

# Effective Components-Targets-Mechanism-Chinese Prescription Strategy (ETMC), A New Strategy for Chinese Prescription Development: A Case Study in Danlou Tablet

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## Research

**Keywords:** ETMC, Chinese prescription, Effective Substance, Targets, Mechanism, Danlou tablet

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# Effective Components-Targets-Mechanism-Chinese Prescription Strategy (ETMC), A New Strategy for Chinese Prescription Development: A Case Study in Danlou Tablet

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8 **Keywords:** ETMC, Chinese prescription, Effective Substance, Targets, Mechanism, Danlou  
9 tablet

## 10 Abstract

11 **Background:** Chinese prescription is a combination medicine used by Chinese medical workers to  
12 treat various diseases and has saved countless human lives. By studying the effective substances,  
13 therapeutic targets, and mechanism of Chinese prescription, Western researchers can better  
14 understand the great value of Chinese prescription.

15 **Methods:** In this study, a new strategy was proposed for the development of Chinese prescription:  
16 the effective components (E)-targets (T)-mechanism (M)-Chinese prescription (C) strategy, namely  
17 the ETMC strategy.

18 **Results:** The proposed strategy used the chemical compositions of the Chinese prescription as the  
19 source of ligands to predict the corresponding targets for discovering the mechanism of Chinese  
20 prescription, which was helpful to further screen out effective substances from the chemical  
21 compositions of Chinese prescription. Here an example on development of Danlou tablet was  
22 performed to introduce the application of ETMC.

23 **Conclusions:** A novel strategy for Chinese prescription development, ETMC, along with its  
24 application was provided in the current study, which contributed to the further development, wider  
25 application and internationalization of Chinese prescription.

26

## 27 Background

28 Chinese prescription (CP), also known as Chinese formula (CF), is a special kind of medicine that  
29 has been clinically used in China for thousands of years, contributed to the prosperity and civilization  
30 of the China and the surrounding countries (Luan et al., 2020). Its advantages of definite curative  
31 effects and acceptable adverse effects have gradually attracted the attention of Western medical  
32 workers on CP (Xiong et al., 2017). But the concern of Western medical workers on CP has not yet

33 turned into full recognition, due to the lack of clear effective components, specific targets and exact  
34 therapeutic mechanism for CP and even Chinese medicine (Chan et al., 2010; Hao et al., 2017). With  
35 the rise of network pharmacology, this disadvantaged situation of CP has been partially improved  
36 (Luo et al., 2020; Tao et al., 2013). However, the range of effective components, targets, and  
37 mechanisms of the CP predicted by the classic network pharmacology strategy is too large to be  
38 further narrowed, which makes it difficult to choose a suitable research direction from it, which  
39 hinders the basic research and modernization of CP. However, the effective ingredients, targets, and  
40 mechanisms determined by this strategy of studying the effective ingredients and mechanisms of  
41 traditional Chinese medicine compound prescriptions through network pharmacology is too large to  
42 be further narrowed, which makes it difficult for medical workers to choose a suitable research  
43 direction from them, thereby Inhibit basic research and modernization of CP. Therefore, it is  
44 imperative to propose a new CP development strategy that can help focus research directions.

45 Here, we propose the effective components (E)-targets (T)-mechanism (M)-Chinese prescription (C)  
46 strategy, abbreviated as ETMC. ETMC refers to a novel CP development strategy and its brief steps  
47 are as follows: Firstly, the chemical compositions of the CP are used as the source of ligands (also  
48 the candidate effective components) to reversely fish the corresponding targets. Targets obtained in  
49 the preceding step are further analyzed by bioinformatics technology to determine a number of  
50 physiological and pathological functions-related signal pathways. Then molecular docking is  
51 performed between all chemical compositions of CP and targets on different signal pathways.  
52 According to the molecular docking results, a few chemical compositions that are superior to the  
53 original ligand of the target are screened out from all chemical component of CP, which considered  
54 as effective components of CP. Similarly, corresponding targets of these effective components are  
55 also screened out from all the targets, which considered as specific targets of CP. Besides, an  
56 example on development of Danlou tablet (DLT), one of the commercial CP against cardiovascular  
57 diseases was performed to better demonstrate the application of ETMC strategy.

58

## 59 **Methods**

### 60 **Collection of Chemical Compositions of DLT**

61 The collection of chemical compositions of DLT was based on the previous research (Dong et al.,  
62 2013). Then compositions mentioned above were further classified according to their herbal source  
63 which determined via Chinese medicine and chemical composition database from Shanghai Institute  
64 of organic chemistry of CAS [<http://www.organchem.csdb.cn>].

### 65 **Target Prediction and Selection**

66 Target prediction of chemical compositions of DLT was performed via SwissTargetPrediction  
67 webtool [<http://www.swisstargetprediction.ch>] (Daina et al., 2019). The most probable  
68 macromolecular targets of chemical compositions of DLT were estimated which founded on a  
69 combination of 2D and 3D similarity with a library of 370'000 known actives on more than 3000  
70 proteins from three different species. Then intersection analysis was performed on targets predicted  
71 by different chemical compositions of DLT, and those targets that had at least five intersections were  
72 selected for subsequent research.

### 73 **Bioinformatics Analysis**

74 Protein-protein interaction (PPI) information of selected targets mentioned above was evaluated by  
 75 the online tool STRING (Search Tool for the Retrieval of Interacting Genes) (Szklarczyk et al., 2015).  
 76 Gene ontology analysis (GO) was performed via STRING and verified by an online bioinformatic  
 77 tool DAVID [<https://david.ncifcrf.gov/home.jsp>] (Huang et al., 2009).

## 78 **Molecular Docking**

79 The three-dimensional geometric coordinates of the X-ray crystal structures of selected targets,  
 80 ACHE (PDB ID: 4ey7), ADORA1 (5n2s), AKR1B1 (1ah3), AKR1B10 (1zua), ALOX15 (1lox),  
 81 CBR1 (1wma), CYP19A1 (3s7s), EGFR (3lzb), ESR1 (1qku), FYN (2dq7), HSD17B1 (6mnc), MET  
 82 (3cth), PTGS1 (2oye), SQLE (6c6p), TTR (6e6z) were obtained from the Protein Data Bank (PDB).  
 83 Molecular docking was performed by Accelrys Discovery Studio (version 3.0; Accelrys, San Diego,  
 84 CA, USA). Chemical compositions of DLT were energy minimized with the CHARMM force field  
 85 and the CDOCKER protocol was performed for semiflexible molecular docking. Based on the  
 86 docking results, chemical compositions whose –CDOCKER Interaction Energy scores were greater  
 87 than the targets' original ligands score were screened out as effective components of DLT.

88

## 89 **Results**

### 90 **Collection of Chemical Compositions of DLT**

91 Dan-Lou Tablet (DLT), a commercial CP developed from *Gualou Xiebai Baijiu Tang* has been  
 92 clinically used for the treatment of cardiovascular diseases such as Coronary disease, myocardial  
 93 infarction and so on (Mao et al., 2016). DLT is composed of ten herbs: *Trichosanthes kirilowii*  
 94 Maxim. (Gualoupi), *Allium macrostemon* Bge. (Xiebai), *Pueraria lobata* ( Willd. ) Ohwi (Gegen),  
 95 *Salvia miltiorrhiza* Bge. (Danshen), *Astragalus membranaceus* ( Fish. ) Bge. var. mongholicus  
 96 ( Bge. ) Hsiao (Huangqi), *Alisma orientalis* ( Sam. ) Juzep. (Zexie), *Drynaria fortune* ( Kunze ) J. Sm.  
 97 (Gusuibu), *Ligusticum chuanxiong* Hort. (Chuanxiong), *Paeonia lactiflora* Pall. (Chishao) and  
 98 *Curcuma longa* L. (Yujin) (Li et al., 2019b). According to the previous report, the chemical  
 99 compositions of DLT were classified according to their herbal source (**Figure 1 and Supplementary**  
 100 **Figure 1**). The classification results showed that the chemical compositions in DLT were mainly  
 101 derived from the Minister drugs Gegen and Danshen, and few chemical compositions from the  
 102 Monarch drug Gualoupi and Minister drug Yujin were detected in DLT.

### 103 **Prediction and Selection of Targets Corresponding to Chemical Compositions of DTL**

104 Next, target fishing was performed to predict corresponding targets of all chemical compositions of  
 105 DLT. The results showed that a total of 529 targets were predicted, including the same target  
 106 predicted by different chemical components. Chinese medicine has the advantage of “multiple targets  
 107 effect”. And the “multiple targets effect” here not only refers to multiple targets corresponding to  
 108 multiple chemical components, but also refers to a single target corresponding to multiple chemical  
 109 components. According to the "single-target superposition" theory of Chinese medicine, different  
 110 chemical compositions of traditional Chinese medicine can be combined with a single target one after  
 111 another in superposition effects of concentration and time differences, which is one of the key  
 112 reasons why traditional Chinese medicine can play a highly effective and long-lasting therapeutic  
 113 effect (Cai et al., 2015). Therefore, intersection analysis was performed on targets predicted by  
 114 different chemical compositions of DLT, and those targets that had at least five intersections were  
 115 selected as key targets for subsequent research (Figure 2).

## 116 **Protein–protein Interaction Network (PPI) and Gene Ontology (GO) Analysis**

117 In order to clarify the connection between key targets selected above and biological functions, PPI  
118 network complex was constructed with selected key targets and their corresponding chemical  
119 compositions of DLT (Figure 3). Gene ontology analysis (GO) is a commonly used for defining  
120 genes or protein product to identify unique biological properties of high-throughput transcriptome or  
121 genome data (Ashburner et al., 2000). Therefore, GO analysis was performed after construction of  
122 PPI network. Results showed that about half of all key targets were particularly enriched in  
123 regulation of lipid metabolic process and the oxidation-reduction process, which were the two main  
124 biological processes that DLT could potentially regulate (Figure 3).

## 125 **Effective Components Screening from Chemical Compositions of DLT**

126 Furthermore, effective components of DLT were screened out from chemical compositions of DLT  
127 via molecular docking. Results showed that certain targets were regulated by corresponding chemical  
128 components of DLT, whose regulatory effects were better than the original ligands of the targets  
129 (Figure 4 and Supplementary Table 1). So far, the number of potential targets of DLT has been  
130 significantly limited, and effective components of certain targets were determined among several  
131 chemical compositions of DLT.

## 132 **The Relationship between Key Targets and Effective Components of DLT against** 133 **Cardiovascular Diseases**

134 Multiple targets were reported to be associated with cardiovascular diseases (CD), including potential  
135 targets of DLT, ALOX15 (myocardial infarction), ACHE (coronary diseases), ESR1 (coronary  
136 diseases), ADORA1 (myocardial ischaemia), AKR1B1 (myocardial ischaemia) and AKR1B10  
137 (myocardial ischaemia) (Halade et al., 2017; Işık et al., 2019; Li et al., 2019a; Louttit et al., 1999;  
138 Ananthakrishnan et al., 2009). Here, by ETMC strategy, it was demonstrated that the anti-  
139 cardiovascular effects of Chinese prescription DLT were due to its regulation on CD-related targets  
140 by eight effective components of DLT (Figure 5).

141

## 142 **Discussion**

143 Chinese medicine has gradually attracted widespread attention from Chinese and Western medical  
144 researchers due to its excellent efficacy and minimal side effects. Revealing the effective components  
145 of Chinese medicine not only helps the further development of Chinese medicine, but also facilitates  
146 the modernization and internationalization of Chinese medicine. In this study, a new strategy for  
147 Chinese prescription development, ETMC, was proposed to screen out the effective components with  
148 potential therapeutic effects from the complex chemical compositions of traditional Chinese medicine,  
149 which contributed to clear directions for Chinese medicine researches.

150 CP is a special administration mode of Chinese medicine based on the theory of traditional Chinese  
151 medicine, which is a combination of different Chinese medicines to exert better, longer-lasting and  
152 safer effects than using a single Chinese medicine alone. However, the unclear pharmacodynamic  
153 mechanism and targets of CP limits the expansion of its indications, and this obstacle can be  
154 eliminated through the proposed ETMC strategy, a combined application of computer simulation  
155 technology, network pharmacology and bioinformatics. ETMC strategy is beneficial for

156 determination of the pharmacodynamic targets corresponding to effective components in CP. And  
157 biological processes that involve these targets are useful to determine the mechanism CP.

158 Multiple studies reported that the association of chemical compositions of CP with potential targets  
159 can be predicted through classic network pharmacology research process. However, it is difficult to  
160 determine fewer but more valuable effective components and key targets from a large number of  
161 chemical components and potential targets in the prediction results of network pharmacology. In the  
162 example of applying the ETMC strategy in this study, a few effective components of DLT and their  
163 targets closely related to cardiovascular diseases were identified. Most of effective components of  
164 DLT are able to regulate multiple targets, which was consistent with "single-target superposition"  
165 theory of Chinese medicine. And targets corresponding to effective components of DLT have also  
166 been reported to be closely related to cardiovascular disease, the indication for DLT. Therefore, the  
167 ETMC strategy, as a simple and important new CP development strategy, needs to be promoted.

168

## 169 **Conslusions**

170 A novel strategy for CP development, ETMC, along with its application was provided in the current  
171 study, which contributed to the further development, wider application and internationalization of CP.

172

## 173 **Abbreviations**

174 CP, Chinese prescription; CF, Chinese formalua; ETMC, effective components-targets-mechanism-  
175 Chinese prescription strategy; DLT, Danlou tablet; PPI, protein-protein interaction; STRING, search  
176 tool for retrieval of interacting genes; GO, gene ontology.

177

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179 Not applicable.

180

## 181 **Author's contributions**

182 YG conceived the concept of this article. YG performed all the experiments and wrote the manuscript.  
183 YJ designed and made all figures. YG gave comprehensive advice and critically revised the  
184 manuscript. All authors approved the final version of this manuscript.

185

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188

**189 Availability of data and materials**

190 The datasets generated and/or analysed during the current study are available from the corresponding  
191 author on reasonable request.

192

**193 Ethics approval and consent to participate**

194 Not applicable.

195

**196 Consent for publication**

197 Not applicable.

198

**199 Competing interests**

200 The authors declared no competing financial or commercial conflict of interest.

201

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## 267 **FIGURE LEGENDS & TABLES**

### 268 **Figure legends**

269 **Figure 1 Collection and classification of chemical compositions of DLT according to their**  
 270 **herbal source.** **1**, adenosine; **2**, gallic acid; **3**, danshensu; **4**, 5-HMF; **5**, protocatechuic acid; **6**,  
 271 daidzein; **7**, 3'-hydroxyl puerarin; **8**, chlorogenic acid; **9**, protocatechuic aldehyde; **10**, puerarin; **11**,  
 272 mirificin; **12**, 3'-methoxy puerarin; **13**, genistein; **14**, paeoniflorin; **15**, ethyl gallate; **16**, calycosin; **17**,  
 273 ferulic acid; **18**, genistin; **19**, salvianolic acid; **20**, formononetin; **21**, 1,3-dicaffeoylquinic acid; **22**, 4'-  
 274 methoxy puerarin; **23**, naringin; **24**, salvianolic acid E; **25**, rosmarinic acid; **26**, tanshindiol C; **27**,  
 275 salvianolic acid B; **28**, salvianolic acid C; **29**, benzoyl paeoniflorin; **30**, 11-anhydro-alisol F; **31**,  
 276 alisol C; **32**, senkyunolide A; **33**, dihydrotanshinone I; **34**, tanshinone I; **35**, cryptotanshinone

277 **Figure 2 The compounds-targets network of chemical compositions of DLT and their key**  
 278 **targets.** The yellow circles represented chemical compositions of DLT. The pink rectangles  
 279 represented the protein targets. Every chemical composition of DLT and its predicted targets were  
 280 connected by grey straight lines.

281 **Figure 3 PPI network constructed with key targets of DLT and GO analysis.** Red cycles  
 282 represented genes involved in lipid metabolic process (GO-term, GO: 0006629; count in gene set, 20  
 283 of 1192; false discovery rate, 9.01E-08), blue cycles represented genes involved in oxidation-  
 284 reduction process (GO-term, GO: 0055114; count in gene set, 16 of 923; false discovery rate, 1.67E-  
 285 06).

286 **Figure 4 Modulatory effects of effective components of DLT on key targets involved in**  
 287 **biological processes regulated by DLT.** All the key targets of DLT were divided into two parts  
 288 based on the biological processes they were involved in. The pink rectangles represented the key  
 289 targets of DLT. The yellow rectangles below each pink rectangle represented effective components  
 290 of DLT that could regulate the key target.

291 **Figure 5 ETMC network diagram of DLT against cardiovascular diseases.** The relationship  
 292 between herbal source, effective components, potential targets and indications of DLT were shown in  
 293 this diagram. Green lines linked effective components of DLT (yellow rectangles) to their herbal  
 294 source (brown rectangles). Red lines linked effective components of DLT (yellow rectangles) to their  
 295 targets (pink rectangles). All the targets were classified according to their related diseases.

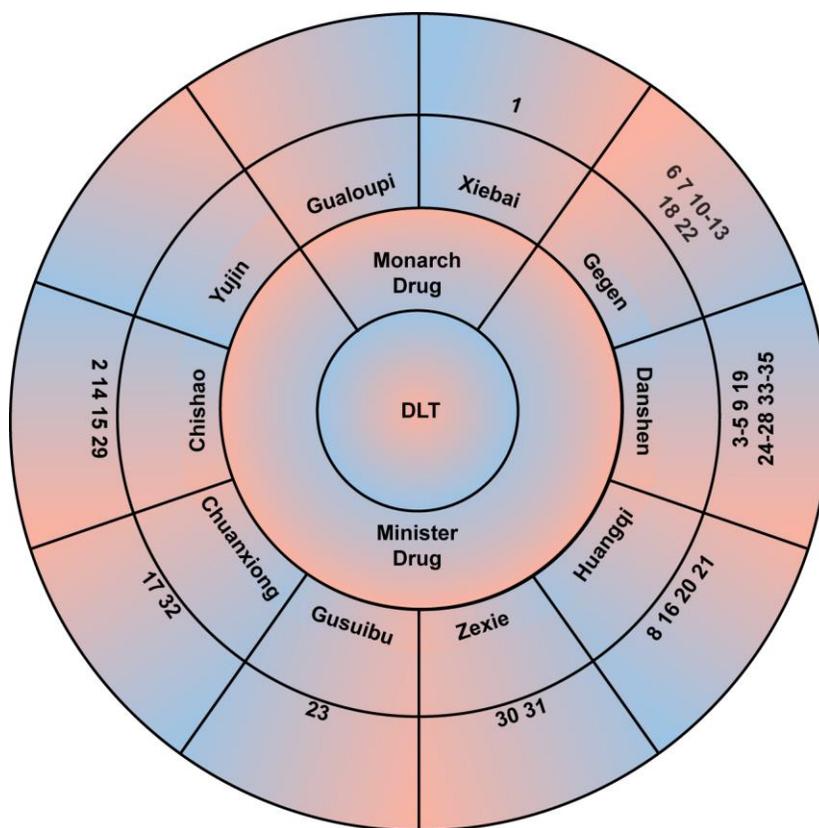
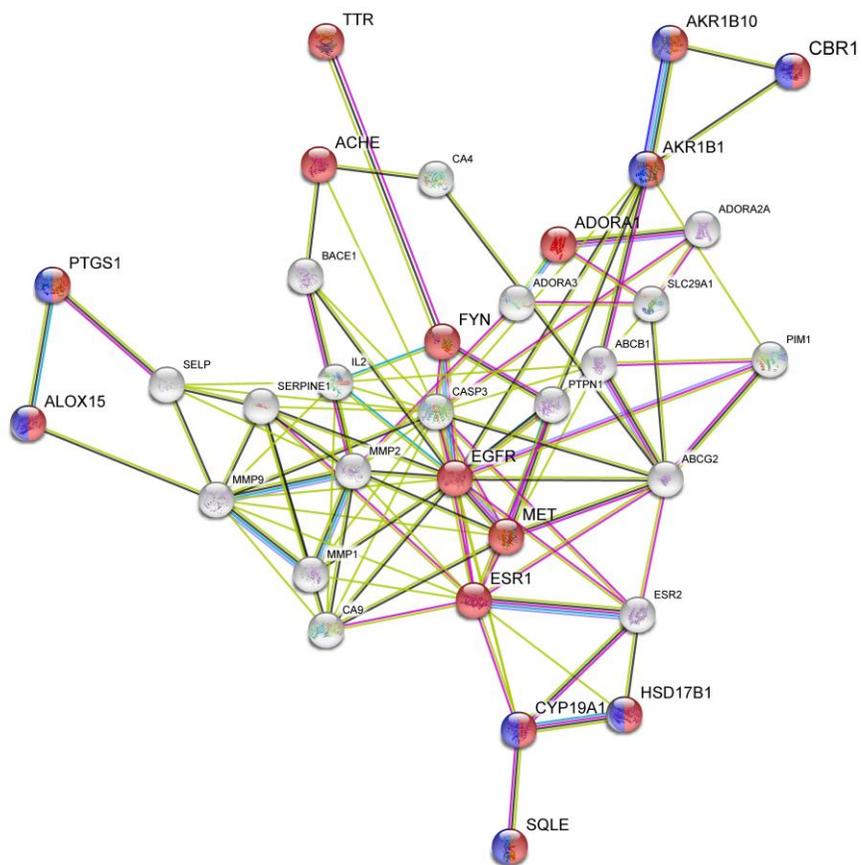


Figure 1

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**Figure 3**

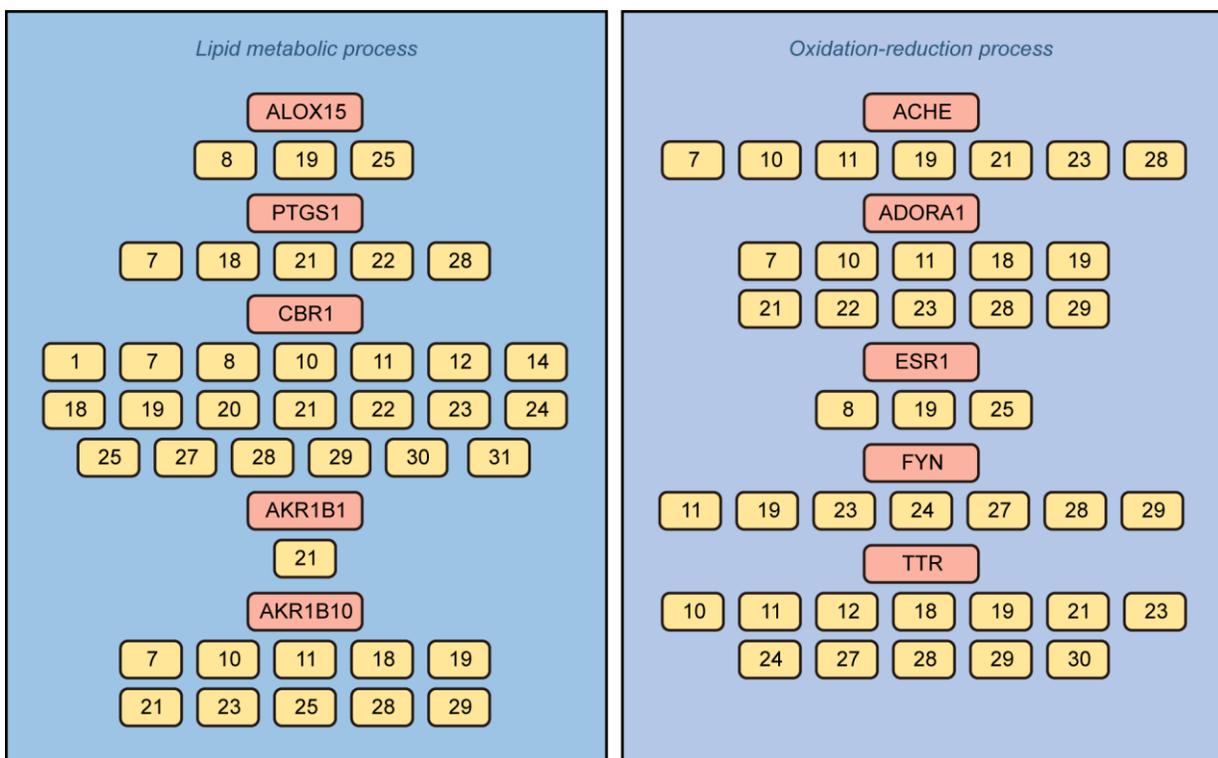


Figure 4

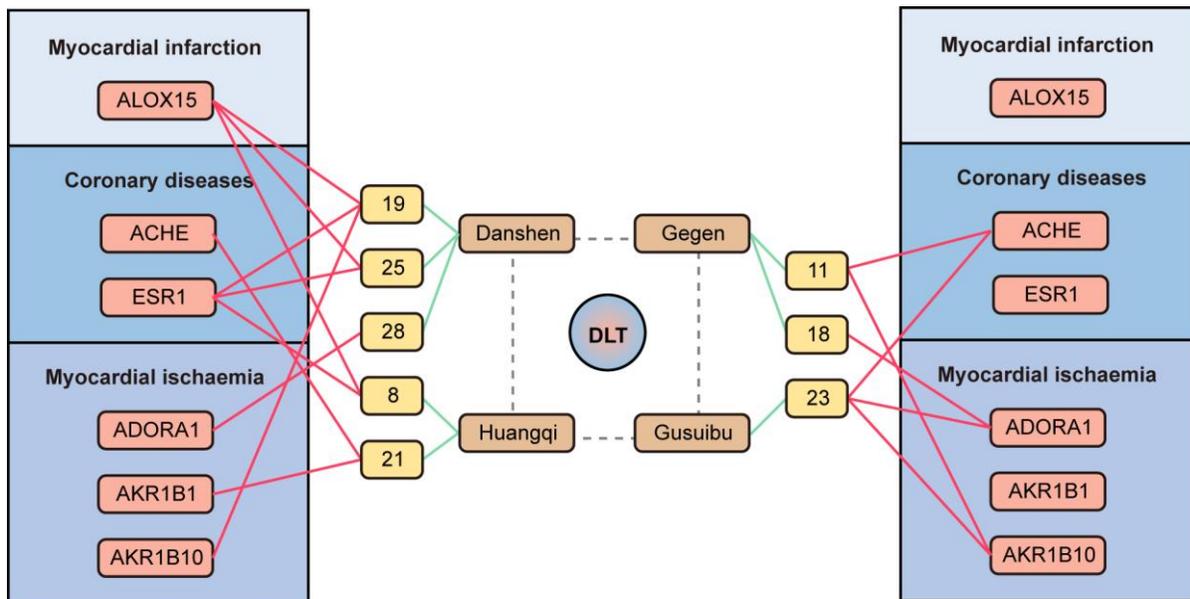


Figure 5

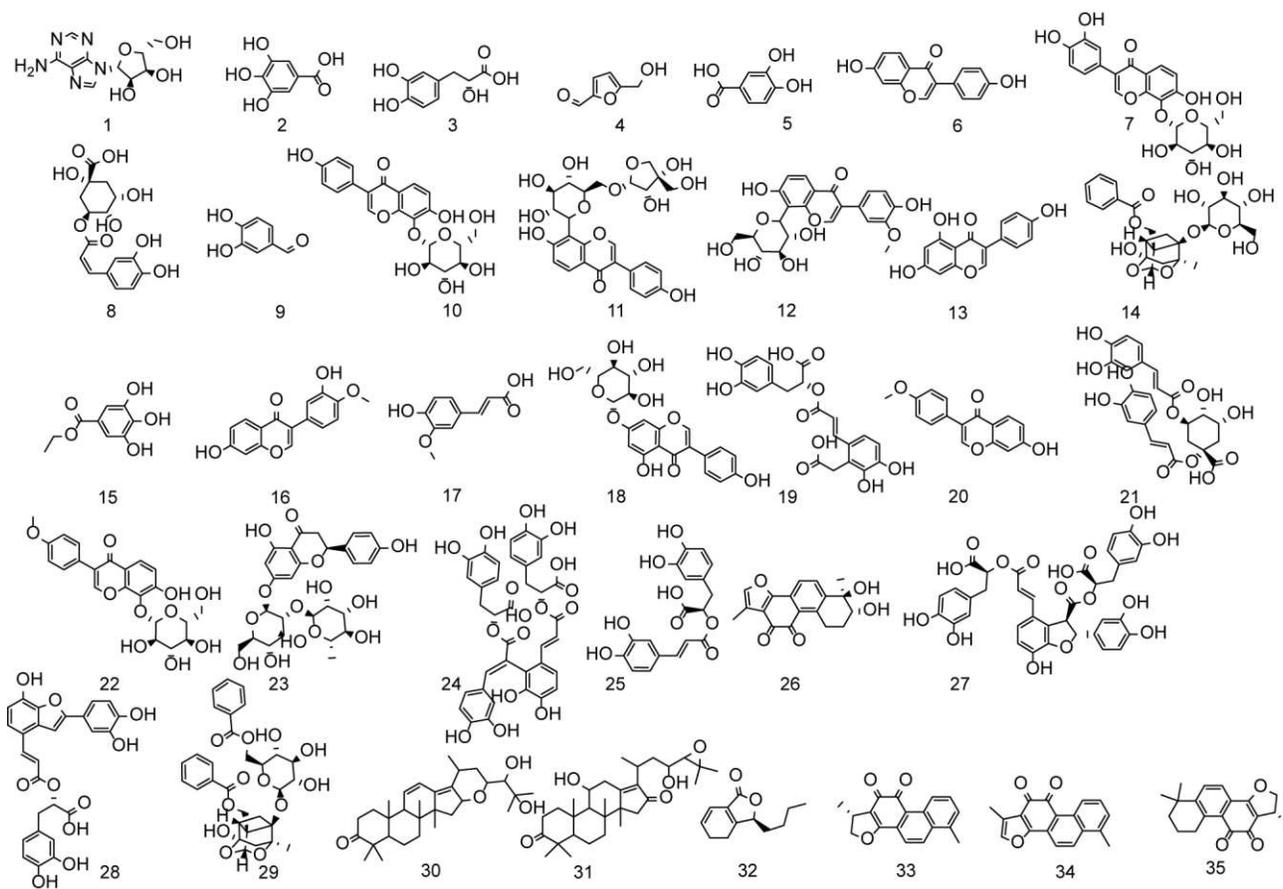
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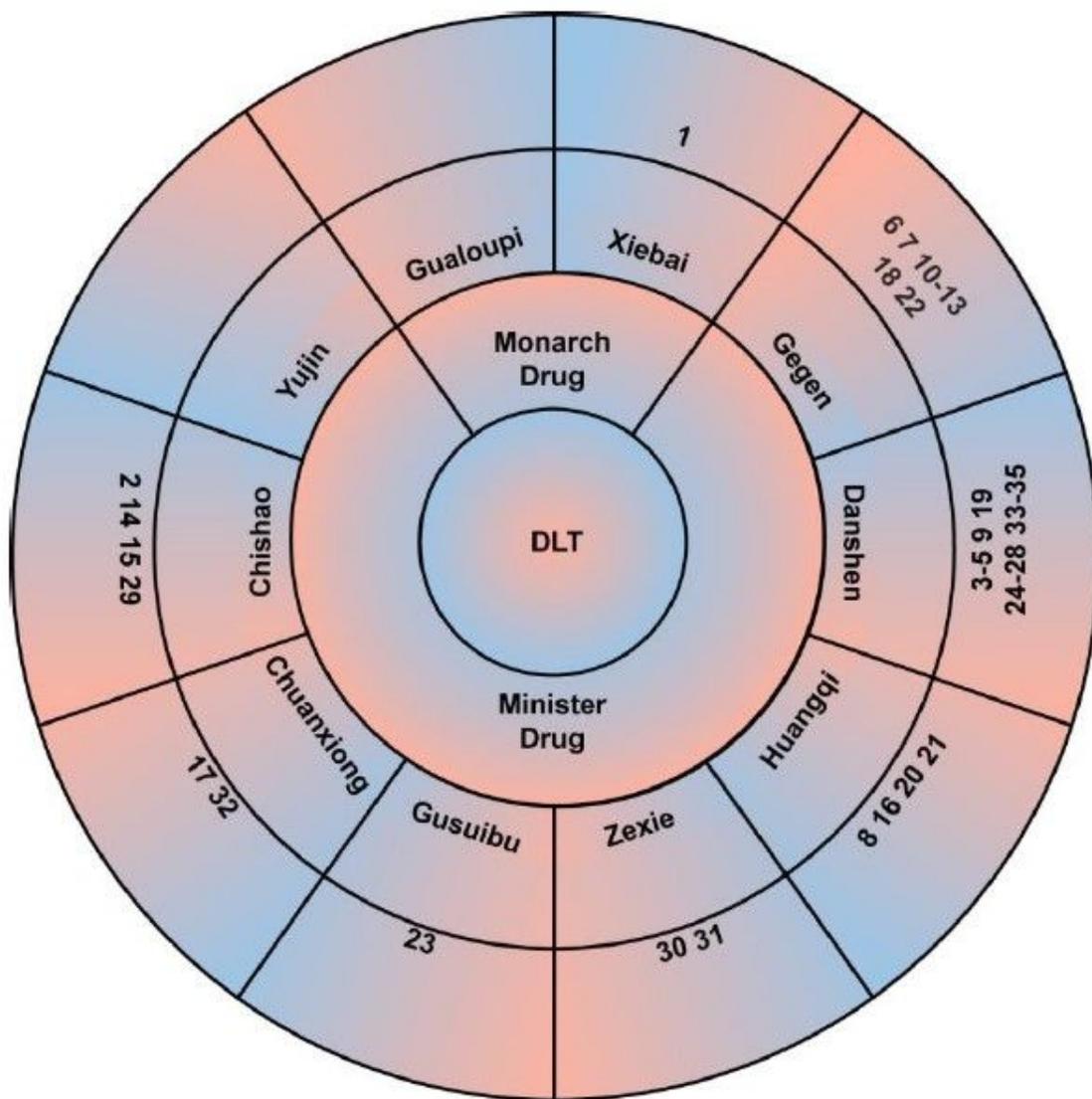


Supplementary Figure 1

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308

# Figures

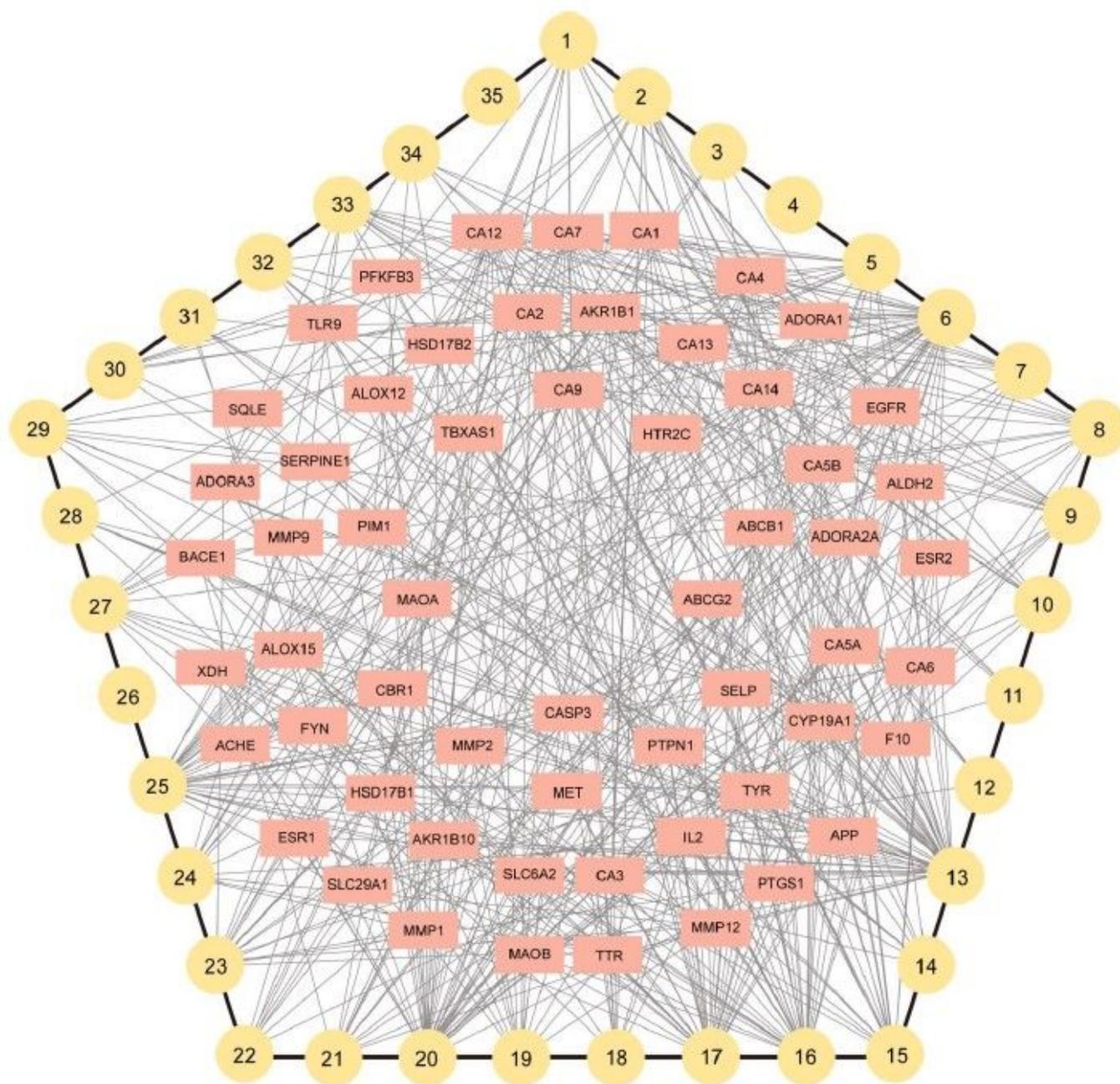


**Figure 1**

## Figure 1

Collection and classification of chemical compositions of DLT according to their herbal source. 1, adenosine; 2, gallic acid; 3, danshensu; 4, 5-HMF; 5, protocatechuic acid; 6, daidzein; 7, 3'-hydroxyl puerarin; 8, chlorogenic acid; 9, protocatechuic aldehyde; 10, puerarin; 11, mirificin; 12, 3'-methoxy puerarin; 13, genistein; 14, paeoniflorin; 15, ethyl gallate; 16, calycosin; 17, ferulic acid; 18, genistin; 19, salvianolic acid; 20, formononetin; 21, 1,3-dicaffeoylquinic acid; 22, 4'-methoxy puerarin; 23, naringin; 24, salvianolic acid E; 25, rosmarinic acid; 26, tanshindiol C; 27, salvianolic acid B; 28, salvianolic acid C; 29,

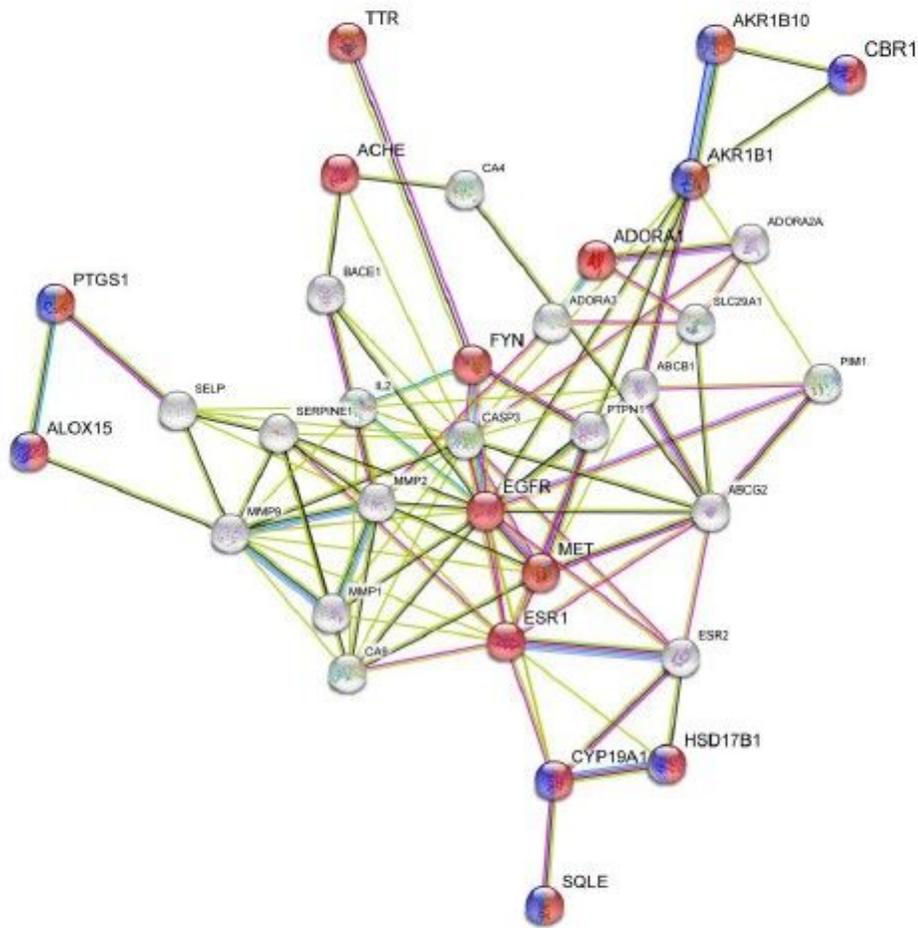
benzoyl paeoniflorin; 30, 11-anhydro-alisol F; 31, alisol C; 32, senkyunolide A; 33, dihydrotanshinone I; 34, tanshinone I; 35, cryptotanshinone



**Figure 2**

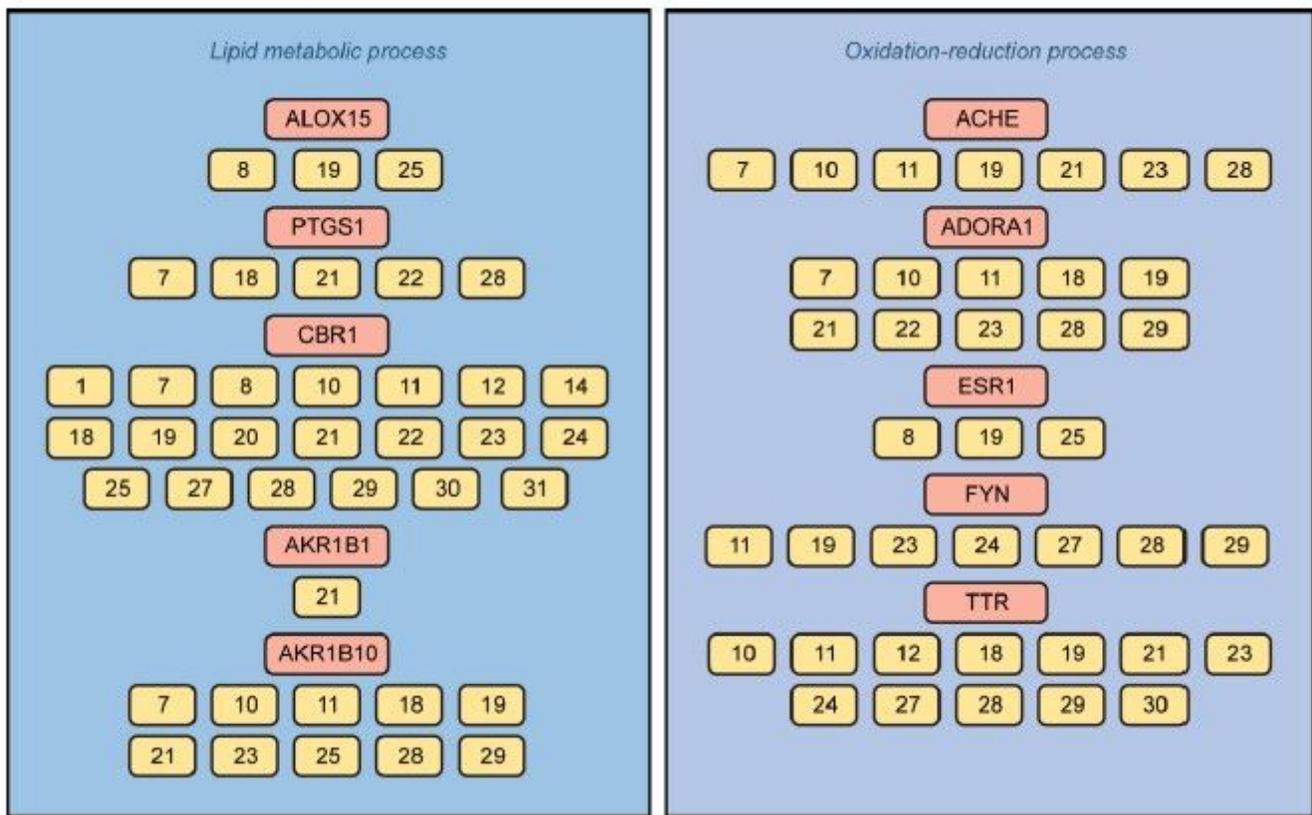
## Figure 2

The compounds-targets network of chemical compositions of DLT and their key targets. The yellow circles represented chemical compositions of DLT. The pink rectangles represented the protein targets. Every chemical composition of DLT and its predicted targets were connected by grey straight lines.



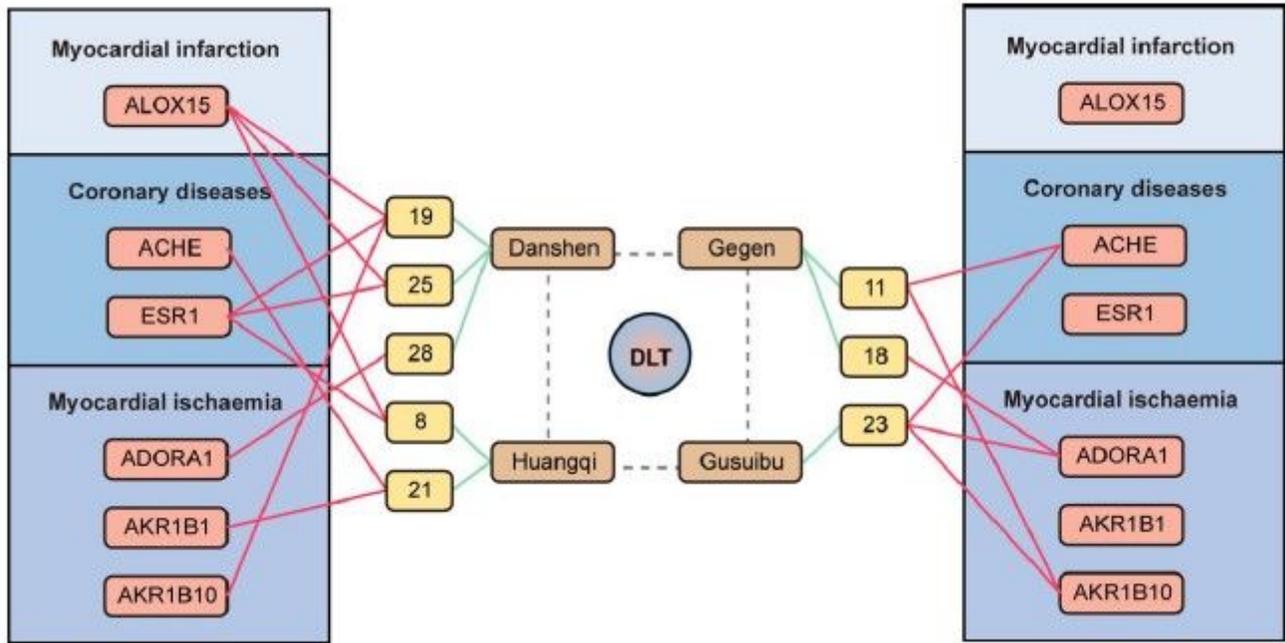
**Figure 3**

PPI network constructed with key targets of DLT and GO analysis. Red cycles represented genes involved in lipid metabolic process (GO-term, GO: 0006629; count in gene set, 20 of 1192; false discovery rate,  $9.01E-08$ ), blue cycles represented genes involved in oxidation-reduction process (GO-term, GO: 0055114; count in gene set, 16 of 923; false discovery rate,  $1.67E-06$ ).



**Figure 4**

Modulatory effects of effective components of DLT on key targets involved in biological processes regulated by DLT. All the key targets of DLT were divided into two parts based on the biological processes they were involved in. The pink rectangles represented the key targets of DLT. The yellow rectangles below each pink rectangle represented effective components of DLT that could regulate the key target.



**Figure 5**

ETMC network diagram of DLT against cardiovascular diseases. The relationship between herbal source, effective components, potential targets and indications of DLT were shown in this diagram. Green lines linked effective components of DLT (yellow rectangles) to their herbal source (brown rectangles). Red lines linked effective components of DLT (yellow rectangles) to their targets (pink rectangles). All the targets were classified according to their related diseases.