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What has traditional Chinese medicine delivered for modern medicine?

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The field of Traditional Chinese Medicine (TCM) represents a vast and largely untapped resource for modern medicine. Exemplified by the success of the antimalarial artemisinin, the recent years have seen a rapid increase in the understanding and application of TCM-derived herbs and formulations for evidence-based therapy. In this review, we summarise and discuss the developmental history, clinical background and molecular basis of an action for several representative TCM-derived medicines, including artemisinin, arsenic trioxide, berberine and *Salvia miltiorrhiza* or Danshen. Through this, we highlight important examples of how TCM-derived medicines have already contributed to modern medicine, and discuss potential avenues for further research.

Introduction

Traditional Chinese Medicine (TCM) refers to the holistic approach to diagnosis, pathophysiology and therapy in the Chinese materia medica, based on over 2000 years of accumulated knowledge and practice (Ref. 1). Major aspects of the practice include herbal medication, acupuncture and other physical therapy such as massage (Refs 2, 3, 4). Outside of China, the practice is generally regarded as a complementary or 'alternative' form of medicine, although the international prevalence of the practice has been steadily increasing (Ref. 5). Acceptance of TCM by the scientific community has been limited at best. Certain guiding principles of TCM such as the concept of 'vital energy' are esoteric and difficult to validate under modern scientific methods, and the TCM approach to both diagnosis and treatment also fundamentally differs from conventional Western methods (Refs 6, 7). Nevertheless, there are select aspects of the practice that hold clear promise for modern evidence-based medicine. The field of Chinese herbals, in particular, has drawn increasing interest as a source of novel drugs and drug leads, which is unsurprising considering that many modern drugs are in fact derivatives of herbal medicines and natural products (Refs 8, 9, 10, 11). The awarding of part of the 2015 Nobel Prize in Physiology or Medicine for Tu Youyou's discovery and development of artemisinin, a potent antimalarial derived from the herbal Artemisia annua, is a clear example and reminder of the potential held by herbal medicine (Ref. 12). Looking ahead, further understanding and appreciation of the current status and successes of TCM herbals will be important for continued developments in the field. In this paper, we seek to provide an overview of some of the major contributions that Chinese herbals have made to modern medicine, focusing on the background and process of discovery, molecular mechanisms, clinical evidence for established treatments as well as novel and promising treatments under development.

Artemisinin in the treatment of malaria

Malaria is caused by the *Plasmodium* genus of endoparasites and remains a global health concern with 212 million new cases and 429 000 deaths as recently as 2015 (Ref. 13). The artemisinin family of sesquiterpene lactone compounds, including artemisinin itself as well as synthetic derivatives such as dihydroartemisinin, artesunate and artemether (Fig. 1), currently serve as the standard treatment and represent the front line of antimalarial drugs with remarkable potency, specificity and safety (Refs 14, 15, 16). ACT (artemisinin-based combination therapies) remain the most effective and recommended treatment regimens for uncomplicated malaria, especially in cases caused by the prevalent *Plasmodium falciparum* strain of parasites (Refs 17, 18). The compound is derived from *A. annua* or 'sweet wormwood', which has been used in TCM (where it is known as 'Qinghao') for the relief of periodic fevers, a symptom of malaria (Ref. 19). The earliest documentation of such usage dates back to the East Jin Dynasty between 317 and 420 A.D., in Ge Hong's *A Handbook of Prescriptions for Emergencies* (Refs 12, 20).

The story of artemisinin is likely the most well-known example of Chinese medicine's contributions to modern medicine, given its outstanding real-world results and the recognition of the 2015 Nobel Prize. Under the Chinese nationwide antimalarial research initiative 'Project 523', the antimalarial agency headed by Tu Youyou in her institute performed extensive

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Figure 1. The structures of artemisinin and its clinically used derivatives.

Artemisinin

Dihvdroartemisinin

Artesunate Artemether

screening without success in hundreds of Chinese medicinederived compounds, including A. annua (Refs 12, 20). With reference to the ancient literature which described preparation conditions for the herb, Tu then attempted a low-temperature approach in extracting the active components of A. annua and in 1971 identified a particular non-toxic extract which displayed close to 100% efficacy in mouse malaria models. The extract was quickly brought to clinical trial where recovery and full parasite clearance was reported in over 90% of cases. The active compound of A. annua was then successfully purified by the same group in 1972 and named artemisinin, or Qinghaosu in Chinese Mandarin. Finally, the full stereochemical structure of artemisinin was elucidated and published in 1977 (Ref. 21). Further work by Tu and others in the modification and derivitisation of artemisinin set the foundation for further understanding of critical functional groups as well as the development of artemisinin derivatives such as dihydroartemisinin, artesunate and artemether, which now serve as indispensable antimalarials worldwide (Refs 22, 23).

Since artemisinin began to draw global attention in the early 1980s, numerous clinical studies have been performed in endemic regions to assess its safety and potency either in combination with other antimalarials or as a monotherapy. In particular, studies from China, Africa and Southeast Asia over the past three decades including recent meta-analytical data have convincingly demonstrated the outstanding clinical properties of artemisinin. Clinical results were marked by rapid parasite clearance and improvement of symptoms, especially for uncomplicated P. falciparum malaria in combination with longer-acting antimalarials such as mefloquine, lumefantrine, benflumetol and piperaquine (Refs 17, 18, 24, 25, 26, 27, 28, 29). Results in severe malaria have likewise been generally positive for artemisinin derivatives compared with other treatments such as quinine, albeit with reduced efficacy compared with uncomplicated cases (Refs 30, 31, 32). Reports of major adverse effects have been minimal, especially considering the volume of available data and the ubiquity of the drug (Refs 15, 33). In addition to its place as the most important antimalarial, limited clinical success has also been reported for the role of artemisinin in other diseases such as colorectal cancer, and research is ongoing for the re-purposing of artemisinin for various nonmalarial roles (Refs 34, 35, 36, 37, 38).

Despite the widespread usage of artemisinin, the current molecular understanding of its mechanism of action remains incomplete. The artemisinin compounds are sesquiterpene lactones sharing the 1,2,4-trioxane pharmacophore, which contains an endoperoxide bridge that is thought to be essential for its pharmacological activity against both malaria as well as cancer (Ref. 39). Artemisinin itself is a prodrug that must be activated via cleavage of this endoperoxide bridge for drug activity, although the mechanism of this activation remains an issue of some debate (Ref. 40). Both free ferrous iron (Fe²⁺) as well as haem released from haemoglobin digestion have been proposed to activate artemisinin (Refs 41, 42, 43). Reductive ion transfer

(from iron or haem) is proposed to induce homolytic cleavage of the endoperoxide bridge, producing oxygen-centred radicals that subsequently isomerise to form reactive carbon-centred radicals (Refs 44, 45, 46). In an alternative theory, free Fe²⁺ functions as a Lewis acid in catalysing the heterolytic cleavage of the endoperoxide, generating cationic intermediates and hydroperoxides that can subsequently generate reactive hydroxyl radicals (Ref. 47). While free ferrous iron has been understood to be responsible for artemisinin activation, recent evidence from mass spectrometry and proteomics approaches have demonstrated that redox active haem could also play an important and possibly principal role in the process (Refs 42, 43, 48, 49). It will be important to clearly elucidate the activation mechanism of artemisinin as this could directly relate to its remarkable specificity.

Following activation, the downstream mechanisms by which artemisinin achieves its antimalarial properties are likewise incompletely understood (Ref. 40). Malaria parasites break down host cell haemoglobins, releasing large amounts of haem in the process (Refs 50, 51). Haem build-up then leads to haematin formation which can induce oxidative and lytic damage to the parasite (Ref. 52). As a defence mechanism, the parasites are able to crystallise toxic haematin to the nontoxic haemozoin (Ref. 53). Activated artemisinin has been shown to be able to bind and alkylate haem both in vitro and in vivo, possibly preventing haemozoin formation and causing a toxic accumulation of haem (Refs 54, 55, 56, 57). Likewise, activated artemisinin is also known to be able to alkylate protein targets, which could interfere with critical biological functions and contribute to toxicity (Refs 58, 59). The identification of such protein targets of artemisinin has been a topic of great interest, and well-described examples include TCTP (translationally controlled tumour protein) and PfATP6 (Refs 60, 61). In particular, PfATP6 (a sarco/ endoplasmic reticulum Ca2+-ATPase or SERCA) was reported to be bound and inhibited by artemisinin, making it a putative target of interest (Ref. 62). Recent reports using mass spectrometry-based proteomics approaches have highlighted a potentially promiscuous and indiscriminate targeting mechanism of artemisinin, where cellular targets including proteins are nonselectively alkylated in a proximity-dependent manner (Refs 42, 43, 63). Our 2015 study identified over 100 binding targets of artemisinin using chemical probes (Fig. 2), implicating multiple cellular functions, including biosynthetic and metabolic pathways (Ref. 42). Under such a model, a large number of proteins may be affected simultaneously and many critical biological pathways could be disrupted. The specificity of artemisinin, in this case, could then be largely attributed to the conditions and the extent of drug activation, rather than the specific targeting of protein effectors (Fig. 3). A combination of haem- and protein alkylation by artemisinin (which is specifically activated in high-haem, high-iron parasite conditions) could possibly explain both the specificity and potency of the drug. Nevertheless, it should be considered that certain proteins or other cellular targets could play a

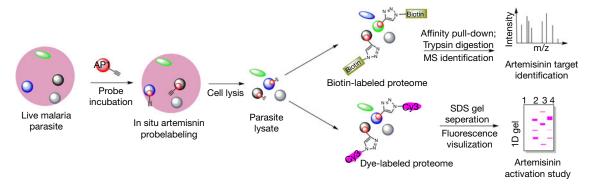


Figure 2. General workflow of a chemical biology approach to study the targets and activation of artemisinin using artemisinin-based chemical probes.

disproportionately large role in the downstream effects of artemisinin. The elucidation of such targets remains a challenge for further understanding and development of this important drug. Ultimately, the story of artemisinin's discovery and development is a classic example of the value held in traditional medicine and traditional literature. Considering the long history of TCM as well as other systems of traditional medicine, it would surely not be surprising that other hidden gems already described and used by our predecessors yet remain to be discovered.

Arsenic trioxide in the treatment of acute promyelocytic leukaemia

Arsenic compounds, though well known for their toxicity, have been used for therapeutic purposes for possibly thousands of years. In accordance with the TCM principle of 'attacking poison with poison', the toxic compound arsenic trioxide (ATO, As₂O₃) or 'Pi Shuang' has been used both topically and orally in TCM as

a medication for fevers, skin conditions and many other diseases (Ref. 64). In TCM as well as pre-modern Western medicine, ATO has also been used in the treatment of cancer, including leukaemia, and ATO was, in fact, a common treatment for chronic myeloid leukaemia in the early 20th century (Ref. 65). With the advent of more sophisticated cancer therapeutics including chemotherapy and the concerns of toxicity, however, the therapeutic use of ATO outside of China was eventually largely phased out (Ref. 66).

In 1996 and 1997, clinical and in vitro data from China reported by Shen et al. and Chen et al. were published in *Blood* detailing the application of ATO to the treatment of acute promyelocytic leukaemia (APL), drawing international attention to the compound which has already been in development in China for over a decade (Refs 67, 68, 69). The efforts could be traced to the work of Dr Zhang Tingdong and colleagues at Harbin Medical University during the 1970s, who studied the first TCM-derived formulation of ATO in the treatment of

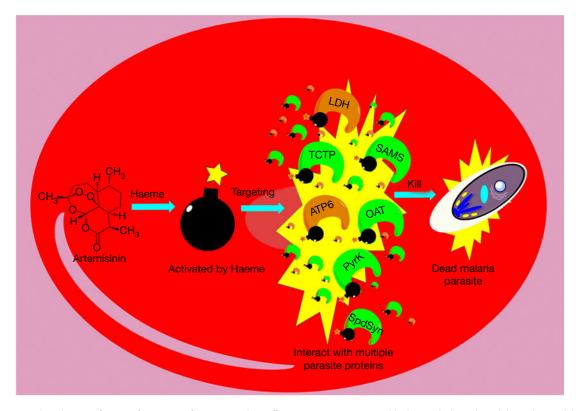


Figure 3. The proposed mechanism of action of artemisinin for its anti-malaria effects. Artemisinin is activated by haem which is released during haemoglobin digestion by the malaria parasite. This generates reactive radicals which alkylate a range of parasite proteins, eventually killing the parasite. Ca²⁺-ATPase (ATP6), the translationally controlled tumour protein (TCTP), ornithine aminotransferase (OAT), pyruvate kinase (PyrK), L-lactate dehydrogenase (LDH), spermidine synthase (SpdSyn) and S-adenosylmethionine synthetase (SAMS).

leukaemia and identified ATO as the principal active component (Ref. 70). APL itself was an acute and aggressive form of leukaemia that saw considerable improvement in treatment outcomes following the development of all-trans retinoic acid (ATRA) therapy, which was also developed and reported in the late 1980s by Chinese researchers (Refs 71, 72). However, ATRA treatment alone suffered from a high rate of relapse, necessitating novel and complementary therapeutic approaches (Ref. 73). The development of ATO for APL treatment by Zhang and others continued in China with promising clinical results, and the 1997 publications were followed up by Soignet et al. in their 1998 clinical report (Ref. 74). Much like artemisinin, the outstanding clinical properties of ATO led to rapid adoption and development by the international community. ATO in combination with ATRA therapy is currently a standard APL treatment, and the clinical success of the treatment in both newly diagnosed and relapsed cases has been well demonstrated and comprehensively reviewed in several authoritative works (Refs 75, 76, 77, 78, 79, 80, 81).

The core mechanisms of ATO in APL treatment are currently understood to be a concentration-dependent dual effect by the induction of cancer cell differentiation and apoptosis. In initial in vitro testing on APL NB4 cell lines, Chen et al. reported robust apoptosis induction by ATO at higher concentrations between $0.5-2~\mu M$ and partial differentiation at lower (0.1-0.25 μM) over longer time periods up to 10 days (Ref. 67). ATO-induced apoptosis is mediated by the canonical mitochondrial pathway, inducing mitochondrial membrane potential collapse by targeting mitochondrial membrane proteins and demonstrating characteristic cytochrome c release, caspase 8 activation and cleavage of poly (ADP-ribose) polymerase (PARP) (Refs 82, 83, 84, 85, 86). Notably, ATO-induced apoptosis is associated with downregulated Bcl-2 expression and can be inhibited by Bcl-2 overexpression (Refs 67, 83). Cellular redox conditions have also been shown to be important for ATO-induced apoptosis, possibly by interfering with the ability of ATO to bind protein effectors through interactions with exposed sulfhydryl (-SH) groups of cysteine residues (Refs 66, 87). Accordingly, modulation of the antioxidant glutathione (GSH) system has been reported to affect cancer cell sensitivity to ATO (Ref. 88). Other reported mechanisms of ATO-induced apoptosis include ROS production and oxidative damage through the modulation of ROS-related genes, inhibition of NF-κB through the binding of IκB kinase (IKK), and further effects on many other signalling pathways, including MAPK, JAK-STAT and JNK which have been extensively reviewed elsewhere (Refs 81, 87, 89, 90).

The mechanisms by which ATO induces differentiation are less well understood but are likely related to its interactions with the PML-RAR α fusion protein which is characteristic of the vast majority of APL cases. This oncoprotein is caused by a particular t(15;17) chromosomal translocation which results in a fusion of the retinoic acid receptor alpha (RAR α) gene and promyelocytic leukaemia (PML) gene (Refs 91, 92). Endogenous PML is a tumour suppressor that localises to the nucleus and forms distinct macromolecular structures known as PML nuclear bodies (PML-NB) which are involved in transcriptional regulation (Refs 93, 94). Functional PML-NB prevents the development of malignancy and is required for terminal differentiation. In the case of APL, expression of the PML-RAR α fusion protein disrupts the normal PML-NB structure, resulting in errant protein-protein interactions, failure of differentiation and eventual leukemogenesis (Ref. 72). Through the induction of ROS formation and direct interaction with PML-RARα, ATO is reported to restore endogenous PML-NB structure and function through the induction of PML-RARα multimerisation, SUMOylation and eventual proteasomal degradation (Refs 95, 96, 97, 98). This ability of ATO to target a key oncogenic protein (as a so-called oncogene-directed therapy) underlies its remarkable specificity and efficacy. In this regard, the mechanism appears to be related to ATRA therapy (and hence the RAR pathway) which also induces differentiation and PML–RAR α degradation, although ATO and ATRA are known to interact with different regions of the fusion protein (Refs 95, 99). This distinction could partly explain the synergism between ATO and ATRA in combination, as well as the effectiveness of ATO in ATRA-resistant and relapsed cases (Ref. 87). In addition to the restoration of PML-NB function and extensive effects on gene regulation and cancer cell stemness, other factors such as miRNA have also recently been reported to contribute to ATO-induced differentiation (Refs 100, 101).

Beyond ATO, the general study of TCM-derived natural products and compound formulations for cancer therapy is an area of immense interest both within and outside China. Apart from the ongoing discovery and characterisation of active antineoplastics, the use of TCM as an effective adjuvant therapy in combination with conventional treatments as well as the role of TCM in pain management and palliative care should also not be overlooked (Refs 102, 103, 104, 105, 106, 107). For instance, the fourherb formulation PHY906 currently undergoing multiple clinical trials has been identified as an effective modulator of chemotherapy toxicity for multiple chemotherapy treatments (Refs 108, 109, 110). Given the recent resurgence of interest in natural products and the success of artemisinin and ATO in their respective roles, it is not difficult to imagine that more drugs and treatments of great potential remain within the TCM materia medica (Ref. 9). It will be of great interest moving forward to observe the potential contributions of TCM to evidence-based cancer therapy.

Berberine in the treatment of type 2 diabetes mellitus

The use of TCM for the prevention and treatment of diabetes in China dates back over 2000 years where diabetic symptoms were known as 'Xiaoke', or increased thirst (Ref. 111). Many herbs and formulations have been developed and remain widely in use in China for TCM treatment of type 2 diabetes mellitus in the present day (Ref. 112). Among those treatments, the isoquinone alkaloid known as berberine (Fig. 4) has stood out for its clinically demonstrated hypoglycemic and hypolipidemic properties. Berberine is an active component of Coptis chinensis rhizomes, a Chinese herbal used for the relief of diabetes as well as gastrointestinal disorders (Ref. 113). Traditionally known for a wide spectrum of antimicrobial activities, the application of berberine to diabetic symptoms was first reported and followed up in a Chinese study in the late 1980s, when berberine used in an antidiarrhoeal capacity for diabetic patients was shown to lower blood sugar (Refs 114, 115).

Compared with well-established treatments such as artemisinin or ATO, clinical study of berberine as an antidiabetic is at a relatively early stage with pilot trials based in China first being reported in 2008. In comparative studies between berberine and the first-line treatment metformin in 36 newly-diagnosed type 2 diabetics, Yin et al. reported comparable effects on the regulation

Figure 4. The structure of berberine.

of glucose metabolism with significant (P < 0.01) decreases in indicators such as haemoglobin A1c, fasting blood glucose and postprandial blood glucose. Berberine outperformed metformin in the regulation of lipid metabolism, effecting a decrease in plasma triglycerides and cholesterol over 13 weeks at a significant improvement over metformin (P < 0.05). In parallel, a study was conducted on the use of berberine in combination therapy with existing antidiabetic treatment for 48 patients with poorly controlled diabetes. The use of berberine significantly improved both glycemic and lipidemic parameters compared with baseline, and treatments were generally well tolerated with limited and transient adverse gastrointestinal reactions in the combination group but no functional liver or kidney damages (Ref. 116). In a double-blind randomised controlled trial in 110 patients published in the same year, Zhang et al. likewise reported the efficacy and safety of berberine in the treatment of newly diagnosed type 2 diabetics. Both glycaemic and lipidaemic indicators exhibited significant improvements compared to the placebo, and secondary outcomes including reduced body weight and blood pressure were also positive (Ref. 117).

As reviewed in meta-analyses published 2012 and 2015, subsequent randomised clinical trials (14 and 27 studies reviewed respectively) continued to provide positive results, with significant support for the efficacy of berberine in improving glycaemic control that is at least comparable or not inferior to current first-line treatments (Refs 118, 119). Considering that berberine is chemically distinct from other first-line options, the compound stands out as a potential novel antidiabetic both as monotherapy or in combination with other treatments for cases which respond poorly to existing treatments. Other plus points for berberine include a low cost of production, safety and a multi-faceted mechanism that appears to target multiple aspects of diabetes (Refs 119, 120). Berberine has been reported to target glucose metabolism through both insulin-dependent and independent pathways, increasing insulin sensitivity (partly through increased insulin receptor expression), insulin secretion, glucose uptake, and stimulating activation of the AMP-activated protein kinase (AMPK) pathway (Refs 121, 122, 123, 124). Other mechanisms including modulating liver metabolic function through regulation of gene expression, reduction of intestinal glucose uptake through the inhibition of α -glucosidase, modulation of gut microbiota composition (through antimicrobial activity) and antioxidant properties have also been reported, among others (Refs 125, 126, 127, 128). In contrast, the antihyperlipidemic efficacy and mechanism of berberine are comparatively less understood and will require additional study. Importantly, the compound suffers from relatively poor bioavailability and novel formulations to improve its pharmacological profile will be of interest (Ref. 129). Further clinical and mechanistic studies, especially high-quality and unbiased clinical trials, will be crucial for the continued development of TCM-derived antidiabetic treatments such as berberine as well as other promising formulations (Refs 130, 131). In any case, the development of TCM for both the management and prevention of diabetes remains widespread and an ongoing subject of great research interest in China, and should be worthwhile of continued observation for contemporary applications.

Salvia miltiorrhiza in the treatment of cardiovascular diseases

Cardiovascular and cerebrovascular diseases, especially ischemic heart disease and stroke, remain consistently among the leading causes of death worldwide. In China, the use of TCM medication and principles in treating such conditions is among the most developed fields in TCM practice and research. Generally termed 'huo xue hua yu' or 'activating blood circulation and removing

blood stasis', this branch of TCM makes use of a wide range of herbs and formulations to achieve antithrombotic, antiplatelet aggregation, vasodilative and cardioprotective effects (Ref. 132). Among the many Chinese medicines used for such purposes, the root of the S. miltiorrhiza or 'Danshen' is particularly known in China as well as other parts of Asia for its historical and contemporary usage in the treatment of many cardiovascular and cerebrovascular diseases (Ref. 133). Remarkably, a complex formulation of Danshen, the Compound Danshen Dripping Pill, holds the distinction of being the first TCM-derived product that was approved in 1997 for phase II clinical trials by the US Food and Drug Administration (FDA) (Ref. 134). Phase II trials for the application of the product (trademarked 'Dantonic') for the prevention and treatment of stable angina were completed with positive results in 2010 (Ref. 134), and phase III trials were completed in 2016. The Compound Danshen Dripping Pill, Danshen products in other forms including Danshen-based drug injections have been the subject of extensive clinical and mechanical study in China. Positive clinical findings have been reported for the application of Danshen-based medication for ischaemic stroke, angina pectoris, intercranial haematoma and acute myocardial infarction, among others (Refs 135, 136, 137, 138). In particular, 16 out of 16 published meta-analyses on the use of Compound Danshen Dripping Pills for the management of coronary heart disease have reported positive findings on efficacy and safety, although additional high-quality and unbiased data remains necessary for further judgment (Ref. 138).

This recognition of Danshen and Danshen-derived formulations by the FDA is noteworthy due to an important distinction between Danshen and other previously recognised TCM-derived drugs. Unlike artemisinin, ATO or even berberine, which are clearly defined singular compounds purified from some herb, mineral or mixture, Danshen extracts and compound formulations contain multiple bioactive components with complex mechanisms of action which is characteristic of most medicinal herbs (Ref. 139). This property is conventionally considered undesirable in modern rational drug design, which values welldefined molecular targets and mechanisms for each drug. This represents a contrast in ideology compared to Chinese medicine, which instead focuses on a holistic approach based on the overall symptoms and presentation of each individual patient and uses complex mixtures of herbs and other forms of treatment (Ref. 6). Compared with previous examples of successful drugs developed from Chinese medicine, the fact that the complex formulation of Danshen-based medication appears to also be the most effective is perhaps indicative that the Chinese philosophy can also be valid, at least on occasion (Refs 140, 141).

To date, at least 50 components have been purified and identified from Danshen extracts, which can be largely categorised into hydrophilic compounds (including various polyphenolic acids) and lipophilic compounds (mostly from the tanshinone family of diterpenes) (Ref. 139). Several of these are relatively wellstudied and believed to contribute significantly to the pharmacological effects of Danshen. The phenolic acids including Danshensu, rosmarinic acid, caffeic acid and the salvianolic acids A and B have been reported to exhibit antioxidant, anticoagulant, vasodilative and cardioprotective effects, in addition to other biological activities including antineoplastic and hepatoprotective effects (Refs 139, 142). In particular, Danshensu (salvianic acid A) and salvianolic acid B have both been proposed as major contributors to the vasodilative and anticoagulant effects of Danshen extracts, acting through the modulation of calcium influx and the immune response, among other mechanisms (Refs 143, 144, 145, 146). Among the lipophilic compounds, the tanshinones which are mostly unique to the Salvia family (represented by tanshinone I, tanshinone IIA and cryptotanshinone) are well-studied and have

been shown to exhibit significant antibacterial, antiinflammatory and antioxidant properties in addition to cardioprotective effects (Ref. 139). Outside these comparatively well-understood compounds, other constituents of Danshen have also been reported to contribute to the overall pharmacological effects of Danshen through multiple mechanisms including calcium modulation and the Akt-eNOS pathway (Refs 147, 148, 149). The chemical and biological properties of the prominent constituents of Danshenbased medicines are comprehensively reviewed in other authoritative publications, including a recent comprehensive characterisation of Danshen-based injection by mass spectrometry (Refs 139, 141, 142, 150). Despite the volume of available research, much remains to be understood with regards to the less-understood constituents and the interactions between the various active components of Danshen. Nevertheless, it appears evident that a combination of many bioactive components is necessary for the full efficacy of Danshen and other such herbal formulations. Pending the results of current and further clinical study, TCM-derived treatments of cardiovascular disease exemplified by Danshen should continue to be an area of great interest.

Conclusion

Evidence-based TCM and the integration of TCM principles and medication with modern science and medicine is an area of tremendous ongoing interest and effort. As an overview of the contributions of the entire TCM practice to modern medicine, we have attempted to select several representative treatments in various stages of development to illustrate the progress that has been made in each respective field. However, it is important to reiterate that this is only a tiny representation of the available data and the ongoing effort in TCM research extending to a wide range of diseases. Where possible, we have included references to comprehensive and authoritative reviews that cover the respective fields in great detail and rigor.

One of the means by which TCM contributes to world medical development is to provide effective monomer chemical drugs. TCM is characterised by individual adjustment of multiple components and multiple targets which enables the body to transform from an abnormal to normal state. More research in this respect is yet to be carried out. Moving ahead, it will be important not only to appreciate the existing successes and the potential of TCM for contributing to modern medicine but also to work towards better realising this potential. Given the widespread nature of TCM usage in China for all manners of diseases, it is unsurprising that clinical data are often available in large abundance. However, the quality of clinical studies in terms of study methodology and the elimination of bias have remained significant areas of concern for TCM clinical research (Ref. 151). High-quality clinical data in conformance with international standards such as the CONSORT and TREND statements are absolutely essential for the further development and acceptance of TCM, and the improvement of clinical data quality will be a critical step for TCM research looking forward. At the same time, the advancement of 'omics' and high-throughput investigative techniques such as proteomics and genomics have led to significant progress in drug target identification and the elucidation of drug mechanisms, which remain crucial to modern rational drug design (Refs 152, 153). Considering the complex, multi-component nature of most TCM herbal treatments, the application of such investigative techniques to more complex formulations should yield great insights not only to drug mechanisms but also to the understanding of TCM principles and philosophies with the potential to greatly contribute to modern medicine. While certain aspects of TCM may remain as fringe topics in the context of modern medicine, the examples covered in this review have

hopefully served to illustrate the potential within TCM herbals and by extension the practice of TCM as well as traditional medicinal systems as a whole. It would be a mistake to completely discount the value of experience accumulated over thousands of years based on specific aspects of the practice. The continued application and integration of scientific principles and techniques with traditional medicine will surely continue to serve modern medicine well.

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