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Abstract

Traditional Chinese medicines (TCMs) have been used in clinic for thousands of years. Their reliable therapeutic efficacies are closely related to their multi-target mechanisms of action (MOAs). Discovery of these targets is important for understanding their clinical features. Danshen is the dried root of *Salvia miltiorrhiza*, a traditional Chinese medicine (TCM) used for the treatment of cardiovascular and cerebrovascular diseases. Although its clinical features are well recognized, targets of its active constituents are poorly understood. Here, label-free cell phenotypic assay was used to investigate the potential targets of phenolic acids in Danshen. Pharmacological profiling of 10 known phenolic acids in Danshen using HT-29 and A431 cells revealed that lithospermic acid, salvianolic acid A, salvianolic acid B, salvianolic acid C and Danshensu all displayed agonistic activity at the GPR35; however, salvianolic acid A and salvianolic acid C at high doses were also active to alter intracellular Ca^{2+} via another unknown target. Since GPR35 has been implicated in inflammation and cardiovascular diseases, the discovery of GPR35 as one target of phenolic acids in Danshen was useful for elucidating their mechanisms of action in the treatment of these diseases. This study also highlights the potential of label-free cell phenotypic assay for discovering multiple targets of TCM.

Key words: label-free cell phenotypic assay, multiple targets, phenolic acids, Danshen

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1. Introduction

Traditional Chinese medicines (TCMs) are attracting considerable attention owing to their reliable clinical efficacy, thus being important resources for lead compounds 51 or drugs.^{1, 2} Most of them provide therapeutic effect by interacting with multiple targets. For instance, paclitaxel (anticancer), reserpine (antihypertensive), artemisinin (antimalarial) and rapamycin (immunosuppression) act on at least two targets. Studying these mechanisms of action (MOAs) is important for understanding their clinical features. Target-based approaches that screen compounds against a single 56 target protein are not effective for discovering multiple targets of compounds.³ While phenotype-based approaches can provide an unbiased way to find active compounds, which are useful for identifying multiple targets or pathways.⁴

As one of new generation phenotype-based assay techniques, label-free cell phenotypic assay has become an attractive approach for determining target 61 engagement of drugs.⁵⁻⁷ This assay uses label-free resonant waveguide grating (RWG) biosensor to convert drug-induced dynamic redistribution of cellular constituents in cells into an integrated and kinetic response, termed dynamic mass redistribution 64 (DMR).¹⁴ The DMR is recorded as a shift in resonant wavelength in picometer (pm) and represents a cell phenotypic response covering a wide range of 66 targets/pathways.⁶⁻⁹ Furthermore, due to its non-invasiveness, DMR assay can be performed in multiple formats and permits intervention with probe molecules, thus 68 enabling mechanistic elucidation of drug pharmacology.¹⁰⁻¹²

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Here we use label-free cell phenotypic assay to discover new targets of phenolic acids from Danshen with a hope to elucidate its *in vivo* therapeutic effects.

2. Experimental

2.1. Materials

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Acetylcholine chloride, cyclopiazonic acid, danshensu, lithospermic acid, salvianolic acid A, salvianolic acid B, salvianolic acid C and thapsigargin were stocked in 100 mM, while all other compounds were stocked in 10 mM. Except for acetylcholine which was prepared in water, all others compounds were dissolved in 100% dimethyl sulfoxide (DMSO). All compounds were diluted freshly by the assay

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2.2. Cell culture

HT-29 cells were cultured using McCoy's 5A Medium (#DY0324S0414L, Sango Biotech, Shanghai, China) with 10% fetal bovine serum (#10099141, Gibco, Life Technologies), 50 µg/mL penicillin and 100 µg/mL streptomycin in a humidified 37° C/5% CO₂ incubator. A431 cells were cultured in Dulbecco's Modified Eagle's Medium (DMEM) (#SH30022.01B, Thermo Scientific HyClone) supplemented with 142 10% fetal bovine serum, 50 μ g/mL penicillin and 100 μ g/mL streptomycin at 37°C 143 under air/5% $CO₂$.

2.3. Dynamic mass redistribution assay

146 All DMR assays were performed using an $Epic^{\circledR}$ BT system (Corning, NY, USA). Cells were directly seeded in Epic 384-well biosensor microplates with a seeding density of 32,000 and 25,000 cells per well for HT-29 and A431 cells, respectively. 149 After cultured for \sim 20 h, these cells formed a monolayer in the cell culture medium with a confluency of ~95%. A431 cells were further starved for 24 h using serum-free medium. After culture, cells were manually washed using the assay buffer and 152 maintained with 30 μ L of the assay buffer for 1 h in the Epic system.

For profiling compounds in HT-29 cells, a 2-min baseline was first established, followed by adding compounds and monitoring the compound-induced DMR signals for 1 h. Then a 2-min baseline was re-established, followed by adding 1 µM zaprinast or 16 µM acetylcholine and monitoring DMR signals for another 1 h. Similarly, for profiling compounds in A431 cells, a 2-min baseline was first established, followed

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by compound addition and DMR recording for 1 h. A 2-min baseline was then re-established, followed by adding 16 nM bradykinin or 1 µM histamine and monitoring DMR signals for 1 h.

For DMR agonist assay, after establishment of a 2-min baseline, phenolic acids at varied doses were added individually and the DMR signals were recorded for 1 h. For DMR desensitization assay, cells were pretreated with these compounds for 1 h, followed by recording a 2-min baseline, adding 1 µM zaprinast and monitoring DMR signals for 1h. For DMR antagonist assay, HT-29 cells were pretreated with an antagonist, SPB05142 for 5 min or ML145 for 10 min, following by recording a 167 2-min baseline, adding a ligand at its indicated concentration $(EC_{20}, EC_{50}$ or $EC_{80})$ and monitoring DMR signals for 1 h.

169 To examine compounds for altering intracellular Ca^{2+} , a two-step assay was also used. Different doses of salvianolic acid A, salvianolic acid C, thapsigargin and cyclopiazonic acid were first assayed, followed by recording the DMR signal induced by cyclopiazonic acid at 16 µM for 1 h.

2.4. Western blot

After cells were treated with compounds for 15 min, whole cell lysates were harvested using lysis buffer (KeyGEN BioTECH, Naijing, China). For each group, 20 µg total protein was electrophoresed through a 12% (w/v) acrylamide gel and electroblotted onto polyvinylidene difluoride (PVDF) membranes. Antibodies were 179 diluted as follows: β-actin (1:300), ERK1/2 (1:300), p-ERK1/2 $\frac{\text{Thr202/Tyr204}}{1.1000}$ and GPR35 (1:500). Detection was performed by GTVTM Ⅲ Detection System/Mo&Rb (GeneTech, Shanghai, China).

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2.5. Data analysis

All DMR data were acquired using Epic Imager software (Corning, NY, USA) and processed by Imager Beta 3.7 (Corning), Microsoft Excel 2010 and GraphPad Prism 6.02 (GraphPad Sofware Inc., San Diego, CA, USA). All DMR signals were 187 background corrected. All EC_{50} or IC_{50} values described were calculated based on the maximal amplitude of DMR signal within 40 min post-stimulation. All DMR data 189 were from two independent measurements, each in duplicate $(n = 4)$.

3. Results and discussion

3.1. Profiling of phenolic acids from Danshen

We first profiled the activity of 10 phenolic acids in Danshen using RWG biosensor-enabled label-free cell phenotypic assay in HT-29 and A431 cells. The concentration of each compound for profiling was 10 µM. These phenolic acids are known to present in Danshen (**Fig. 1**). A two-step DMR assay was performed. The first step was DMR agonist assay to characterize all compounds for their agonist activities in HT-29 and A431 cells. The second step was DMR desensitization assay to examine the ability of each compound to desensitize the cellular response induced by a panel of agonist probe molecules, each at its saturating dose. Since HT-29 cells 202 endogenously express $GPR35^{24}$ and muscarinic M3 receptor,^{12, 25} zaprinast²⁶ and 203 acetylcholine¹² were chosen as their agonist probes, respectively. Similarly, A431 204 endogenously express bradykinin B_2 receptor^{25,26} and histamine H₁ receptor²⁷, bradykinin and histamine were chosen as agonist probes, respectively. Based on the 206 EC₁₀₀ reported in literature, the doses of probe agonists examined were set to be 1 μ M, 16 µM, 16 nM and 1 µM for zaprinast, acetylcholine, bradykinin and histamine,

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respectively. Additionally, the expression level of GPR35 in HT-29 and A431 cells was examined using western blot and DMR assay (**Fig. S1**). Results indicated that the expression level of GPR35 in HT-29 cells was much higher than that in A431 cells.

Results showed that in HT-29 cells salvianolic acid A and salvianolic acid C triggered a large DMR, while lithospermic acid, salvianolic acid B and danshensu led to a relatively small DMR, and all other five compounds resulted in negligible DMR (**Fig. 2a**). Interestingly, the five active compounds, including salvianolic acid A, salvianolic acid B, salvianolic acid C, lithospermic acid and danshensu, all attenuated the zaprinast-induced signal (**Fig. 2b**). Furthermore, the ability of salvianolic acid A and salvianolic acid C to attenuate the zaprinast-induced signal was greater than that of lithospermic acid, salvianolic acid B and danshensu. However, all compounds had no or little effect on the acetylcholine-induced signal (**Fig. 2c**). Combining the agonistic activity with the desensitization pattern suggests that lithospermic acid, salvianolic acid A, salvianolic acid B, salvianolic acid C and danshensu may have agonistic activity at the GPR35.

Different label-free pharmacological patterns were observed in A431 cells. The DMR responses at 30 min post-stimulation showed that salvianolic acid A and salvianolic acid C triggered relatively strong DMR signal, while lithospermic acid and salvianolic acid B led to small DMR signal, and other compounds were inactive (**Fig. 2d**). Furthermore, both salvianolic acid A and salvianolic acid C attenuated the bradykinin- and histamine-induced DMR signals (**Fig. 2e** and **2f**, respectively). Given 229 that in A431 cells there is little GPR35 expressed (Fig. S1), histamine activates the H_1 230 receptor triggering G_q -signaling,⁵ and bradykinin activates B_2 receptor triggering both 231 G_s- and G_q-pathways²⁷, these results suggest that salvianolic acid A and salvianolic 232 acid C may alter intracellular Ca^{2+} via another unknown target.

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3.2. Activity of phenolic acids at the GPR35

We next examined the potential agonistic activity of lithospermic acid, salvianolic acid A, salvianolic acid B, salvianolic acid C and danshensu at the GPR35 using three different DMR assays. DMR agonist assay was employed to determine their agonist activities. DMR desensitization assay and DMR antagonist assay were used to examine the specificity of these five compounds to activate GPR35.

First, DMR agonist assay showed that lithospermic acid, salvianolic acid A, salvianolic acid B, salvianolic acid C and danshensu with the concentration range of $0.03 \sim 250 \mu M$ all triggered dose-dependent positive DMR signals, but with different amplitudes (**Fig. 3**). According to DMR characteristics, these five compounds can be divided into two groups. The first group consists of salvianolic acid A and salvianolic acid C, both triggering a quite large DMR with almost identical characteristics (**Fig. 3a**). The second group consists of lithospermic acid, salvianolic acid B and danshensu, all triggering a moderate DMR (**Fig. 3b**). Except for salvianolic acid B and danshensu, whose DMR signals seem not reach saturation at the maximal concentration of 250 µM examined, the dose responses of salvianolic acid A, salvianolic acid C and lithospermic acid were best fitted with a single-phase non-linear regression (**Fig. 3c** 251 and **3d**). The EC₅₀ values of these compounds are showed in **Table 1**. Furthermore, the maximal DMR of lithospermic acid, salvianolic acid B and danshensu all shared similarity to those of zaprinast and kynurenic acid (Fig. 4). Zaprinast²⁶ and kynurenic acid²⁸ are two known GPR35 agonists and often used as probe agonists for GPR35. Of note, salvianolic acid A or salvianolic acid C at 16 µM gave rise to a DMR also similar to that of zaprinast, but at high doses triggered a DMR that is much greater

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257 than the maximal DMR of zaprinast, suggesting that both compounds may also 258 activate another unknown target or pathway, besides GPR35.

259 Second, DMR desensitization assay was performed against the DMR of 1 μ M zaprinast. Results showed that these five compounds all dose-dependently 261 desensitized the DMR of zaprinast, leading to an apparent IC₅₀ of 3.48 \pm 0.38 μ M, $72.74 \pm 22.96 \mu M$, $2.70 \pm 0.31 \mu M$, $109 \pm 23 \mu M$ and $30.42 \pm 2.25 \mu M$ (n = 4 for all) for salvianolic acid A, salvianolic acid B, salvianolic acid C, lithospermic acid and danshensu, respectively (**Fig. 5**; **Table 1**). These results suggest that these compounds activate GPR35.

266 Third, DMR antagonist assay was performed to confirm the specificity of these 267 compounds to activate GPR35. Known GPR35 antagonists $SPB05142^{29}$, 30 and 268 ML145 $^{31, 32}$ were used to pretreat HT-29 cells at different doses, followed by detecting 269 DMR signal induced by each compound at its respective dose close to EC_{80} , EC_{50} or 270 EC₂₀. Results showed that SPB05142 displayed different inhibitory effects on the 271 DMR of different compounds at different doses. When the dose of each compound 272 was assayed at its EC_{80} , SPB05142 dose-dependently and completely blocked the 273 DMR of 125 µM lithospermic acid or 100 µM danshensu, leading to an IC₅₀ of 2.29 \pm 274 0.52 μ M and 2.98 \pm 1.35 μ M (n = 4), respectively (**Fig. 6a**). However, SPB05142 275 only partially blocked the DMR of 100 μ M salvianolic acid B with an IC₅₀ of 0.71 \pm 276 0.42 μ M (n = 4) (**Fig. 6a**), but had little effect on the DMR of 125 μ M salvianolic acid 277 A, or 100 µM salvianolic acid C (**Fig. 6b**). On the other hand, SPB05142 weakly 278 attenuated the DMR of salvianolic acid A and salvianolic acid C at their EC_{50} (**Fig.** 279 **6c**), but dose-dependently inhibited their DMR at their respective EC_{20} (15 μ M and 4 280 µM, respectively) with an IC₅₀ of 2.52 \pm 0.48 µM and 1.57 \pm 0.64 µM (n = 4),

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respectively (**Fig. 6d**). Almost identical inhibition pattern was observed using ML145 as an antagonist (**Fig. S2, Table 1**).

We further examined the activity of salvianolic acid A, salvianolic acid C, salvianolic acid B, lithospermic acid and danshensu using ERK phosphorylation assay. The activation and signaling of many GPCRs induce ERK phosphorylation, including 286 GPR35.³³ As in Fig. 7, it shows that salvianolic acid A, salvianolic acid C, salvianolic acid B, lithospermic acid and danshensu lead to ERK phosphorylation (**Fig. 7a)**. As controls, the known GPR35 agonist zaprinast also triggers ERK phosphorylation (**Fig. 7b**). Moreover, the GPR35 antagonist ML145 attenuated ERK phosphorylation induced by these compounds (**Fig. 7c**). These results suggest that salvianolic acid A, salvianolic acid C, salvianolic acid B, lithospermic acid and danshensu resulted in the phosphorylation of ERK via the activation of GPR35.

Together, these results suggest that all five compounds display agonistic activity at the GPR35. Lithospermic acid and danshensu are mostly specific to activate GPR35 in HT-29 cells. However, salvianolic acid B may also activate another unknown target, and both salvianolic acid A and salvianolic acid C at high doses may activate multiple targets/pathways, triggering a quite large DMR.

3.3. Deconvolution of possible target(s), besides GPR35, of salvianolic acid A and salvianolic acid C

Both salvianolic acid A and salvianolic acid C appear to display polypharmacology. This was evidenced by that both triggered a larger DMR consisting of contributions from GPR35 as well as other unknown pathway(s) in HT-29 cells. Furthermore, both compounds also triggered a large DMR in A431 cells and attenuated the bradykinin-and histamine-induced DMR signals, probably due to heterologous desensitization

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306 mechanism. The activation of G_q -coupled receptors is known to sequentially activate 307 trimeric G proteins and phospholipase C (PLC), the latter of which then hydrolyzes 308 the membrane lipid phosphatidylinositol bisphosphate, yielding diacylglycerol (DAG) 309 and inositol triphosphate (IP3). IP3 binds to and opens a calcium channel in the 310 endoplasmic reticulum, causing calcium mobilization. On the other hand, Danshen is 311 known to cause vasorelaxant actions due to the inhibition of Ca^{2+} flux.^{34, 35} Therefore, 312 we hypothesized that salvianolic acid A and salvianolic acid C may alter intracellular Ca^{2+} via an unknown target. To validate this hypothesis, we used thapsigargin and 314 cyclopiazonic acid as the probe molecules. Thapsigargin is a non-competitive 315 inhibitor of sarco/endoplasmic reticulum Ca^{2+} -ATPase (SERCA),³⁶ while 316 cyclopiazonic acid is a specific inhibitor of SERCA.³⁷ Both inhibitors can raise 317 intracellular calcium concentration by blocking the ability of the cell to pump calcium 318 into the sarcoplasmic and endoplasmic reticula.³⁵

DMR agonist assay showed that salvianolic acid A and C triggered robust DMR with similar characteristics in A431 cells (**Fig. 8a** and **b**), and salvianolic acid C exhibited slightly higher potency than salvianolic acid A (**Fig. 8c**). On the other hand, thapsigargin and cyclopiazonic acid triggered comparable DMR in A431, whose characteristics were different from those of salvianolic acid A and C (comparing **Fig. 8d** and **e** with Fig. 8a and **b**). As expected, 35 thapsigargin exhibited higher potency 325 than cyclopiazonic acid (**Fig. 8f**). The IC_{50} values to trigger DMR were found to be 326 122.46 \pm 11.75 μ M, 60.75 \pm 3.01 μ M, 0.30 \pm 0.02 μ M and 2.08 \pm 0.09 μ M (n = 4) for salvianolic acid A, salvianolic acid C, thapsigargin and cyclopiazonic acid, respectively.

329 DMR desensitization assay showed that all four compounds caused dose-dependent 330 desensitization of cells responding to the second stimulation with 16 μ M

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331 cyclopiazolic acid (**Fig. 9**). The IC₅₀ values to desensitize the cyclopiazolic acid DMR 332 were found to be 71.43 \pm 9.53 μ M, 26.06 \pm 1.70 μ M, 0.023 \pm 0.002 μ M and 5.12 \pm 333 0.50 μ M (n = 4 for all) for salvianolic acid A, salvianolic acid C, thapsigargin and cyclopiazonic acid, respectively. These results suggest that salvianolic acid A and salvianolic acid C alter intracellular Ca^{2+} via an unknown target and share similar mechanism to those of thapsigargin and cyclopiazonic acid. Further elucidation of the exact target intervened by salvianolic acid A and salvianolic acid C is warranted and currently under investigation.

4. Conclusion

Danshen water extract has clinical benefits for treating cardiovascular and cerebrovascular diseases and contains several phenolic acids. Here we used label-free cell phenotypic assay to investigate the possible targets of these phenolic acids from Danshen. We found that lithospermic acid, salvianolic acid A, salvianolic acid B, salvianolic acid C and danshensu all displayed agonistic activity at the GPR35, but with different potency. We also found that salvianolic acid A and salvianolic acid C 346 also alter intracellular Ca^{2+} via a mechanism that is similar to two known SERCA inhibitors. The multi-target activity of these phenolic acids may be related to the clinical features of Danshen products. Specifically, GPR35 has been implicated in inflammation, hypertension, coronary artery disease and cancer. Thus, the agonistic activity of multiple phenolic acids in Danshen may be useful for elucidating the mechanism of Danshen products in the treatment of cardiovascular diseases.

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References

- 1. D. J. Newman and G. M. Cragg, *J. Nat. Prod.*, 2012, **75**, 311-335.
- 2. M. S. Butler, *Nat. Prod. Rep.*, 2008, **25**, 475-516.
- 3. D. C. Swinney and J. Anthony, *Nat. Rev. Drug Discov.*, 2011, **10**, 507-519.
- 4. J. Lee and M. Bogyo, *Curr. Opin. Chem. Biol.*, 2013, **17**, 118-126.
- 5. E. Tran and Y. Fang, *J. Biomol. Screen*, 2008, **13**, 975-985.
- 6. Y. Fang, *Expert. Opin. Drug Discov.*, 2011, **6**, 1285-1298.
- 7. Y. Fang, *J. Pharmacol. Toxicol. Methods*, 2013, **67**, 69-81.
- 8. Y. Fang, A. M. Ferrie, N. H. Fontaine, J. Mauro and J. Balakrishnan, *Biophys. J.*, 2006, **91**, 1925-1940.
- 9. Y. Fang, *Sensors*, 2007, **7**, 2316-2329.
- 10. A. M. Ferrie, H. Y. Sun and Y. Fang, *Sci Rep*, 2011, **1**.
- 11. H. Y. Deng, H. B. Hu and Y. Fang, *Sci Rep*, 2012, **2**.
- 12. H. Y. Deng, C. M. Wang, M. Su and Y. Fang, *Anal. Chem.*, 2012, **84**, 8232-8239.
- 13. L. M. Zhou, Z. Zuo and M. S. S. Chow, *J. Clin. Pharmacol.*, 2005, **45**, 1345-1359.
- 14. Y. G. Li, L. Song, M. Liu, H. Zhi Bi and Z. T. Wang, *J. Chromatogr. A*, 2009, **1216**, 1941-1953.
- 15. X. P. Chen, J. J. Guo, J. L. Bao, J. J. Lu and Y. T. Wang, *Medicinal Research Reviews*, 2014, **34**, 768-794.
- 16. M.-H. Li, J.-M. Chen, Y. Peng, Q. Wu and P.-G. Xiao, *J. Ethnopharmacol.*, 2008, **120**, 419-426.
- 17. M. Liu, Y. G. Li, F. Zhang, L. Yang, G. X. Chou, Z. T. Wang and Z. B. Hu, *J. Sep. Sci.*, 2007, **30**, 2256-2267.
- 18. R.-W. Jiang, K.-M. Lau, P.-M. Hon, T. C. Mak, K.-S. Woo and K.-P. Fung, *Curr. Med. Chem.*, 2005, **12**, 237-246.
- 19. J. H. C. Ho and C. Y. Hong, *J. Biomed. Sci.*, 2011, **18**.
- 20. J. Huang, Y. Qin, B. Liu, G. Y. Li, L. Ouyang and J. H. Wang, *Cell Prolif.*, 2013, **46**, 595-605.
- 21. D. G. Kang, H. Oh, H. T. Chung and H. S. Lee, *Phytother. Res.*, 2003, **17**, 917-920.
- 22. B. Jiang, J. Chen, L. Xu, Z. Gao, Y. Deng, Y. Wang, F. Xu, X. Shen and D.-a. Guo, *BMC Pharmacology*, 2010, **10**, 10.
- 23. B. Sperl, M. H. J. Seifert and T. Berg, *Bioorg. Med. Chem. Lett.*, 2009, **19**, 3305-3309.
- 24. H. Y. Deng, H. B. Hu and Y. Fang, *FEBS Lett.*, 2011, **585**, 1957-1962.
- 25. H. Y. Deng, H. Y. Sun and Y. Fang, *J. Pharmacol. Toxicol. Methods*, 2013, **68**, 323-333.
- 26. Y. Taniguchi, H. Tonai-Kachi and K. Shinjo, *FEBS Lett.*, 2006, **580**, 5003-5008.
- 27. C. Liebmann, A. Graness, B. Ludwig, A. Adomeit, A. Boehmer, F. D. Boehmer, B. Nurnberg
- and R. Wetzker, *Biochem. J*, 1996, **313**, 109-118.

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415 **Table 1** The active phenolic acids and their pharmacological characteristics in HT-29

416 cells.

417 $\overline{C_{50}}$ to desensitize the DMR of 1 µM zaprinast in HT-29 cells.

418 $*$ IC₅₀ of SPB05142 to block the DMR of a phenolic acid compound at a dose of 125 μ M, 100 μ M,

419 100 µM, 15 µM and 4 µM for lithospermic acid, salvianolic acid B, danshensu, salvianolic acid A and

420 salvianolic acid C, respectively. The HT-29 cells were pre-treated with the antagonist for 5 min.

^{***} IC₅₀ of ML145 to block the DMR of compounds at the same doses as above. The HT-29 cells were

422 pre-treated with the antagonist for 10 min.

423

- **Fig. 1.** Chemical structures of 10 known phenolic acids in Danshen.
-

428	Fig. 2 A label-free cell phenotypic profiling technique to identify potential
429	polypharmacology of phenolic acid compounds in HT-29 cells (a-c) and A431 cells
430	(d-f). (a) The DMR amplitudes at 30 min post-stimulation of 10 μ M compounds as a
431	function of compounds in HT-29 cells. (b) The DMR response of 1μ M zaprinast after
432	the cells were pre-stimulated with each compound for 1 h. (c) The DMR response of
433	16 µM acetylcholine after the cells were pre-stimulated with each compound for 1 h.
434	(d) The DMR amplitudes of 10 μ M compounds as a function of compounds in A431
435	cells. (e) The DMR response of 16nM bradykinin after the cells were pre-stimulated
436	with each compound for 1 h. (f) The DMR response of 1 μ M histamine after the cells
437	were pre-stimulated with each compound for 1 h. The compounds are lithospermic
438	acid (1), salvianolic acid A (2), salvianolic acid B (3), salvianolic acid C (4),
439	salvianolic acid D (5), danshensu (6), caffeic acid (7), isoferulic acid (8), carnosol (9),
440	protocatechuic aldehyde (10) and control (11). All data represents mean \pm s.d. from 2
441	independent measurements, each in duplicate $(n = 4)$.
442	

Fig. 3 DMR dose responses of compounds in HT-29 cells. (a and b) Real time DMR signals of salvianolic acid A (a) and danshensu (b); (c and d) DMR amplitudes as a function of their doses: salvianolic acid A and salvianolic acid C (c), salvianolic acid

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452 $(n = 4)$.

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446 B, lithospermic acid and danshensu (d). All data represents mean \pm s.d. from 2 447 independent measurements, each in duplicate $(n = 4)$. 449 **Fig. 4** DMR dose responses of probes in HT-29 cells. (a, b) Real time DMR signals of 450 zaprinast (a) and kynurenic acid (b); (c) DMR amplitudes as a function of their doses. 451 All data represents mean \pm s.d. from 2 independent measurements, each in duplicate 454 **Fig. 5** The DMR amplitudes of 1 µM zaprinast as a function of compounds doses in 455 HT-29 cells after pre-stimulation with these compounds for 1 h. (a) Salvianolic acid A 456 and salvianolic acid C; (b) salvianolic acid B, lithospermic acid and danshensu. All 457 data represents mean \pm s.d. from 2 independent measurements, each in duplicate (n =

460 **Fig. 6** The DMR amplitudes of compounds as a function of SPB05142 doses in HT-29 461 cells after pre-treated with the antagonist for 5 min. (a) Lithospermic acid, salvianolic 462 acid B and danshensu at their EC_{80} of 125 μ M, 100 μ M and 100 μ M, respectively; (b) 463 salvianolic acid A and salvianolic acid C at their EC_{80} of 125 μ M and 100 μ M, 464 respectively; (c) salvianolic acid A and salvianolic acid C at their EC_{50} of 35 μ M and 465 20 μ M, respectively; (d) salvianolic acid A and salvianolic acid C at their EC₂₀ of 15 466 μ M and 4 μ M, respectively. All data represents mean \pm s.d. from 2 independent 467 measurements, each in duplicate $(n = 4)$.

480 from 2 independent measurements, each in duplicate $(n = 4)$.

Fig. 9 The DMR amplitudes of 16 µM cyclopiazonic acid as a function of compounds doses in A431 cells after pre-stimulation with these compounds for 1 h. (a) Salvianolic acid A and salvianolic acid C; (b) thapsigargin and cyclopiazonic acid. All 485 data represents mean \pm s.d. from 2 independent measurements, each in duplicate (n = 4).

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Fig. 3

