MDA Content *in vitro* and *vivo*

LDH is a key enzyme involved in tumor glycolysis and associated with tumor proliferation and metastasis. SOD and MDA are markers of oxidation and antioxidation and are also related to tumor apoptosis and metastasis. Therefore, we examined their expression. After treatment with AMD, RTD, and their "cocktail" for 48 h in A375 cells, the LDH activity of herb-only groups decreased, and the "cocktail" groups (AMD IC_{25} + RTD IC_0) and (AMD IC_{25} + RTD IC_{50}) had an additive effect with a statistical difference (**Figure 8A**). *In vivo*, compared with the NCG, the LDH activity of the MCG significantly increased, and that in other groups decreased (**Figure 8B**). In addition, the MDA content increased and the SOD activity decreased, in a dose-dependent manner. The "cocktail" groups, especially (AMD IC_0 +RTD IC_{50}) and (AMD IC_{25} + RTD IC_{50}), had an additive effect (**Figure 8C, D**).

AMD, RTD, and Their Compatibility Could Alter the Expression of Protein (Akt and p-Akt) in A375 Cells

In vitro, we found that after herb treatment for 48 h, certain doses of AMD and RTD led to a decrease in Akt and p-Akt expression.

For 72 h, herbs, especially "cocktail" groups (AMD IC_0+RTD IC_{50}) and (AMD $IC_{25} + RTD IC_{50}$), could reduce the expression of Akt and p-Akt. *In vivo*, their compatibility also could reduce the Akt protein expression in melanoma tissues (**Figure 9D**).

DISCUSSION

As important qi-tonifying herbs are to be widely used in clinics, AM and RT were originally published in the Chinese medical ancient book Shen Nong's Herbal Classic. Currently, there is much evidence that suggested that AM and RT have antitumor, anti-inflammatory, and antiviral effects, with low expenditure and few side effects (Zhou et al., 2007; Li et al., 2020). LC-MS and network pharmacology analysis pointed out these herbs have multiple components and multiple targets which may exert an antitumor effect. Compared with AM and RT, the cocktail of them possesses some different compounds, such as berberine, which may promote the effects on melanoma and against tumor (Huang et al., 2021). According to our early studies, AMD and RTD possess an antitumor effect because they can inhibit MM growth and metastasis by inhibiting the proliferation and migration of mouse malignant melanoma B16 cells (Zhang



et al., 2020). Furthermore, the PI3K-Akt signaling pathway may play a crucial role in the treatment, and the aforementioned data, which we researched, also confirmed that AMD and RTD could have a restraining effect on mice melanoma and human malignant melanoma A375 cells.

Tumor occurrence is closely associated with unrestricted proliferation, which is also one of the root causes of tumor development. Based on that, directly killing tumor cells or inhibiting proliferation is the main way for antitumor drugs to take effect. In this study, it was confirmed that AMD and RTD could both play a role in inhibiting the A375 cell proliferation while not affecting or slightly affecting L-02 cells. This made it clear that AMD and RTD are noncytotoxic agents at a certain dose, which has great significance in the promotion of their application. Furthermore, *in vivo*, the best compatibility of AMD and RTD could inhibit melanoma growth while not affecting their livers.

Combining with the results of the MTT assay, we found that AMD (below 4 mg/ml) and RTD (below 1.28 mg/ml) had no effect on the normal cell proliferation but caused cell cycle arrest in different phases of A375 cells. Cells that arrest in G_1 and G_2 phases may return to a stationary state (G_0 phase) or undergo apoptosis when the DNA damage is too serious to be repaired, with an irregular nuclear membrane, condensed chromatin, and crumpled nuclear membrane (Hustedt and Durocher 2016;

Lezaja and Altmeyer 2018). These changes could be the featured phenomena of early apoptosis. This research showed that AMD, RTD, and their compatibility could promote early apoptosis in A375 cells, especially the "cocktail" groups.

Antitumor drugs could induce apoptosis by promoting the expression of Bax and p53, or inhibiting the expression of Bcl-2. Similarly, we demonstrated that AMD, RTD, and their "cocktail" could promote apoptosis in A375 cells via modulating Bax, Bcl-2, and p53 expression. *In vivo*, the best compatibility of AMD and RTD also could influence tumors via promoting p53 expression or inhibiting Bcl-2 expression. In summary, we could know that AMD and RTD may achieve a better therapeutic effect on MM by inhibiting proliferation, regulating the cell cycle, and inducing apoptosis. The "cocktail" had an additive antitumor effect.

MM had high metastatic potential. The epithelial-mesenchymal transition (EMT) was a crucial step in the cell invasion and metastasis, with two makers of E-cadherin and N-cadherin. It was confirmed that the upregulation of E-cadherin and suppression of N-cadherin were effective in inhibiting cell migration and invasion (Fenouille et al., 2012; John et al., 2012). In our study, AMD and RTD could inhibit the A375 cell migration by affecting the expression of E-cadherin, N-cadherin, and snail protein. Their "cocktail" also could regulate the expression level of E-cadherin and N-cadherin in mice tissues.



Additionally, one study found that *Astragalus Membranaceus–Curcuma zedoaria* can obviously inhibit tumor cell migration; this was similar to the results from our study (Tan et al., 2019). "Cocktail" of AMD and RTD could markedly inhibit the A375 cell migration, which showed an additive effect.

The growth of cancer cells required glycolysis, and excessive glycolysis could promote cell proliferation and metastasis (Vander Heiden et al., 2009). LDH was a key enzyme during glycolysis to promote malignant behavior via the activation of EMT in cancer cells (Jurisic et al., 2015; Hou et al., 2019), and drugs could decrease glycolysis by reducing the LDH activity, thereby inhibiting cell proliferation and migration. We found that AMD and RTD could reduce the LDH activity of A375 cells, which might be one of the mechanisms for inhibiting the A375 proliferation and migration. The serum LDH of the MCG was higher than that of the NCG, and the abnormal increase in LDH was reduced in each administration group. It meant that the

"cocktail" of AMD and RTD could effectively inhibit melanoma metastasis.

SOD was an important antioxidant biomarker in the organism, and MDA was frequently used as an indicator of cell membrane oxidative damage. Studies had demonstrated that antioxidants can be effective against cell apoptosis and promote the migration of MM cells (Le Gal et al., 2015). In our study, we found that the activity of SOD was decreased, whereas the MDA content was increased in drug groups of A375 cells, especially the "cocktail" groups. This resulted in increased oxidative stress and cell damage, leading to apoptosis.

Akt was an important oncogene closely related to cell proliferation, migration, and metabolism, and there was a significant difference in Akt expressions between MM and common pigmented nevi. The death of MM cells could be promoted by reducing Akt activity and inhibiting Akt phosphorylation (Bhattacharya et al., 2011; Dang et al., 2018). Activated Akt played an important role in both cell apoptosis and



migration. It could not only regulate apoptosis by affecting Bax and Bcl-2 expression but also inhibit the expression of E-cadherin to promote EMT occurrence (Qu et al., 2015; Qin et al., 2017). These all had been confirmed to occur in MM progression. In this study, the results showed that AMD and RTD, especially their "cocktail," could reduce the expressions of Akt and p-Akt protein to restrain their activity and phosphorylation. It may be the mechanism for AMD, RTD, and their "cocktail" to regulate the proliferation, cell cycle, apoptosis, and migration of A375 cells. Moreover, activated Akt can also affect the production of ROS, thereby affecting the survival of MM cells (Hambright et al., 2015). In conclusion, the present study explored the antitumor effect on MM *in vitro* and *in vivo* of AMD, RTD, and their compatibility. The results clarified that AMD, RTD, and their "cocktail" could exert an antitumor effect by regulating the Aktrelated signaling pathway to influence the cell proliferation cycle, apoptosis, migration, LDH activity, SOD activity, and MDA content (**Figure 10**). The additive effect of compatibility *in vitro* was confirmed, and the best compatibility group obtained by *in vitro* experiments could indeed inhibit the growth of mouse melanoma *in vivo*. These provide the experimental basis and new ideas for the clinical treatment with TCM in MM.

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article are available from corresponding author upon reasonable request.

ETHICS STATEMENT

The animal study was reviewed and approved by the Medical and Experimental Animal Ethics Committee of the Beijing University of Traditional Chinese Medicine.

AUTHOR CONTRIBUTIONS

QYZ: validation, formal analysis, and writing—original draft. LG: validation and formal analysis. SLH: formal analysis and writing—original draft. YXL: investigation. JYH: investigation.

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SUPPLEMENTARY MATERIAL

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