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Review Article

Arsenic, Cadmium, Lead, and Mercury in Sweat: A Systematic Review

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Arsenic, cadmium, lead, and mercury exposures are ubiquitous. These toxic elements have no physiological benefits, engendering interest in minimizing body burden. The physiological process of sweating has long been regarded as "cleansing" and of low risk. Reports of toxicant levels in sweat were sought in Medline, Embase, Toxline, Biosis, and AMED as well as reference lists and grey literature, from inception to March 22, 2011. Of 122 records identified, 24 were included in evidence synthesis. Populations, and sweat collection methods and concentrations varied widely. In individuals with higher exposure or body burden, sweat generally exceeded plasma or urine concentrations, and dermal could match or surpass urinary daily excretion. Arsenic dermal excretion was severalfold higher in arsenic-exposed individuals than in unexposed controls. Cadmium was more concentrated in sweat than in blood plasma. Sweat lead was associated with high-molecular-weight molecules, and in an interventional study, levels were higher with endurance compared with intensive exercise. Mercury levels normalized with repeated saunas in a case report. Sweating deserves consideration for toxic element detoxification. Research including appropriately sized trials is needed to establish safe, effective therapeutic protocols.

1. Introduction

No person is without some level of toxic metals in their bodies, circulating and accumulating with acute and chronic lifetime exposures. An individual may take numerous measures to minimize exposures and to optimize metabolism and excretion of toxic elements in the stool and urine with diet, supplements, and chelation therapy [1, 2]; however, an often overlooked route of excretion of toxicants is via the process of sweating [3].

Sweating with heat and/or exercise has been viewed throughout the ages, by groups worldwide, as "cleansing." As part of a scoping review regarding arsenic, cadmium, lead, and mercury, we reviewed the scientific literature pertaining to toxicant excretion in sweat.

1.1. Arsenic, Cadmium, Lead, and Mercury: Background. While many chemical elements are essential for life, arsenic,

cadmium, lead, and mercury have no known beneficial effect in humans. On the contrary, all four elements are confirmed or probable carcinogens, and they exhibit wideranging toxic effects on many bodily systems, including the nervous, endocrine, renal, musculoskeletal, immunological, and cardiovascular systems [4–7].

Children and the fetus are most at risk of harm, with early exposures potentially predisposing the youngster over his/her lifetime to multisystem ailments, as well as lower IQ and dysfunctional behavior. In older populations there is increased likelihood of early cognitive decline, as well as a range of conditions including kidney and cardiovascular disease, diabetes, and osteoporosis [4–7].

Some populations are exposed to elevated levels of toxic elements by virtue of geochemistry, resulting in groundwater or foods with elevated levels of toxic elements (e.g., elevated arsenic in groundwater, most famously in parts of Asia such as Bangladesh but also elsewhere; cadmium that accumulates

in foods grown in particular locations with high levels in soils or from fertilizers, including shellfish [8], grains [9], and brassicas [10]; and mercury in fish and seafoods). Tobacco avidly accumulates cadmium and lead from soil, making smoking a major source of exposure. In addition, valuable and unique properties of arsenic, cadmium, lead, and mercury have made them integral in many products, including electronics, batteries, and alloys. Modern environmental exposures arise from mining, refining, and industrial processes (e.g., arsenic from precious metal mining and refining, mercury from chloralkali production, or lead and cadmium from mining, refining, and recycling these and other metals such as zinc); the vestiges of older products (e.g., pesticides, leaded gasoline, paint and plumbing, mercurycontaining switches and thermometers, and arsenical wood preservatives); ongoing uses (e.g., arsenical veterinary drugs, and mercury-containing dental amalgams, preservatives, and lamps); as well as emissions from burning coal and other incineration (including cremation).

With toxic elements ubiquitous in our air, water, food, and the physical environment, as well as in many consumer products, prudent avoidance is not always possible. Although signs and symptoms of chronic disease are consistent with effects of arsenic, cadmium, lead, and/or mercury, physicians commonly have a low index of clinical suspicion, and therefore levels of toxic elements are seldom investigated. Diagnosis may be challenging because multiple chemicals may contribute to subtle effects in chronic illnesses of an individual, and the effects may be synergistic. A recent review called for mercury assessment in all patients presenting with hypertension or any vascular disease [11], but other toxic elements such as lead [12] may also be implicated at levels commonly observed in the population. "Interaction Profiles" [13] compiled by the US Agency for Toxic Substances and Disease Registry report that renal toxicities of mixtures of lead plus mercury are greater than would be predicted knowing the toxicity dose response of the individual elements. Similarly, neurological toxicities of mixtures of lead plus arsenic, lead plus methylmercury, and lead plus cadmium are supra-additive [13].

1.2. Sweating: Background. Increasing the thermal load on the body activates heat loss mechanisms including increased circulation throughout the skin and sweating [14], with blood flow to the skin increasing from a baseline of 5–10%, to 60–70% of the cardiac output [15]. Maximal sweating occurs within 15 minutes and the fluid loss may be as high as 2 L/h in an "acclimatized" person who regularly sweats [16].

Eccrine sweat is produced in tubular coil glands under the skin surface in response to heat and, or work stress. Capillaries as well as adjacent adipose tissue may contribute to secretions from sebaceous and apocrine glands, as has been seen in research using sweat patches to detect drugs of abuse [17]. Sweat arises from the blood supply to the sweat gland, but is not simply an ultrafiltrate of blood plasma; sodium and chloride are lower in sweat than in serum, as salt loss is restricted by reabsorption in the gland [18]. Both the concentration and total loss of salt (sodium chloride) in sweat vary widely among individuals [19], as well as with acclimatization to exercise and heat [20]. In an early study, Robinson et al. demonstrated that with serum salt depletion the kidneys responded within hours by restricting excretion into the urine, while the sweat glands responded only after days with decreased concentrations in the sweat [21]. Potassium, urea, ammonia, and lactic acid concentrations are higher in sweat than in plasma, although these levels are also regulated to some extent by reabsorption in the ductal tubule of the sweat gland [22]. In one study of successive exercise sessions with cool-down breaks, over the short-term sodium, potassium, calcium, and magnesium excretion in sweat remained constant, while zinc excretion fell [23]. It is unclear whether reabsorption or depletion of plasma supply resulted in diminishing zinc losses.

Children, with greater surface area in comparison to body mass, have been observed in research studies to sweat less than adults, with sweating increasing through puberty [24]. Although some research has indicated that children's thermoregulation and heat tolerance may be less robust than adults, these findings may be at least in part an artifact of study designs and models for interpretation [25]. In research involving exercise and heat, it may be a challenge to maintain ongoing, consistent motivation among children.

2. Methods

2.1. Search Strategy. Medline, Embase, Toxline, Biosis, and AMED were searched, with no restriction on date or language, to March 22, 2011. These records were supplemented with searches for other research by key authors, searches of citations and reference lists of key reports, and "related articles."

Neither sweating nor toxic elements are exclusively modern topics of research, so in order to search older literature for all chemical forms, the online version of the Chemical Rubber Company Handbook was searched for all arsenic, cadmium, lead, and mercury compounds, and lists of keywords were extracted from these lists. Searches using these keywords yielded records that were not identified in searches using the four chemical abstracts service (CAS) numbers or the medical subject headings (MeSHs) for arsenic, cadmium, lead, and mercury. CAS numbers and MeSHs are intended for specific individual chemicals or records referring to unspecified compounds—the tool cannot simultaneously be both specific and general. Toxic element records were searched for terms related to sweating, perspiration, sauna, steam baths, exercise, depuration, and secretion or excretion from skin. Bibliographic records were imported, duplicates were removed, and reports were screened using Zotero 2.03 (http://www.zotero.org/).

2.2. Report Screening and Inclusion. Titles and abstracts were screened by one investigator (MS), for primary reports with data on one or more of the toxic elements in sweat, with at least a substantial abstract in English. Reviews were included at this level, to search reference lists. Two investigators (MS and KK) independently screened studies

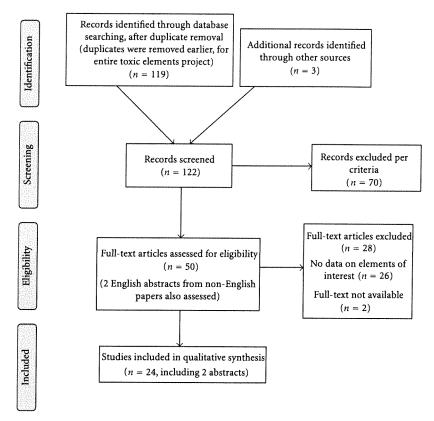


FIGURE 1: PRISMA flow diagram of evidence searches and inclusion.

for inclusion, and extracted and verified data. All studies presenting quantitative human data on levels of arsenic, cadmium, lead, and/or mercury were included, regardless of experimental design, or methods of sweat collection or chemical analysis.

3. Results

Of 122 bibliographic records identified, 70 did not meet inclusion criteria at first screening, 52 full-text articles were sought for full-text screening, and 50 were obtained and screened. Data from the extended abstract of a report in German [26] and the conclusion from the abstract of one report in Russian [27] that were not obtained in full text were noted. Twenty-four reports of 22 or 23 trials or studies (it is unclear if two studies from one institution reported results twice for a subset of participants [22, 28]) were included in evidence synthesis. Searching, screening, and study inclusion are summarized in the modified PRISMA flow diagram, Figure 1.

3.1. Excretion of Toxic Elements in Sweat. Along with essential minerals, sweat is an acknowledged excretory route for toxic metals. For instance, it is recommended to sample hair close to the scalp because content of toxic elements may be elevated along the shaft, from either environmental

contamination or excreted toxins in sweat and sebum [32, 42]. The minerals generally arise from blood serum [28], with contribution from dermally absorbed occupational exposures, which might not be reflected in blood or urine [35, 37]. Sweating was induced by sauna, exercise, or pilocarpine iontophoresis to measure the concentration of the heavy metals in the sweat, while sauna and exercise were used for therapy. Study participants included workers with occupational exposures and individuals with no occupational exposures who were well or experiencing chronic ill health, and in two studies participants were intentionally dosed with lead [34, 37]. Studies that have examined the presence of toxic metals in sweat are summarized in Tables 1, 2, 3, and 4, for arsenic, cadmium, lead, and mercury, respectively.

Arsenic accumulates highly in the skin, and causes characteristic skin lesions, but little information is available on levels in sweat. Yousuf et al. recently found that excretion of arsenic was greatest from the skin of patients with skin lesions, slightly but not statistically significantly lower from arsenic-exposed controls, and severalfold lower from nonexposed controls [29]. Genuis et al. measured numerous toxic elements in blood plasma, urine, and sweat of 20 study subjects (10 healthy and 10 with chronic health problems) [3]. The maximum sweat arsenic concentration was $22 \,\mu g/L$. On average, arsenic was 1.5-fold (in males) to 3-fold (in females) higher in sweat than in blood plasma; however,

TABLE 1: Studies of excretion of arsenic in sweat.

Study	Country, participants	Study design and intervention	Key findings (concentrations of μ g/L unless otherwise indicated)
Yousuf et al. 2011 [29]	Bangladesh 20 arsenicosis patients with melanosis and leucomelanosis 20 controls with As in drinking water 20 unexposed controls	Secretions from chest, back, and abdomen collected for 24 h, on gauze pads (8-fold; 2 × 3 inches) attached to fitted T-shirt	As secretion severalfold greater for As-exposed groups No significant difference between patients and As-exposed controls 2 zinc atoms excreted per As atom Vitamin E excreted with As
Genuis et al. 2010 [3]	Canada 10 with chronic conditions 10 healthy	Simultaneous measurement of As in blood plasma, urine, and sweat Sweating induced by exercise or sauna, collected directly into bottle	17 participants with As detected in all comples

Table 2: Studies of cadmium excretion in sweat.

Study	Country, participants	Study design and intervention	Key findings (concentrations μ g/L unless otherwise indicated)
Genuis et al., 2010 [3]	Canada 10 with chronic conditions 10 healthy	Simultaneous measurement of toxic trace elements in blood plasma, urine, and sweat Exercise or sauna Sweat collected directly into bottle	3 participants with cadmium detected in all samples Blood plasma mean: 0.03 (range 0.02–0.07) $(n = 11)$ Urine mean: 0.28 $(0.18-0.39)$ $(n = 3)$ Sweat mean: 5.7 $(0.36-36)$ $(n = 18)$
Omokhodion and Howard, 1994 [30]	UK 15 healthy participants	Sweat collected using modified arm bag (hand excluded) Participants exercised at room temperature	Cadmium detected in 13 sweat samples Mean 1.9 Range 1.1–3.1
Stauber and Florence, 1988 [28]	Australia 24 males 13 females taking oral contraceptives 26 females not taking oral contraceptives	Forearm sweat induced by pilocarpine iontophoresis and collected on a membrane filter	Males mean sweat cadmium 1.4 (range <0.5–10) Females not taking contraceptives 2.6 (<0.5–18) Females taking contraceptives 2.4 (<0.5–5.5)
Stauber and Florence, 1987 [22]	Australia 9 males 7 females taking oral contraceptives 6 not taking oral contraceptives (unclear overlap with 1988 participants)	Forearm sweat induced by pilocarpine iontophoresis and collected on a membrane filter	Cadmium not detected in sweat (0.5 detection limit) Mean blood cadmium 0.8
Robinson and Weiss, 1980 [31]	USA 28 males (university faculty members)	Exercise and shower preceded sauna for sweat collection. Sweat collected as drips from forehead or nose	Sweat cadmium (range 11–200) Urine cadmium (range ND–67) Sweat/urine ratio (range 1.0–16) No correlation between the concentrations in urine and sweat
Robinson and Weiss, 1980 [32] (companion to previous)	USA 2 males (university faculty members)	As previous, cadmium also measured in hair segments.	Daily excretion of cadmium estimated as follows: (i) 30 µg/day in urine (ii) 120 µg/day in sweat (iii) 0.2 µg/day in hair Cadmium concentrations in hair and sweat were lower in one participant than the other
Cohn and Emmett, 1978 [33]	USA 6 males 3 females	Total body washdown and arm bag techniques	Mean concentration of cadmium in sweat > urine Arm bags yielded lower levels than whole body measurements

TABLE 3: Studies of lead excretion in sweat.

Study	Country, participants	Study design and intervention	Key findings (concentrations μ g/L unless otherwise indicated)
Genuis et al., 2010 [3]	Canada 10 with chronic health conditions 10 healthy	Analyses of blood plasma, urine, and sweat Sweating induced by exercise or sauna, collected directly into bottle	Sweat mean 31 (range 1.5–94) ($n = 20$) Blood plasma mean 0.12 (0.39–1.7) ($n = 20$) Urine mean 1.8 (0.91–7.5) ($n = 20$)
Omokhodion and Crockford, 1991 [34]	UK 2 participants	Blood, urine, and sweat lead measured before and following ingestion of lead chloride: 1 or 2 doses of lead chloride (20 mg PbCl ₂ total, in 1 or 2 divided doses).	Blood lead peaked at 4 h Sweat concentrations did not increase significantly (range 0–11) Blood concentration range 6–51 Urine concentration range 10–97 Arm sweat collections varied by more than 2-fold between arms at the same time on the same person
Omokhodion and Howard, 1991 [35]	Unidentified "tropics" 19 workers in a lead battery factory 8 controls (medical students)	Measured lead in sweat, blood, and urine simultaneously Sweating induced by exercising at room temperature. Sweat collected in arm bags.	Workers: (i) blood lead 13–36 (ii) urine lead 28–290 μg/g creatinine (iii) sweat lead 72–260 Controls: (i) blood lead 90–120 (ii) urine lead 9–20 μg/g creatinine (iii) sweat lead 9–30
Omokhodion and Crockford, 1991 [36]	UK 24 normal, healthy subjects	Measured lead in sweat, urine, blood, and saliva Sweat collected in arm bags, sitting in a hot chamber	 (i) Blood lead 86 (range 60–140) (ii) Urine lead 18 μg/g creatinine (range 7.7–44 μg/g creatinine) (iii) Mean sweat lead 5.2 (2.5–13) (iv) Saliva lead 4.8 (2.5–10)
Parpaleĭ et al., 1991 [27] (in Russian—English abstract only)	Russia NR in abstract	NR in abstract	" sauna increased excretion with sweat fluid of toxic substances [lead] that penetrated the body during work. Sauna is recommended."
Lilley et al., 1988 [37]	Australia 9 lead workers volunteers had lead applied to skin	Lead dust 6 h/day for 4 days 20 mg Pb dust on L arm of volunteer PbNO ₃ 24 h of 60 mg PbNO ₃ on L arm of volunteer.	Sweat lead in workers: 71–18,000 Following exposure, sweat lead from R arm increased approximately by 10x, returning to baseline after approximately by 2–4 days. Saliva increased approximately 5-6x. Urine and blood levels were unchanged
Stauber and Florence, 1988 [28]	Australia 24 males 13 females taking oral contraceptives 26 not taking oral contraceptives	Sweating induced on the forearms by pilocarpine iontophoresis and collected on a membrane filter	Mean sweat lead: (i) males: 41 (range 6–87) (ii) females not taking contraceptives: 24 (<5–66) (difference with males <i>P</i> < 0.01) (iii) females taking contraceptives: 36 (<5–70)
Stauber and Florence, 1987 [22]	Australia 9 males 7 females taking oral contraceptives 6 not taking oral contraceptives (unclear overlap with 1988 participants)	Sweating induced in the forearms by pilocarpine iontophoresis and collected on a membrane filter	No significant differences among groups Mean blood lead 200 Mean blood plasma lead 10 Mean sweat lead 15

Table 3: Continued.

Study	Country, participants	Study design and intervention	Key findings (concentrations μ g/L unless otherwise indicated)
Haber et al., 1985 [26] (in German-used extended abstract)	Germany 4 groups of 8 males 2 groups with occupational lead exposure 2 control groups	Comparison of precisely defined physical work (intensive cycling and extended rowing in a pool), examining lead excretion in persons with elevated blood levels compared with nonexposed controls	Aerobic endurance training (rowing) caused a significant drop in the blood lead level in the occupationally exposed group (mean 430 (range 320–580) decreased to 370 (240–450)) (P < 0.05) Endurance training was more effective than shorter, more intensive training (cycling) Urine lead levels were not significantly affected by training
Cohn and Emmett, 1978 [33]	USA 6 males 3 females	Total body washdown and arm bag techniques	The mean concentration of lead in sweat was similar to that in urine (1) Total body sweat lead mean: (i) males: 24 (SD 16) (ii) females: 53 (range 40–60) (2) Body minus 1 arm/arm bag sweat lead 60 (SD 16) (40–120)/83 (86) (20–250)
Hohnandel et al., 1973 [38]	33 healthy males 15 females	15 min of arm bag collection	Mean sweat lead: (i) males: 51 (range 8–180) (ii) females: 120 (SD 72) (49–280)

Table 4: Studies of mercury excretion in sweat.

Study	Country, participants	Study design and intervention	Key findings (concentrations μ g/L unless otherwise indicated)
Genuis et al., 2010 [3]	Canada 10 with chronic conditions 10 healthy	Sweating induced by exercise or sauna, collected directly into bottle	16 participants had mercury detected in all samples Blood plasma mercury mean 0.61 (range 0.26–1.6) ($n = 16$) Urine mean 0.65 (range 0.32–1.3) ($n = 16$) Sweat mean 0.86 (range 0.48–1.5) ($n = 20$)
Robinson and Skelly, 1983 [39]	USA 21 males at university 7 sampled more than once	Mercury in sweat dripping from forehead or nose, compared with urine	Sweat mean 0.5 (range 0.1–1.4)
Sunderman 1978 [40]	USA 1 case with mercury intoxication	Case report of chelating agents to treat mercury intoxication, followed by a regimen of daily sweat and physiotherapy for a protracted period of several months	Appreciable quantities of mercury were excreted in sweat. With the sweating regimen mercury, levels in sweat decreased to within the normal range
Lovejoy et al., 1973 [41]	USA 3 mercury-exposed workers 3 nonexposed workers 1 control	Participants wore rubber chest waders from 7:30 to 9:00 am Sweat accumulated in the feet was collected, as well as a 16-hour urine sample	Exposed workers: 1.5 h sweat: 120–350 ng mercury 16 h urine: 160–190 ng mercury Unexposed workers: 1.5 h sweat: 5–8 ng mercury 16 h urine: 5–7 ng mercury Internal controls: 1.5 h sweat: 43–70 ng mercury 16 h urine: 30–46 ng mercury Mercury concentrations in sweat > urine for exposed workers; similar for controls

arsenic was excreted at lower concentrations in sweat than in urine [3].

Cadmium in sweat was examined in six studies [3, 22, 28, 30–33], with concentrations in sweat ranging from <0.5–10 μ g/L [28] to 0.36–35.8 μ g/L [3]. Stauber and Florence concluded that sweat may be an important route for excretion of cadmium when an individual is exposed to high levels [22, 28], a finding that was confirmed by observing that the total daily excretion of cadmium was greater in sweat than in urine [3, 32]. The maximum cadmium concentration observed in sweat was 35.8 μ g/L [3].

Lead was examined in eleven studies [3, 22, 26-28, 33-38]. In 1973, Hohnadel et al. suggested that "sauna bathing might provide a therapeutic method to increase elimination of toxic trace metals" [38]. In two males, 36% and 50% of sweat lead was of molecular weight > 30,000, as measured by ultrafiltration, suggesting excretion of organic complexes rather than simple ions [22]. Lead excretion was lower in females taking birth control medications compared with females not taking medications, or males [28]. Haber et al. found that prolonged endurance workouts (rowing) ameliorated elevated blood lead levels in exposed workers but did not alter levels in control subjects and did not affect urine levels [26]. They suggested that the elimination route was not urine, but potentially sweat or/and bile. Omokhodion and Crockford carried out several studies of trace elements in sweat, including a study of lead ingestion by two human participants [34]. Sweat lead levels did not increase immediately with elevated blood lead, although the authors make reference to an older study with longer followup wherein lead in underarm pads doubled in the five days following ingestion. Omokhodion and Howard also reported higher lead in sweat of exposed workers compared with unexposed controls [35], and in another study that sweat and blood lead levels were the only two variables that correlated among blood, urine, sweat, and saliva [36]. The English abstract of a 1991 case report in Russian indicated that sauna increased excretion of toxic elements and resulted in clinical improvements [27]. Sweat lead levels up to $283 \mu g/L$ have been observed in nonoccupationally exposed subjects [38] and up to $17,700 \mu g/L$ in workers [37], where it is noted that lead in sweat may partially originate from material absorbed within the skin that was not removed by pretest cleaning protocols [35]. Indeed, although dermal application of lead via hair follicles, sweat ducts, and diffusion does not result in immediate increases in blood or urine lead concentrations, dermal absorption was demonstrated using the Pb-204 isotope [43], lead powder, and salt [37].

Mercury. In 1973, Lovejoy et al. noted that exposure to mercury does not always correlate with urine mercury levels and that elimination by other routes such as sweat may be an explanation [41]. They suggested, "sweating should be the initial and preferred treatment of patients with elevated mercury urine levels." In a 1978 case report, a severely poisoned worker was rescued with chelation therapy, followed by a regimen of daily sweat and physiotherapy over several months during which the sweat mercury level returned to normal and the patient recovered [40]. Robinson

measured mercury in sweat repeatedly in two volunteers, observing sweat to urine concentration ratios ranging from less than 0.1 to greater than 5. Sweat mercury concentrations varied widely from day to day, and there was no correlation with urine levels. Sweat mercury levels of $1.5 \,\mu\text{g/L}$ were observed by Genuis et al. [3] and $1.4 \,\mu\text{g/L}$ by Robinson and Skelly [39].

4. Discussion

Arsenic, cadmium, lead, and mercury may be excreted in appreciable quantities through the skin, and rates of excretion were reported to match or even exceed urinary excretion in a 24-hour period. This is of particular interest should renal compromise limit urinary excretion of toxic elements.

Most of the research identified was over 20 years old, and collection methods varied widely. Although authors described thorough precleaning methods, sweat concentrations measured in research settings are not well validated and varied according to the location on the body, collection method, and from day to day according to other variables such as hydration. Sweat contains metals not only from the blood plasma, but also evidently originating from dermal layers (particularly with significant dermal exposures, as for workers in welding, smelting, or battery manufacturing). It would appear that large variabilities in measured concentrations, apart from collection methods as mentioned above, were likely the result of differences in excretion amongst widely varying individuals with ranges of body burdens, genetic polymorphisms affecting detoxification efficiency, and physiological states, coupled with necessarily crude if simple experimental techniques. These variations were very much greater than would be expected due to limitations of analytical methods. Although analytical methods have improved over the years, analysis of these metals was commonplace at the time of the studies. Authors generally reported analytical methods rigorously or provided references to thorough descriptions and included internal standards and some indication of sensitivity.

The observation that between a third and a half of lead in sweat may be associated with high-molecular-weight molecules [22] merits replication, including examination of additional toxic elements and characterization of the associated molecules previously observed. Excretion of these large molecules also suggests that sweating may be a means of excretion of metals complexed with natural or synthetic chelating agents.

Yousuf et al.'s recent study demonstrating a 2:1 molar ratio of zinc: arsenic and increased vitamin E in skin secretions suggests potential therapeutic supplementation to accommodate these biochemical requirements. Vitamin E, zinc, and other nutrients are required for methylation and detoxification of arsenic within the body, and vitamin E supplementation improves the skin manifestations in arsenicosis [29].

From an occupational health perspective, lead, and presumably other toxic elements, may be absorbed via the skin, which supports showering at work and further suggests the possibility of purging workers' skin by washing with a chelating agent (e.g., EDTA rinses extracted lead from workers' skin in methods validation experimentation [38]). It is unknown if sweating during the workday may affect dermal absorption, or if forced sweating at the end of the workday would be beneficial. It is also unknown if increased blood flow to the skin could possibly enhance absorption into the bloodstream, or if worker health could be optimized by a combination of workplace skin cleaning and sweating interventions.

Sweating has long been perceived to promote health, not only accompanying exercise but also with heat. Worldwide traditions and customs include Roman baths, Aboriginal sweat lodges, Scandinavian saunas (dry heat; relative humidity from 40% to 60%), and Turkish baths (with steam). Infrared saunas heat exposed tissues with infrared radiation, while air temperatures remain cooler than in other saunas.

Sweating is a long-standing, if recently forgotten, aspect of mercury detoxification. Various strategies used to maintain the mercury mining workforce have been explored over the centuries. In Spain and colonies, long the western world's primary sources of mercury, sending ill workers to warmer climes away from the exposure to drink weak beer (the hydrogen peroxide catalase oxidation of elemental mercury to ionic mercury is competitively inhibited by alcohol, increasing mercury in exhaled breath [44]) and to work in the heat (presumably to sweat out the "vapors") was a common and effective strategy centuries ago; tremors, salivation, and mouth ulcers resolved generally within a few weeks [45].

With acclimatization and regular use, the sauna is generally well tolerated by all ages [46], though medical supervision may be recommended during initial sessions for children, the elderly, or those with compromised health. Varying qualities of evidence indicate potential short- and long-term improvements for cardiovascular, rheumatological and respiratory conditions; contraindications include unstable angina pectoris, recent myocardial infarction, severe aortic stenosis, and high-risk pregnancy [15, 46]. Sweating is not only observed to enhance excretion of the toxic elements of interest in this paper, but also may increase excretion of diverse toxicants, as observed in New York rescue workers [47], or in particular persistent flame retardants [48] and bisphenol-A [49].

Optimizing the potential of sweating as a therapeutic excretory mechanism merits further research. To date, the large body of research into homeostasis of the most common metals (sodium, potassium, and to a lesser extent, magnesium, calcium, and zinc) and conditioning or adaptation to regular sweating by athletes has not been matched with studies of excretion of trace elements. Limited research suggests indirectly that conditioning may not restrict excretion of nonessential elements. Combination therapies, such as administration of *n*-acetyl cysteine, vitamin C, a chelating agent, or low doses of ethanol (for mercury), to name a few possibilities, along with sauna and/or exercise therapy to induce sweating, may be fruitful avenues of investigation.

It has been noted that among people whose health is compromised by toxicants, heat regulatory mechanisms of the autonomic nervous system are often affected, resulting in a failure to sweat readily [3]. In these cases, along with diet and nutritional supplementation to remediate biochemical imbalances, interventions to consider include brushing the skin, niacin to assist with vasodilation, and exercise prior to sauna use [50]. Clinical experience is that with persistence and ample hydration patients do eventually start to sweat. This is often a sign that the autonomic nervous system function is beginning to improve. With enhanced ability to sweat, detoxification is facilitated, which can ultimately result in clinical improvement.

For biomonitoring and research purposes, modern validated methods are desirable to collect and measure elements in sweat, so this means of excretion may be considered in the context of other measures such as urine, blood, feces, and hair concentrations. Considerations for dry and wet collection methods were recently discussed in the context of essential solutes [51, 52].

Undoubtedly further research in this area would improve understanding, but the available evidence suggests that physicians could consider recommending sweating as tolerated via exercise (preferred) and/or use of a sauna as a lowrisk, potentially beneficial treatment for individuals who may be experiencing effects of toxic elements, or for individuals with regular exposure to or accretion of toxicants.

5. Conclusions

Sweating offers potential and deserves consideration, to assist with removal of toxic elements from the body. As toxic elements are implicated in many serious chronic conditions, research is needed in patients with select conditions to evaluate the body burden and to test the efficacy of source removal, dietary choices and supplements, interventions that induce sweating, and treatments with drugs, all to enhance excretion of toxic elements with the goal of clinical improvement. There is a clear need for robust trials, appropriately sized to assess clinical outcomes, from which therapeutic protocols can be derived. Both biochemical and clinical outcomes should be examined in order to develop and monitor clinical interventions that are both safe and effective.

Conflicts of Interests

The authors declare that they have no conflicts of interests.

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Theme Section: Emerging Therapeutic Aspects in Oncology

REVIEW

Curcumin: an orally bioavailable blocker of TNF and other pro-inflammatory biomarkers

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TNFs are major mediators of inflammation and inflammation-related diseases, hence, the United States Food and Drug Administration (FDA) has approved the use of blockers of the cytokine, TNF- α , for the treatment of osteoarthritis, inflammatory bowel disease, psoriasis and ankylosis. These drugs include the chimeric TNF antibody (infliximab), humanized TNF-α antibody (Humira) and soluble TNF receptor-II (Enbrel) and are associated with a total cumulative market value of more than \$20 billion a year. As well as being expensive (\$15 000-20 000 per person per year), these drugs have to be injected and have enough adverse effects to be given a black label warning by the FDA. In the current report, we describe an alternative, curcumin (diferuloylmethane), a component of turmeric (Curcuma longa) that is very inexpensive, orally bioavailable and highly safe in humans, yet can block TNF- α action and production in in vitro models, in animal models and in humans. In addition, we provide evidence for curcumin's activities against all of the diseases for which TNF blockers are currently being used. Mechanisms by which curcumin inhibits the production and the cell signalling pathways activated by this cytokine are also discussed. With health-care costs and safety being major issues today, this golden spice may help provide the solution.

LINKED ARTICLES

This article is part of a themed section on Emerging Therapeutic Aspects in Oncology. To view the other articles in this section visit http://dx.doi.org/10.1111/bph.2013.169.issue-8

Abbreviations

ACR, American College of Rheumatology; AP-1, activator protein-1; ATF2, activating transcription factor 2; C/EBP, CCAAT/enhancer binding protein; CD, Crohn's disease; COPD, chronic obstructive pulmonary disease; CREB, cAMP response element binding protein; DAS, Disease Activity Score; DNBS, dinitrobenzene sulfonic acid; IBD, inflammatory bowel disease; LITAF, LPS-induced TNF-α factor; MCP-1, monocyte chemotactic protein-1; MD-2, myeloid differentiation protein-2; NFAT, nuclear factor of activated T-cell transcription factor; OA, osteoarthritis; PhK, phosphorylase kinase; RA, rheumatoid arthritis; SLCP, solid lipid curcumin particle; TLRs, toll-like receptors; TNBS, trinitrobenzene sulfonic acid; UC, ulcerative colitis

Introduction

Extensive research during the past century has revealed that inflammation plays a major role in most chronic diseases. It was Cornelius Celsus, a physician in first century Rome, who first attempted to describe inflammation as heat (calor), pain (dolor), redness (rubor) and swelling (tumour). Rudolf Virchow, a German scientist from Wurzburg, in 1850, was the first to observe a link between inflammation and various chronic diseases, which include cancer, atherosclerosis, arthritis, diabetes, asthma, multiple sclerosis and Alzheimer's disease (Heidland et al., 2006). More than 200 different types of inflammatory disease have been described. When the name of a disease ends with 'itis', this means inflammation of the affected organ. Thus, arthritis is inflammation of the joints, whereas bronchitis, sinusitis, gastritis, oesophagitis, pancreatitis, meningitis, rhinitis and gingivitis are, respectively, inflammation of the bronchi, sinuses, stomach, oesophagus, pancreas, brain, nose and gums. Acute inflammation is thought to be therapeutic as it helps an organism to heal. Chronic inflammation, however, can lead to a disease; inflammation of the colon (colitis), for example, when



persistant, for as long as 30 years, can finally lead to colon cancer.

During the past three decades, molecular mechanisms that lead to inflammation have been extensively examined. Various enzymes, cytokines, chemokines and polypeptide hormones have been identified, which can mediate inflammation. These include COX-2, 5-lipooxygenase (LOX), TNF- α , IL-1, IL-6, IL-8, Il-17, IL-21, IL-23 and monocyte chemotactic protein-1 (MCP-1). Among these, TNF- α is a major mediator of inflammation, which is the primary focus of this review.

Discovery of TNFs

TNF has at various times been called tumour necrosis serum, cachectin, lymphotoxin or monocyte cytotoxin based on work from our laboratory and others. It is now clear that TNF is a 25 kDa transmembrane protein (17 kDa when secreted) produced primarily by activated macrophages. The ability of tumours to undergo haemorrhagic necrosis after injection of endotoxin was first shown by Shear and Perrault (1944). O'Malley et al. (1962) reported that endotoxin injection into normal mice resulted in the appearance of tumour necrotizing activity in the circulating blood. This activity was renamed tumour necrosis factor by Carswell et al. (1975). The true chemical identity of TNF, however, was unclear until our group isolated two different molecules: one from macrophages, which we named TNF-α (Aggarwal et al., 1985b), and the other from lymphocytes, which we named TNF-B (Aggarwal et al., 1984). The current review primarily deals with TNF-α.

Because of the amino acid sequence homology between human TNF- α and endotoxin-induced murine cachectin, a

protein linked to endotoxin-mediated cachexia and shock (Beutler et al., 1985), it became clear that TNF-α and cachectin were identical. Soon thereafter, numerous groups independently identified the same molecule by using a variety of approaches (Haranaka et al., 1984; Old, 1985; Wang et al., 1985; Fiers et al., 1986; Wallach, 1986). TNF-α is now known to bind to two different receptors, TNFRSF1A and TNFRSF1B, and to activate caspase-mediated apoptosis, NF-κB, activator protein-1 (AP-1), JNK, p38 MAPK and ERK signalling (Figure 1). Our group demonstrated that both TNF-α and TNF-β bind to identical receptors and with similar affinities (Aggarwal et al., 1985a). Although much is known about TNF-α, very little is understood about TNF-β (Aggarwal, 2003; Aggarwal et al., 2012). Both overlapping and nonoverlapping activities of the two molecules have been reviewed (Stone-Wolff et al., 1984; Kuprash et al., 2002; Liepinsh et al., 2006).

Aside from originating in monocytes, it is now clear that TNF- α is also produced by a variety of other cell types including Kupffer cells in the liver, astrocytes in the brain, T-cells and beta cells in the immune system, and ovarian cells. In general, under appropriate conditions, most cell types have the potential to produce TNF- α .

TNF- α and inflammation

It is only within the past few decades that the mechanisms by which inflammation is mediated at the molecular level have become apparent. Although the role of macrophages in inflammation has been known for quite some time, the first indication of the pro-inflammatory activity of TNF emerged in 1985 when it was found to stimulate collagenase and PGE₂ production by isolated human synovial cells and dermal fibroblasts (Dayer *et al.*, 1985; Caput *et al.*, 1986), thus sug-

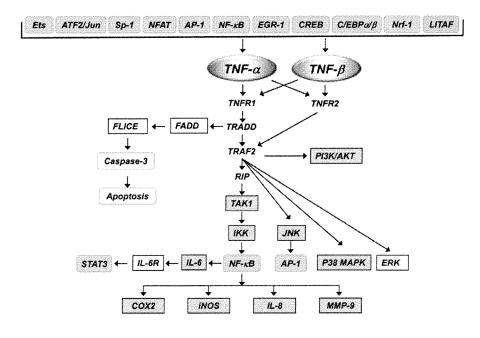


Figure 1
Regulation of the production and action of TNF by curcumin. TNFR1 and TNFR2 are TNF receptors TNFRSF1A and TNFRSF1B respectively. Targets highlighted as yellow are down modulated by curcumin.

gesting that TNF may play a role in the tissue destruction and remodelling associated with inflammatory diseases.

TNF-associated diseases

TNF dysregulation has been linked to a wide variety of diseases including cancer, obesity, cardiovascular diseases, pulmonary diseases, metabolic diseases, neurological diseases, psychological diseases, skin diseases and autoimmune diseases (Aggarwal, 2003; Aggarwal et al., 2012). Thus, blockers of TNF have been approved for the treatment of various autoimmune disorders such as rheumatoid arthritis (RA), ankylosing spondylitis, Crohn's disease (CD), psoriasis, hidradenitis suppurativa and refractory asthma. The inhibition of TNF can be achieved with monoclonal antibodies such as infliximab (Remicade; Janssen Biotech Inc., Horsham, PA, USA), adalimumab (Humira; Abbott Laboratories, North Chicago, IL, USA), certolizumab pegol (Cimzia; UCB, Brussels, Belgium) and golimumab (Simponi; Janssen Biotech) or with a circulating receptor fusion protein such as etanercept (Enbrel; Amgen, Thousand Oaks, CA, USA). Their potential use in other pro-inflammatory diseases is currently being explored. Some of the important adverse effects most extensively associated with TNF blockers include lymphoma, infections, congestive heart failure, demyelinating disease, a lupus-like syndrome, induction of auto-antibodies, injection site reactions and systemic adverse effects (Scheinfeld, 2004).

Suppression of TNF-α production by curcumin *in vitro*

Numerous reports have suggested that the production of TNF from macrophages activated by various stimuli can be suppressed by curcumin (Table 1). Studies have supported findings that LPS is one of the major inducers of TNF- α in macrophages and monocytes and that curcumin can downregulate the expression of TNF-α (Chan, 1995; Abe et al., 1999; Jang et al., 2001; Gao et al., 2004; Strasser et al., 2005; Woo et al., 2007; Liang et al., 2008; 2009; Cheung et al., 2009; Jain et al., 2009; Nishida et al., 2010; Zhao et al., 2010). Besides being expressed by myeloid cells, TNF is also expressed by microglial cells, adipocytes and other cell types. Curcumin, however, has been shown to down-regulate TNF expression (Jin et al., 2007; Lee et al., 2007; Zhang et al., 2008; 2010a). In addition to being induced by LPS, TNF is also upregulated by a variety of other stimuli including phorbol ester, palmitate and other inflammatory cytokines, and curcumin has been shown to block the expression of TNF induced by all of these stimuli (Abe et al., 1999; Lee et al., 2007; Jain et al., 2009; Wang et al., 2009).

How curcumin down-regulates TNF expression in different cell types and in response to a variety of stimuli has been examined extensively. TNF suppression primarily occurs at the transcriptional level. The factors known to be involved in TNF transcription include the transcription factor ETS (Kramer *et al.*, 1995), activating transcription factor 2 (ATF2)/Jun (Leitman *et al.*, 1991; Newell *et al.*, 1994; Tsai *et al.*, 1996a,b), Sp1 (Kramer *et al.*, 1994), nuclear factor of activated

T-cell transcription factor (NFAT) (McCaffrey et al., 1994; Tsai et al., 1996a,b), NF-kB (Udalova et al., 1998; Kuprash et al., 1999), early growth response protein-1 (Kramer et al., 1994), cAMP response element binding protein (CREB) (Geist et al., 1997), CCAAT/enhancer binding protein β (C/EBPβ) (Pope et al., 1994; Wedel et al., 1996; Zagariya et al., 1998), NF-E2related factor 1 (Novotny et al., 1998; Prieschl et al., 1998) and LPS-induced TNF-α factor (LITAF) (Takashiba et al., 1995; Myokai et al., 1999) (Figure 1). Hence, different transcription factors appear to be involved in the stimulation of TNF expression by various stimuli and in different cell types. For instance, the transcription factor LITAF is involved in LPS-stimulated TNF expression. Whereas ATF2/Jun and NFATp are involved in TNF expression in activated beta and T-cells, C/EBPβ is involved in human monocytes. Several of these transcription factors have been shown to be modulated by curcumin.

Curcumin can mediate its effect on TNF expression by inhibiting p300/CREB-specific acetyl transferase, leading to repression of the acetylation of histone/non-histone proteins and histone acetyl transferase-dependent chromatin transcription (Balasubramanyam *et al.*, 2004). It is also well known that curcumin can down-modulate the activation of NF-κB by a variety of agents (Singh and Aggarwal, 1995) and this down-regulation of NF-κB by curcumin plays a major role in suppressing the expression of TNF. In addition, in a number of studies it has been shown that methylation of a TNF promoter may affect the promoter's function (Kochanek *et al.*, 1990; 1991; Muiznieks and Doerfler, 1994; Takei *et al.*, 1996). Thus, curcumin could affect TNF expression by affecting the methylation of a TNF promoter (Reuter *et al.*, 2011).

It is possible that the effects of curcumin on LPS-induced TNF production are mediated in part through its LPS signalling. Two of the toll-like receptors (TLRs), TLR2 and TLR4, mediate responsiveness to LPS. LPS-mediated TLR2 mRNA induction has been shown to be attenuated by pretreatment with curcumin (Matsuguchi et al., 2000). In addition, there is biochemical evidence indicating that curcumin can inhibit both ligand-induced and ligand-independent dimerization of TLR4 (Youn et al., 2006). The beneficial effect of curcumin is partly mediated by reducing the expression levels of TNF through inhibition of the expression of TLR2, TLR4 and TLR9 in mouse liver (Tu et al., 2012). Curcumin also binds with sub-micromolar affinity to the myeloid differentiation protein-2 (MD-2), which is the LPS-binding component of the endotoxin surface receptor complex MD-2/TLR4 (Gradisar et al., 2007). The binding site for curcumin overlaps with that of LPS; this results in the inhibition of MyD88dependent and MyD88-independent signalling pathways of LPS signalling through TLR4, indicating that MD-2 is an important target of curcumin involved in its suppression of the innate immune response to bacterial infection.

Suppression of TNF- α -mediated signalling by curcumin in vitro

There are numerous reports suggesting that curcumin can not only block the production of TNF but also block the cell signalling mediated by TNF in a variety of cell types (Table 1). Our group was the first to show that curcumin can inhibit TNF-mediated NF- κ B action in variety of cell types (Singh and Aggarwal, 1995). We also showed that TNF-mediated expression of various cell surface adhesion molecules in endothelial



Table 1

Curcumin inhibits the production and action of TNF in vitro

Production of TNF

- Inhibited LPS-induced TNF and IL-1 release from macrophages (Chan, 1995).
- Inhibited production of IL-8, MIP-1α, MCP-1, IL-1β and TNF-α by PMA- or LPS-stimulated human monocytes and alveolar macrophages (Abe et al., 1999).
- Inhibited LPS-induced TNF-α release from macrophages (Jang et al., 2001).
- Inhibited the expression/production of IL-12 and TNF-α by peritoneal macrophages (Gao et al., 2004).
- Decreased NF-κB activation and TNF-α secretion after LPS exposure in U-937 cells (Strasser et al., 2005).
- Inhibited the production of IL-1, IL-6 and TNF-α in LPS-stimulated BV2 microglia (Jin et al., 2007).
- Exhibited neuroprotective effects through suppression of NO, TNF-α, IL-1α and IL-6 from Abeta (25–35)/IFN-γ- and LPS-stimulated microglia cells (Lee *et al.*, 2007).
- Inhibited inflammatory responses of adipose tissue in obesity by suppressing release of TNF-α, NO and MCP-1 from adipocytes (Woo et al., 2007).
- Inhibited NO and TNF-α production in rat primary microglia induced by LPS (Zhang et al., 2008).
- Inhibited LPS-induced TNF-α and IL-6 synthesis in macrophages (Liang et al., 2008).
- Down-regulated TNF, IL-1, NO and PGE₂ in Raw 264.7 cells possibly through induction of phase II/antioxidant enzymes including HO-1 and NQO-1 (Cheung *et al.*, 2009).
- Reversed palmitate-induced insulin resistance through suppression of NF-κB, TNF-α and IL-6 in adipocytes (Wang et al., 2009).
- Inhibited LPS-induced production of TNF-α, IL-1β, MCP-1, COX-2, iNOS and p65 NF-κB in the macrophages (Liang et al., 2009).
- Inhibited the high glucose-induced secretion of IL-6, IL-8, MCP-1 and TNF-α in U937 monocytes (Jain et al., 2009).
- Inhibited secretion of TNF-α and IL-6 in vitro (Tham et al., 2010).
- Inhibited the release of TNF- α and IL-6 in LPS-stimulated RAW 264.7 macrophages (Zhao et al., 2010).
- Inhibited IκB phosphorylation, NF-κB activation and TNF-α production induced by LPS in mouse macrophages (Nishida et al., 2010).
- Decreased LPS-induced TNF-α and IL-1β expression at both transcriptional and protein level in microglial cells (Zhang et al., 2010a).

Action of TNF-a

- Inhibited TNF-induced NF-κB activation in human myeloid cells (Singh and Aggarwal, 1995).
- Reduced TNF-induced endothelial tissue factor by inhibiting AP-1 and NF-kB in endothelial cells (Bierhaus et al., 1997).
- Blocked the activation of AP-1 and NF-κB induced by IL-1α and TNF-α in stromal cells (Xu et al., 1997).
- Inhibited TNF-α-induced expression of ICAM-1, VCAM-1 and E-selectin in HUVEC (Gupta and Ghosh, 1999; Kumar et al., 1998).
- Suppressed TNF-α-induced VEGF secretion in U937 and Raji cells. Reduced the expression of VEGF165 and VEGF121 mRNA induced by TNF-α (Chen et al., 2005).
- Inhibited TNF-mediated constitutive NF-κB activation linked to proliferation of mantle cell lymphoma cells (Shishodia et al., 2005).
- Blocked TNF-α-induced endothelial dysfunction in HUVEC (Nan et al., 2005).
- Down-regulated TNF-induced expression of cell proliferation and anti-apoptotic and metastatic gene products (Aggarwal et al., 2006).
- Inhibited TNF-\alpha-stimulated Gb3 synthase (GalT6) mRNA expression in intestinal epithelial cells (Moon et al., 2006).
- Inhibited TNF-α-induced expression of IL-1β, IL-6, TNF-α and cyclin E, but not IL-8, in HaCaT cells (Cho et al., 2007).
- Inhibited TNF-α-induced NF-κB activation in MCF-7 cells by inhibiting the proteasomal activities (Yoon and Liu, 2007).
- Suppressed TNF-α-induced expression of ICAM-1 and VCAM-1, and secretion of IL-6, IL-8 and MCP-1 in HUVEC (Kim et al., 2007).
- Inhibited TNF- α -induced NF- κ B activation in chronic myeloid leukaemia cells through modulation of redox status of the cells (Sandur et al., 2007).
- Down-regulated the expression of 29 out of 84 TNF-α-activated NF-κB-associated genes in leukaemia cells (Reuter et al., 2009).
- Inhibited TNF-induced NF-kB activation in leukaemia cells (Yadav et al., 2010).
- Inhibited TNF-α-induced cell migration, intracellular ROS generation, MMP-9 expression, MMP-9 activity and NF-κB in human aortic smooth muscle cells (Yu and Lin, 2010).
- Attenuated TNF-α-induced enhancement of TRPC1 expression, and COX-2-dependent PGE₂ production in colonic myofibroblasts (Hai et al., 2011).
- Inhibited NF-kB-mediated inflammation in human tenocytes through suppression of the PI3K/Akt pathway (Buhrmann et al., 2011).

AKT, AKT8 virus oncogene cellular homologue; AP-1, activator protein-1; HO-1, haeme oxygenase-1; ICAM-1, intercellular adhesion molecule-1; MCP-1, monocyte chemotactic protein-1; MIP-1α, macrophage inflammatory protein-1α; MMP-9, matrix metallopeptidase-9; NQO1, NADH quinone oxidoreductase 1; TRPC1, transient receptor potential channel 1; VCAM-1, vascular cell adhesion molecule-1.

cells is down-regulated by curcumin (Kumar *et al.*, 1998). Since then, a wide variety of cell signalling pathways activated by TNF have been shown to be down-regulated by curcumin; these include JNK, MAPK, PI3K/Akt.

In addition, curcumin has also been shown to modulate TNF- α function by directly binding to the ligand (Gupta *et al.*, 2011). Wua *et al.* (2010) performed molecular docking

studies with TNF- α and curcumin to predict and analyse the ability of curcumin to inhibit TNF- α by binding to it. The protein-ligand interactions were analysed by simulating the docking of the curcumin using Autodock 4.0. They identified three main binding regions for curcumin and found that curcumin is a potent inhibitor of TNF- α . They also observed that curcumin docked at the receptor-binding sites of TNF- α .



Covalent π - π aromatic interactions or π -cation interactions were found between curcumin and TNF- α . The authors predicted that curcumin is a strong inhibitor of TNF- α because of the covalent bonds it forms with Cys¹²⁹ in TNF- α . In contrast to its interaction with TNF- α , it is unclear whether curcumin can interact or affect the expression of TNF- β or lymphotoxin.

Suppression of TNF by curcumin in vivo

The anti-inflammatory effect of curcumin was first demonstrated in acute and chronic models of inflammation in rats and mice (Srimal and Dhawan, 1973). The authors showed that curcumin (50-200 mg·kg⁻¹) suppressed carrageenaninduced oedema in mice. Furthermore, curcumin was found to be as potent as phenylbutazone and exhibited minimal ulcerogenic activity. No mortality in mice was noted at doses as high as 2 g·kg⁻¹ bodyweight (Srimal and Dhawan, 1973). In the same study, the authors showed that curcumin suppresses formaldehyde-induced arthritis in rats at a dose of 40 mg·kg⁻¹ and inhibited granuloma formation at 80-160 mg·kg⁻¹. However, the mechanism by which curcumin mediates these anti-inflammatory effects in animals was not revealed until several years later, when our group showed that curcumin can suppress TNF-induced NF-κB activation (Singh and Aggarwal, 1995) and other groups showed that curcumin blocked TNF production in cell culture (Chan, 1995) and the expression of pro-inflammatory genes (Jobin et al., 1999). Since then, numerous mechanisms by which curcumin can exhibit antiinflammatory activity have been proposed (Figures 1 and 2).

Numerous reports have been published suggesting that oral administration of curcumin down-regulates $TNF-\alpha$

expression both in the serum and in the tissue of animals (Nanji et al., 2003; Yao et al., 2004; Sharma et al., 2007a; Billerey-Larmonier et al., 2008; Larmonier et al., 2008; Ung et al., 2010; El-Moselhy et al., 2011; Gutierres et al., 2012) (Table 2). Attenuation of TNF- α levels by curcumin has been noted in mice (Leyon and Kuttan, 2003), rats (Siddiqui et al., 2006) and rabbits (Yao et al., 2004; Huang et al., 2008). A dose of curcumin of 50-500 mg·kg⁻¹·day⁻¹ was used for most of these studies. Endotoxin has been shown to induce septic shock in animals, in part, through the production of TNF, and this condition has been shown to be reversed by curcumin (Siddiqui et al., 2006; Chen et al., 2007; 2008; Huang et al., 2008; Nishida et al., 2010). Decreased TNF- α levels have also been noted in tumour-bearing animals treated with this polyphenol (Leyon and Kuttan, 2003). In addition to cancer, down-regulation of TNF-α by curcumin has been associated with protection from various pro-inflammatory diseases, including sub-chronic inflammation (Nandal et al., 2009; Nishida et al., 2010), cardiovascular diseases (Yao et al., 2004; 2005; Mito et al., 2011; Avci et al., 2012), diabetes (Jain et al., 2009; El-Azab et al., 2011; El-Moselhy et al., 2011), acute pancreatitis (Gulcubuk et al., 2006), enterocolitis (Jia et al., 2010), enteritis (Song et al., 2010), prostatitis (Zhang et al., 2010b), diabetic neuropathy (Sharma et al., 2007b), hepatic injury (Yun et al., 2010), Th1-type ileitis (Bereswill et al., 2010), hepatic fibrosis (Shu et al., 2007; Zeng et al., 2011), radiationinduced lung fibrosis (Lee et al., 2010), asthma (Ammar el et al., 2011), alcohol-induced liver disease (Nanji et al., 2003), non-alcoholic steatohepatitis (Ramirez-Tortosa et al., 2009), concanavalin A-induced liver injury (Tu et al., 2012), renal injury (Hashem et al., 2008; Pan et al., 2012), infection (Allam, 2009), fatigue (Gupta et al., 2009), bone turnover (Yang et al., 2011) and high-fat diet-induced hyperglycaemia (El-Moselhy et al., 2011). Curcumin has also been found to

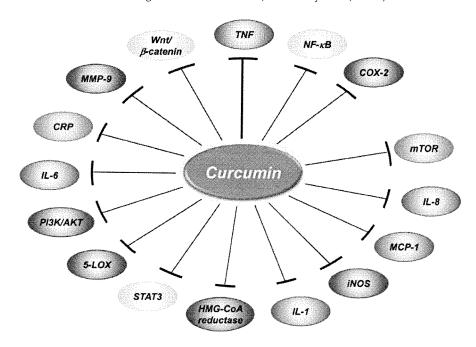


Figure 2 Inflammatory targets modulated by curcumin.



Table 2

Curcumin inhibits TNF production in animals

- Prevented alcohol-induced liver disease in rats by inhibiting the expression of NF-κB-dependent genes including TNF-α (Nanji *et al.*, 2003)
- Reduced the serum level of TNF-α and NO in B16F-10 melanoma cells bearing C57BL/6 mice (Leyon and Kuttan, 2003).
- Suppressed the myocardial TNF- α and MMP-2 expression and improved left ventricular function in pressure overloaded rabbits (Yao et al., 2004).
- Decreased the elevations in plasma IL-8, IL-10 and TNF-α in rabbits after cardiopulmonary bypass and cardiac global ischaemia (Yeh et al., 2005).
- Significantly lowered the serum TNF-α and IL-6 levels in rat model of acute pancreatitis (Gulcubuk et al., 2006).
- Decreased the expression of TNF-α and reduced the mortality in rat model of sepsis (Siddiqui et al., 2006).
- Reduced the mortality rate of LPS-infused rats by decreasing the circulating TNF-α levels and the consumption of peripheral platelets and plasma fibrinogen (Chen *et al.*, 2007).
- Significantly inhibited TNF-α and NO levels in rat model of diabetic neuropathy (Sharma et al., 2007b).
- Down-regulated the expressions of TNF-α and IL-8 in the copper-overloaded rats (Wan et al., 2007).
- Decreased the levels of NO, TGF-β1 and TNF-α in rat model of hepatic fibrosis (Shu et al., 2007).
- Inhibited expression of TNF-α and IL-1β stimulated by LPS in murine macrophages through inhibition of NF-κB pathway (Chen *et al.*, 2008).
- Significantly reduced the LPS-induced overproduction of circulating TNF-α, IL-1β and IL-6, brain glutamate, PGE₂, and hydroxyl radicals in rabbit (Huang *et al.*, 2008).
- Significantly decreased TNF-α mRNA and caspase-8 that probably contributes to the protective role of the turmeric-based diet against renal injury in rat (Hashem *et al.*, 2008).
- Reduced TNF-α levels in a rabbit model of non-alcoholic steatohepatitis (Ramirez-Tortosa et al., 2009).
- Decreased the levels of TNF-α in a rat model of subchronic inflammation (Nandal et al., 2009).
- Exhibited anti-fibrosis activity by decreasing the levels of TNF-α and TGF-β1 in serum and lung tissue of SiO₂-induced fibrosis mice model (Jiang *et al.*, 2009).
- Prevented the injurious effects of DSS and ameliorated release of TNF-α and NO in a rat model (Arafa et al., 2009).
- Decreased serum levels of IL-12 and TNF-α in mice infected with Schistosoma mansoni cercariae (Allam, 2009).
- Significantly attenuated oxidative stress and TNF-α levels in a mouse model of immunologically induced fatigue (Gupta et al., 2009).
- Significantly decreased the blood levels of IL-6, MCP-1, TNF-α, glucose, HbA₁ and oxidative stress in streptozotocin-induced diabetic rat model (Jain et al., 2009).
- Decreased LPS-induced TNF-α production in lungs of mice. At 5% concentration, curcumin significantly improved survival of mice and decreased radiation-induced lung fibrosis (Lee *et al.*, 2010).
- Exhibited protective effects against necrotizing enterocolitis in neonatal rats, possibly by inhibiting COX-2, reducing TNF-α and increasing IL-10 contents (Jia et al., 2010).
- Significantly decreased the levels of TNF-α and IL-8 in the serum and prostate tissues in a rat model of prostatitis (Zhang et al., 2010b).
- Significantly decreased the production of TNF- α in a mouse model of acute inflammation (Bansal and Chhibber, 2010).
- Protected mice from LPS/GalN-induced hepatic injury and inflammation by blocking TNF-α production (Yun et al., 2010).
- Increased IFN-γ, IL-12 and IL-13 levels, but decreased TNF-α level in rats intoxicated with 2,3,7,8-tetrachlorodibenzo-p-dioxin (Ciftci et al., 2010).
- Lowered the production of IL-23p19, IFN-γ, TNF-α, IL-6 and MCP-1 in a murine model of hyperacute Th1-type lleitis (Bereswill *et al.*, 2010).
- Suppressed LPS stimulated TNF-α production in mice (Nishida et al., 2010).
- Reduced the aluminum-induced inflammatory response as indicated by down-regulation of NF-κB and TNF-α in glial cells (Sood et al., 2011).
- Improved the lipid metabolism and delayed the progression of hepatic fibrosis in rats with experimental steatohepatitis through suppression of TNF-α, NF-κB and HMG-CoA reductase (Zeng et al., 2011).
- Inhibited mRNA expression of TNF-α in a murine model of asthma (Ammar el et al., 2011).
- Suppressed inflammation by reducing levels of TNF-α, NF-κB and IL-6 in CCl₄-treated rats (Bassiouny et al., 2011).
- Reduced cardiac inflammation through suppression of IL-1β, TNF-α, GATA-4 and NF-κB in a rat model of experimental autoimmune myocarditis (Mito et al., 2011).
- Suppressed serum levels of TNF-α and IL-1β in a streptozotocin-induced diabetic mouse model (El-Azab et al., 2011).
- Attenuated TNF-α levels and exhibited anti-hyperglycaemic effect and improved insulin sensitivity in high-fat diet-fed rats (El-Moselhy et al., 2011).
- Prevented deterioration of the bone structure and produced beneficial effects in bone turnover in transgenic mice possibly through modulation of TNF-α and IL-6 (Yang *et al.*, 2011).
- Protected against ischaemia/reperfusion injury in rat skeletal muscle through inhibition of plasma TNF-α levels (Avci et al., 2012).
- Inhibited the high glucose-induced plasma TNF-α production and macrophage infiltration and prevented renal injury in diabetic rats (Pan et al., 2012).
- Attenuated concanavalin A-induced liver injury in mice by inhibition of TNF expression through TLR-2, TLR-4 and TLR-9 expression (Tu et al., 2012).

CCl₄, carbon tetrachloride; DSS, dextran sulfate sodium; GalN, D-galactosamine; HbA1, haemoglobin α 1; HMG-CoA, 3-hydroxy-3-methylglutaryl-coenzyme A; MCP-1, monocyte chemotactic protein-1; SiO₂, silicon dioxide.

down-regulate NF-κB-regulated gene products such as inducible NOS (iNOS), IL-1, IL-6, IL-8, MCP-1, MMP-2 and MMP-9 in animals (Yao *et al.*, 2004; Yeh *et al.*, 2005; Jain *et al.*, 2009). Curcumin also protects against the toxic effects of copper overload (Wan *et al.*, 2007), dextran sulfate sodium (Arafa *et al.*, 2009), *p*-dioxin (Ciftci *et al.*, 2010) and aluminum (Sood *et al.*, 2011) through down-modulation of TNF- α .

A reduction in the production of iNOS mRNA was observed when BALB/c mouse peritoneal macrophages cultured ex vivo were treated with 1-20 µM curcumin (Chan et al., 1998); in vivo, two oral treatments of 0.5 mL of a 10 μ M solution of curcumin (92 ng·g⁻¹ bodyweight) reduced iNOS mRNA expression in the livers of LPS-injected mice by 50-70% (Chan et al., 1998). This suggests that curcumin is potent at nmol g-1 bodyweight. This efficacy was associated with two modifications of the schedule of dosing: firstly, an aqueous solution of curcumin was prepared by initially dissolving the compound in 0.5 N NaOH and then diluting it immediately with PBS; secondly, mice were fed curcumin at dusk after fasting. Inhibition of iNOS mRNA expression was not observed in the mice that were fed ad libitum, suggesting that food intake may interfere with the absorption of curcumin.

Oral bioavailability and safety of curcumin

Curcumin usually manifests its biological response when given orally to mice at about 50-500 mg·kg⁻¹ bodyweight (Farombi and Ekor, 2006). These doses, however, are too low to detect significant levels of curcumin in the serum. The reason for this discrepancy is not clear; however, there are several possible explanations. Firstly, curcumin is known to bind to numerous proteins present in the serum including albumin (Gupta et al., 2011; Kim et al., 2012). Secondly, curcumin is rapidly transported across the cells and tissues (Anand et al., 2007). Thirdly, tetrahydrocurcumin, a metabolite of curcumin, was found to be more active than curcumin for treating chloroquine-induced hepatotoxicity in rats (Pari and Amali, 2005). Fourthly, in another study it was shown that when curcumin was dissolved in 0.1 N NaOH it manifested its effects in animals in the μg·kg-1 range (Chan et al., 1998).

In one recent study the potential of a novel solid lipid curcumin particle (SLCP) preparation to produce adverse effects in rats after acute and sub-chronic administration was investigated (Dadhaniya et al., 2011). The oral LD50 of the preparation in rats as well as in mice was found to be greater than 2000 mg·kg⁻¹ bodyweight. In the sub-chronic toxicity study, 180, 360 and 720 mg·kg⁻¹ bodyweight day⁻¹ of SLCP preparation was administered via oral gavage to Wistar rats (10 per sex per group) for 90 days. Administration of the curcumin preparation did not result in any toxicologically significant treatment-related changes in clinical (including behavioural) observations, ophthalmic examinations, bodyweights, bodyweight gains, food consumption and organ weights. No adverse effects of the curcumin preparation were noted on the haematology, serum chemistry parameters and urinalysis. Terminal necropsy did not reveal any treatmentrelated gross or histopathology findings. On the basis of these study results, the no observed-adverse-effect level for this standardized novel curcumin preparation was determined as 720 mg·kg⁻¹ bodyweight day⁻¹.

In humans, as little as 150 mg of curcumin has been shown to be effective in reducing serum levels of proinflammatory cytokines (Usharani et al., 2008) (Table 3). The effect of curcumin administration, 500 mg of curcumin day-1 for 7 days, on serum levels of cholesterol and lipid peroxides was studied in 10 healthy human volunteers (Soni and Kuttan, 1992). A significant decrease in the level of serum lipid peroxides was noted, along with an increase in highdensity lipoprotein cholesterol and a decrease in total serum cholesterol. In another study, curcumin was given orally at up to 8000 mg·day-1 to 25 patients (Cheng et al., 2001). The serum concentration of curcumin usually peaked at 1-2 h after oral intake of curcumin and gradually declined within 12 h. The average peak serum concentrations after oral intake of 4000, 6000 and 8000 mg of curcumin were 0.51 \pm 0.11, 0.63 ± 0.06 and $1.77 \pm 1.87 \,\mu\text{M}$ respectively. , However, urinary excretion of curcumin was undetectable.

Vareed et al. (2008) examined the pharmacokinetics of a curcumin preparation in healthy human volunteers at 0.25-72 h after a single oral dose. Curcumin was administered at doses of 10 g (n = 6 subjects) and 12 g (n = 6 subjects). Using HPLC with a limit of detection of 50 ng·mL⁻¹, only one subject had detectable free curcumin at any of the 14 time points assayed, but curcumin glucuronides and sulfates were detected in all subjects. Based on the pharmacokinetic model, the area under the curve for the 10 and 12 g doses was 35.33 \pm 3.78 and 26.57 \pm 2.97 μ g·mL⁻¹ \times h, respectively, whereas $C_{\rm max}$ was 2.30 \pm 0.26 and 1.73 \pm 0.19 $\mu g \cdot {\rm mL^{-1}}$. The $T_{\rm max}$ and $t_{1/2}$ were estimated to be 3.29 \pm 0.43 and 6.77 \pm 0.83 h. The ratio of glucuronide to sulfate was 1.92:1. The curcumin conjugates were present as either glucuronide or sulfate, not as mixed conjugates. The group concluded that curcumin is absorbed after oral dosing in humans and can be detected as glucuronide and sulfate conjugates in plasma. Another study evaluated the efficacy of oral curcumin (4 g·day-1) in 26 patients with monoclonal gammopathy of undefined significance (Golombick et al., 2009). They found that oral curcumin was bioavailable as it decreased paraprotein load. Thus, all of these studies clearly demonstrate that although serum levels of curcumin administered orally are very low, it can still manifest its effect in vivo.

Table 3 Curcumin inhibits TNF production in humans

- Improved endothelial function and reduced levels of malondialdehyde, IL-6, TNF-α and endothelin-1 in diabetic patients (Usharani et al., 2008).
- Had non-significant effects on the production of IL-8, IL-1β, TNF-α and COX-2 in gastric mucosa from Helicobacter pylori-infected gastritis patients (Koosirirat et al., 2010).
- Improved bodyweight, reduced serum TNF-α and induced p53 expression in patients with colorectal cancer (He et al., 2011)



Suppression of TNF- α by curcumin in patients

At least two studies have suggested that orally administered curcumin can down-modulate the expression of TNF- α in patients (Usharani *et al.*, 2008; He *et al.*, 2011) (Table 3). In addition, several other pro-inflammatory biomarkers are decreased by curcumin in human subjects (Hanai and Sugimoto, 2009; Khajehdehi *et al.*, 2011; Koosirirat *et al.*, 2010). In most of these studies, 150–500 mg of curcumin was sufficient to manifest a response.

The interest in curcumin research in human participants has increased markedly over the years (Table 4). To date, over 60 clinical trials have evaluated the safety and efficacy of this polyphenol in humans, whereas another 35 clinical trials are further evaluating its efficacy. Curcumin was found to be effective in TNF-associated human diseases such as cancer, cardiovascular diseases, metabolic diseases, neurological diseases, skin diseases, RA, CD and psoriasis. However, whether curcumin exerts its effects through modulation of TNF in these patients is, at present, unclear. Given the fact that most of the currently available TNF blockers produce adverse effects in patients and are very expensive, this orally bioavailable polyphenol represents an important therapeutic for TNF-associated diseases.

In addition to its efficacy in TNF-associated human diseases, curcumin has been found to be effective in a number of other human diseases. Readers interested in such studies should refer to one of the recent reviews published from this laboratory (Gupta *et al.*, 2013).

Role of curcumin in TNF-related diseases

Rheumatoid arthritis

Numerous reports have suggested that TNF plays a major role in RA. Thus, TNF blockers have been found to be beneficial for patients with RA. Therefore, curcumin has been tested as a treatment for RA. One of the earliest indications of its potential efficacy was obtained in 1973, when curcumin was found to suppress formaldehyde-induced arthritis in rats at a dose of 40 mg·kg⁻¹ and inhibit granuloma formation at 80-160 mg·kg⁻¹ (Srimal and Dhawan, 1973). Later, Joe et al. (1997) showed that curcumin can lower the elevated serum acidic glycoprotein levels present in adjuvant-induced arthritis. Also, oral administration of curcumin has been shown to prevent streptococcal cell wall-induced arthritis in mice (Funk et al., 2006) and to suppress MMP-1 and MMP-3 production and attenuate the inflammatory response in a collagen-induced arthritis model in mice (Moon et al., 2010). In addition, curcumin has been found to down-regulate the expression of TNF-α and IL-1β in ankle joints and decrease NF-κB activity, PGE2 production, COX-2 expression and MMP secretion in synoviocytes. Furthermore, curcumin has been shown to have a synergistic effect with methotrexate in decreasing adjuvant-induced arthritis in mice and in minimizing liver damage (Banji et al., 2011).

Other *in vitro* findings indicate that the protective effects of curcumin against RA are mediated through inhibition of

neutrophil activation, suppression of synoviocyte proliferation and inhibition of angiogenesis as suggested by curcumin's ability to inhibit collagenase and stromelysin in chondrocytes (Jackson *et al.*, 2006). Further, the suppression of NF-κB by curcumin has been found to be associated with its inhibition of the expression of COX-2, NO, PGE₂, IL-1β, IL-6, IL-8, MMP-3 and MMP-9 in human chondrocytes (Shakibaei *et al.*, 2007; Mathy-Hartert *et al.*, 2009). Curcumin has also been found to suppress IL-8 expression in human synovial fibroblasts (Tong *et al.*, 2008).

One of the earliest studies demonstrating that curcumin has anti-rheumatic activity in humans appeared almost three decades ago (Deodhar et al., 1980). In a more recent, the efficacy of a proprietary complex of curcumin with soy phosphatidylcholine (Meriva®; Throne Research Inc., Dover, ID, USA) was investigated in 50 patients with osteoarthritis (OA) at a dosage of 200 mg of curcumin day-1 (Belcaro et al., 2010b). OA symptoms were evaluated by the use of Western Ontario and McMaster Universities Osteoarthritis Index (WOMAC) scores. After 3 months of treatment with this complex, the global WOMAC score was found to be decreased by 58% (P < 0.05), walking distance in the treadmill test was prolonged from 76 to 332 m (P < 0.05) and C-reactive protein levels decreased from 168 \pm 18 to 11.3 \pm 4.1 mg·L⁻¹ in the subpopulation with high C-reactive protein levels. In comparison, the control group experienced only a modest improvement in these parameters. These results show that curcumin is clinically effective in the management and treatment of OA. In another study, the same investigator examined the efficacy and safety of Meriva in 100 patients with OA after long-term administration (8 months) (Belcaro et al., 2010a). The clinical end points were WOMAC score, Karnofsky performance scale index score and treadmill walking performance, and were complemented by the evaluation of a series of inflammatory markers including IL-1β, IL-6, sCD40L, soluble vascular cell adhesion molecule-1 and erythrocyte sedimentation rate. Significant improvements in both the clinical and the biochemical end points were observed for the Meriva group compared with the control group.

In another randomized pilot study, the efficacy of curcumin alone and in combination with diclofenac sodium was assessed in patients with active RA (Chandran and Goel, 2012). Forty-five patients diagnosed as having RA were randomized into three groups: patients receiving curcumin alone (500 mg), those receiving diclofenac sodium alone (50 mg) and those receiving combinations of curcumin and diclofenac sodium. The primary end points were reduction in Disease Activity Score (DAS) 28. The secondary end points included American College of Rheumatology (ACR) criteria for reduction in tenderness and swelling of joint scores. Patients in the curcumin group showed the highest percentage of improvement in overall DAS and ACR scores, which were significantly better than those of patients in the diclofenac sodium group. To our knowledge, this is the first evidence showing the potential of curcumin as a therapeutic for patients with active RA.

Inflammatory bowel disease

Inflammatory bowel disease (IBD) consists of two separate diseases, CD and ulcerative colitis (UC); both characterized by chronic recurrent ulceration of the bowel (Kozuch and

Xerox WorkCentre **SMTP Transfer Report**



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Submission Date:

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Images Scanned:

Attachment Name: Format:

Image-Only PDF Encrypted E-mail:

Message Settings:

Subject: From: Reply To: To:

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 Table 4

 Chronological listing of curcumin studies in human participants

Year	Disease	Reference
1937	Cholecystitis	Oppenheimer, 1937
1972	Diabetes	Srinivasan, 1972
1980	Rheumatoid arthritis	Deodhar <i>et al.</i> , 1980
1986	Post-operative inflammation	Satoskar <i>et al.</i> , 1986
1987	Cancer lesions	Kuttan <i>et al.</i> , 1987
1992	Lung cancer	Polasa et al., 1992
1992	Atherosclerosis	Soni and Kuttan, 1992
1993	Gastric ulcer	Kositchaiwat et al., 1993
1996	Acquired immunodeficiency syndrome	James, 1996
1997	Cancer lesions	
1999	Biliary dyskinesia	Hastak et al., 1997
	Gallbladder contraction	Niederau and Gopfert, 1999
	Chronic anterior uveitis	Rasyld and Lelo, 1999
.000	Idiopathic orbital inflammatory pseudotumour	Lal et al., 1999
	Psoriasis	Lal et al., 2000
001	Cancer lesions	Heng et al., 2000
	Colorectal cancer	Cheng <i>et al.</i> , 2001
		Sharma <i>et al.,</i> 2001
004	Peptic ulcer	Prucksunand et al., 2001
004	Colorectal cancer	Sharma <i>et al.</i> , 2004
005	Irritable bowel syndrome	Bundy <i>et al.</i> , 2004
003	Colorectal cancer	Garcea et al., 2005
	Pancreatic cancer	Durgaprasad et al., 2005
	Crohn's disease	Holt <i>et al.</i> , 2005
	Ulcerative proctitis	Holt <i>et al.</i> , 2005
	Alzheimer's disease	Ringman et al., 2005
	Renal transplantation	Shoskes <i>et al.</i> , 2005
006	Colorectal cancer	Cruz-Correa et al., 2006
	Ulcerative colitis	Hanai <i>et al.,</i> 2006
00 <i>7</i>	Cancer lesions	Chainani-Wu et al., 2007
	Multiple myeloma	Vadhan-Raj <i>et al.,</i> 2007
	Helicobacter pylori infection	Di Mario et al., 2007
800	Pancreatic cancer	Dhillon et al., 2008
	Psoriasis	Kurd <i>et al.</i> , 2008
	Alzheimer disease	Baum <i>et al.</i> , 2008
	Acute coronary syndrome	Alwi et al., 2008
	Diabetes	Usharani et al., 2008
	Hepatoprotection	Adhvaryu <i>et al.</i> , 2008
009	Multiple myeloma	Golombick et al., 2009
	Irritable bowel syndrome	Shimouchi et al., 2009
	DejerineSottas disease	Burns <i>et al.</i> , 2009
	Recurrent respiratory tract infections	
	Chronic bacterial prostatitis	Zuccotti et al., 2009
110	Cancer lesions	Cai <i>et al.</i> , 2009
	Pancreatic cancer	Rai et al., 2010
	Breast cancer	Epelbaum <i>et al.,</i> 2010
	Prostate cancer	Bayet-Robert et al., 2010
	Inflammatory bowel disease	lde et al., 2010
	Recurrent anterior uveitis	Epstein et al., 2010
		Allegri <i>et al.,</i> 2010
	Helicobacter pylori infection	Koosirirat <i>et al.</i> , 2010
	Osteoarthritis	Belcaro et al., 2010a; Belcaro et al., 2010
	Diabetes	Wickenberg et al., 2010
	Vitiligo	Asawanonda and Klahan, 2010
	β-Thalassaemia	Kalpravidh et al., 2010
	Chronic arsenic exposure	Biswas <i>et al.</i> , 2010
	Osteosarcoma 	Gota <i>et al.</i> , 2010
11	Colorectal cancer	Carroll et al., 2011; He et al., 2011
	Pancreatic cancer	Kanai <i>et al.,</i> 2011
	Head and neck cancer	Kim <i>et al.,</i> 2011
	Ulcerative colitis	Lahiff and Moss, 2011
	Diabetic nephropathy	Khajehdehi <i>et al.</i> , 2011
	Diabetic microangiopathy	Appendino et al., 2011
	Alcohol intoxication	Sasaki <i>et al.</i> , 2011
12	Rheumatoid arthritis	
	Diabetes	Chandran and Goel, 2012
	Lupus nephritis	Chuengsamarn et al., 2012
		Khajehdehi et al., 2012



Hanauer, 2008). It is likely that the pathogenesis of these diseases involves genetic, environmental and immunological factors (Hanauer, 1996). The expression of several cytokines, including TNF-α, IL-1β, IL-6, IL-8 and chemokines, all regulated by NF-kB, is increased in IBD (Ferretti et al., 1994; Jijon et al., 2000; Yamamoto et al., 2000; Jobin, 2008). All of these critical proteins are up-regulated by NF-kB and suppression of NF-κB by anti-sense can attenuate experimental colitis in mice (Neurath et al., 1996). Among the various gut immune factors, TNF-α is a major pro-inflammatory cytokine in IBD (Brown and Mayer, 2007; Louis, 2001). Mucosal levels of TNF- α are elevated in patients with IBD (Murch et al., 1991; Braegger et al., 1992), and its inhibition (Papadakis and Targan, 2000) or neutralization can improve both UC (Jarnerot, 1989) and CD (Ardizzone and Bianchi Porro, 2005). Conventional therapies for UC include sulfasalazine, 5-aminosalicylic acid, salazosulfapyridine, azathioprine, mercaptopurines, cyclosporine, corticosteroids and TNF blockers (Kozuch and Hanauer, 2008; Ng and Kamm, 2009). All of these treatments have significant toxic side effects and are partly or completely ineffective in a significant number of

Now there are numerous lines of evidence to suggest that curcumin has enormous potential against both CD and UC (Table 5). Firstly, epidemiological studies indicate that turmeric (which contains 2-8% curcuminoids) may contribute to the lower incidence of cancer, especially large-bowel cancers in Indians (Mohandas and Desai, 1999; Sinha et al., 2003). Secondly, curcumin, when administered orally in the diet, prevented trinitrobenzene sulfonic acid (TNBS)- or dinitrobenzene sulfonic acid (DNBS)-induced colitis in mice (Sugimoto et al., 2002; Salh et al., 2003; Venkataranganna et al., 2007; Ung et al., 2010). Thirdly, curcumin improves both the wasting and histopathological signs of colonic inflammation. Fourthly, curcumin inhibits CD4+ T-cell infiltration and NF-κB activation in colonic mucosa. Fifthly, curcumin manifests its effects against colitis by suppressing the expression of inflammatory cytokines such as TNF- α (Camacho-Barquero et al., 2007; Mouzaoui et al., 2012), IFN-γ (Ung et al., 2010), IL-17 (Ung et al., 2010) and enzymes, such as p38 MAPK, iNOS, COX-2, myeloperoxidase and MMP-9, in the colonic mucosa (Camacho-Barquero et al., 2007). Sixthly, curcumin has a therapeutic effect on DNBS-induced colitis in mice induced by its agonistic action on the vanilloid receptor TRPV1 (Martelli et al., 2007). Seventhly, curcumin suppresses colonic inflammation induced by deletion of the mdr1 gene in mice (Nones et al., 2009). The effects of curcumin in TNBS-induced colitis in mice were found to be strain-dependent: BALB/c mice were protected, whereas SJL/J mice were not protected (Billerey-Larmonier et al., 2008). The effect of curcumin against colitis was also limited in Th1-driven colitis in IL-10-deficient mice (Larmonier et al., 2008; Ung et al., 2010). Curcumin failed to inhibit NF-κB in these mice, but when combined with IL-10, curcumin inhibited NF-κB quite effectively. Eighthly, curcumin decreases TNF-α-induced oxidative stress and colitis in mice (Mouzaoui et al., 2012). Ninthly, curcumin combined with resveratrol and simvastatin decreases acute small intestinal inflammation in mice by down-regulating the Th1-type immune response (Bereswill et al., 2010). Tenthly, curcumin suppresses TNBS-induced

colonic inflammation in mice by down-regulation of NFκB, TLR4 and MyD88 (Lubbad et al., 2009a). Eleventhly, curcumin has been shown to inhibit colitis by inducing the production of tolerogenic dendritic cells that promote differentiation of T-cells into Treg, which include CD4⁺CD25⁺Foxp3⁺Treg and IL-10-producing Tr1 cells, and by producing TGF-β (Cong et al., 2009). Twelfthly, curcumin can reverse TNF-α-mediated reduction in Phex protein in mice, which is responsible for the inhibition of osteoblast mineralization linked to the abnormal bone metabolism associated with IBD (Uno et al., 2006). For this, the authors examined calvaria of 6- to 7-week-old mice given TNBS with or without neutralizing TNF-α antibody, dietary curcumin or systemically with recombinant TNF-α. They found that compared to control animals, Phex mRNA expression decreased by 40-50% in both TNBS colitis and TNF-αinjected mice. Dietary curcumin and TNF-α antibody counteracted these detrimental effects of TNBS on Phex gene expression.

Thus, the above findings in animals clearly indicate that orally administered curcumin has the potential to protect from the development of IBD. Additional evidence suggests its potential in humans. Firstly, as little as 150 mg of curcumin twice daily can suppress the levels of expression of inflammatory cytokines TNF-α and IL-6 in serum (Usharani et al., 2008; He et al., 2011). Secondly, a dose of 2 g-day-1 of curcumin can suppress NF-κB activation in human peripheral blood mononuclear cells (Vadhan-Raj et al., 2007). Thirdly, curcumin was found to suppress p38 MAPK, reduce IL-1ß and MMP-3, and enhance IL-10 in mucosa of children and adults with IBD (Epstein et al., 2010). Fourthly, more than 65 different trials have been conducted with orally administered curcumin in humans. Fifthly, promising results were obtained from a small open-label study examining the use of curcumin to treat IBD (Holt et al., 2005). A pure curcumin preparation was administered to five patients with ulcerative proctitis and to five patients with CD. All patients with proctitis improved, and four had their concomitant medications reduced; four of the five CD patients had lowered CDAI scores and sedimentation rates. Sixthly, in a randomized multicentre doubleblind, placebo-controlled trial, curcumin was examined as a maintenance therapy for UC and found to produce favourable effects (Hanai et al., 2006). Of 89 patients with UC, 45 received 1 g of curcumin after breakfast and 1 g after their evening meal, plus sulfasalazine and mesalamine for 6 months. Of the 43 patients who received curcumin, 2 (4.65%) experienced relapse during the 6 months of therapy, whereas 8 (20.51%) of the 39 patients in the placebo group experienced relapse. Recurrence rates in the curcumin-treated and placebo groups were significantly different. Furthermore, curcumin use resulted in an improvement in both the clinical activity index and the endoscopic index and suppressed UC-associated morbidity. Thus, curcumin appears to be a promising and safe medication for maintaining remission in patients with quiescent UC. These two small studies have shown promising results for IBD. Seventhly, orally administered curcumin was found to have a therapeutic effect against colorectal cancer (He et al., 2011). In an open-label study in which 126 patients were treated with 360 mg, p.o., curcumin three times a day, curcumin's effects were noted within 10-30 days.

Table 5

Effect of curcumin on models of inflammatory bowel disease

- Prevented TNBS-induced colitis in mice; inhibited CD4+ T-cell infiltration and NF-κB activation, and expression of TNF-α, IFN-γ, IL-6 and IL-12 in colonic mucosa (Sugimoto et al., 2002).
- Inhibited DNB-induced colitis in mice, prevented tissue damage, reduced MPO and IL-1β expression and inhibited NF-κB activation in the mucosal tissue (Salh *et al.*, 2003).
- Inhibited mucosal injury in TNBS-induced colitis in mice, reduced NO and ROS levels, inhibited neutrophil infiltration and inactivated NF-κB in colonic mucosa (Ukil et al., 2003).
- Inhibited TNBS-induced colitis in rats, inhibited IL-1 expression, increased IL-10 expression in colonic mucosa and decreased NF-κB activation (Jian et al., 2004).
- Attenuated TNBS-induced chronic colitis through inhibition of MPO and COX-2 in rats and improved survival (Jiang et al., 2006).
- Prevented TNBS-induced chronic colitis, decreased Th1 (IL-12, IFN-γ/TNF-α, IL-1) and increased Th2 (IL-4 and IL-10) cytokines in colon mucosa; and increased IL-4 and IFN-γ in splenocytes and circulation (Zhang et al., 2006).
- Reversed TNF-α-mediated reduction in Phex protein responsible for the inhibition of osteoblast mineralization linked abnormal bone metabolism in IBD (Uno *et al.*, 2006).
- Prevented the development of DSS-induced experimental colitis in BALB/c mice through inhibition of MPO and NF-κB (Deguchi et al., 2007).
- Protected against DNCB-induced colitis through down-regulation of MPO, ALP, LPO, NF-κB and iNOS (Venkataranganna et al., 2007).
- Prevented DNBS-induced colitis in mice by interaction with vanilloid receptor TRPV1 (Martelli et al., 2007).
- Attenuated the TNBS-induced colitis in rats through inhibition of MPO, TNF-α, COX-2, iNOS, p38 MAPK in colonic mucosa (Camacho-Barquero et al., 2007).
- Attenuated the TNBS-induced colitis in rats through inhibition of down-regulation of hepatic CYP3A2 (Masubuchi et al., 2008).
- Inhibited the TNBS-induced colitis and splenocyte proliferation in BALB/c mice but not in NKT-deficient SJL/J mice (Billerey-Larmonier et al., 2008).
- Exhibited protective effect on Th1-driven colitis in IL-10 deficient mice, with no effect on NF-κB (Larmonier et al., 2008).
- Attenuated the TNBS-induced colitis in rats through reversal of carbachol-induced contraction of the colon and modulating NF-κB
 activation (Lubbad et al., 2009b).
- Attenuated the TNBS-induced colitis in rats through suppression of expression in TLR-4, MyD88 and NF-κB proteins in inflamed tissue (Lubbad et al., 2009a).
- Prevented the DSS-induced colitis in mice through suppression of serum TNF-α levels, NO and colonic MPO expression (Arafa et al., 2009).
- Protected from IBD in mdr1a-KO mice through inhibition of TNF-α, IFN-γ, chemokine, p38, TLR2, CD14 and up-regulation of xenobiotic metabolism (Nones et al., 2009).
- Protected from IBD by inducing the tolerogenic dendritic cell that promotes differentiation of intestine-protective regulatory T-cells in vivo (Cong et al., 2009).
- Inhibited pro-inflammatory cytokine release in the IL-10-deficient mouse model of IBD (Ung et al., 2010).
- Ameliorated small intestinal inflammation by down-regulating Th1 cell-associated cytokines (IFN-γ, TNF-α, IL-6, MCP-1) (Bereswill et al., 2010).
- Protected intestinal mucosal barrier function of rat enteritis via activation of MKP-1 and attenuation of p38 and NF-κB activation (Song et al., 2010).
- Inhibited DSS-induced colitis in mice via inhibition of beta catenin translocation, and down-regulation of TNF-α and IFN-γ levels (Villegas et al., 2011).
- Attenuated TNF-α-induced oxidative stress, acute colitis and hepatotoxicity in mice (Mouzaoui et al., 2012).
- Suppressed p38, reduced IL-1β and MMP-3, and enhanced IL-10 in mucosa of children and adults with IBD (Epstein et al., 2010).

ALP, alkaline phosphatase; DNBS, dinitrobenzene sulfonic acid; DNCB, 2,4-dinitrochlorobenzene; DSS, dextran sulfate sodium; IBD, inflammatory bowel disease; LPO, lipid peroxidation; MCP-1, monocyte chemotactic protein-1; MKP-1, mitogen-activated protein kinase phosphatase-1; MPO, myeloperoxidase; MyD88, myeloid differentiation primary response gene 88; NKT, natural killer T-cells; Phex, phosphate regulating gene with homologies to endopeptidases on the X chromosome; Th, T helper cell type; TLR, toll-like receptor; TNBS, trinitrobenzene sulfonic acid.

Psoriasis

Like IBD and RA, psoriasis is a common chronic inflammatory disease of the skin and joints that affects about 2% of the general population is another indication for which TNF blockers have been approved. Depending on the stage of disease, current treatment options include UVB, UVA plus psoralen, methotrexate, acitretin, cyclosporine, infliximab, etanercept, adalimumab, efalizumab and alefacept. Whereas infliximab, etanercept and adalimumab are specific TNF blockers, all of the others are immunosuppressive agents that

could increase the risk of infections and malignancies, especially with long-term use (Greaves and Weinstein, 1995; Kurd et al., 2007). Psoriasis is a chronic disease, which requires long-term treatment and 51% of patients with psoriasis use complementary and alternative therapies (Fleischer et al., 1996). Thus, safe, affordable and effective agents are needed to treat this condition.

There are several reasons to believe that curcumin may have potential for treating psoriasis. Firstly, on irradiation with visible light, curcumin has been proven to be phototoxic for *Salmonella typhimurium* and *Escherichia coli*, even at very

low concentrations (Tonnesen et al., 1987). This observed phototoxicity makes curcumin a potential photosensitizing drug, which could be used in phototherapy of psoriasis. Secondly, when curcumin was tested as an anti-psoriatic drug in the modified mouse tail test, an animal model of psoriasis, it exhibited some activity (Bosman, 1994). Thirdly, curcumin has been shown to inhibit the proliferation of human keratinocytes through suppression of pro-inflammatory pathways (Pol et al., 2003; Cho et al., 2007). Curcumin inhibited the expression of TNF-α-induced IL-1β, IL-6, TNF-α, cyclin E, MAPKs (JNK, p38 MAPK and ERK) and NF-κB in HaCaT cells. Because curcumin can reverse the anti-apoptotic function of TNF- α in skin cells, it may have potential for the treatment of psoriasis (Sun et al., 2012). Fourthly, as TNF blockers have been successfully used to treat psoriasis and since curcumin can block both the production and the action of TNF, curcumin may have potential as a treatment of psoriasis. Fifthly, our laboratory has shown that curcumin is a potent inhibitor of phosphorylase kinase (PhK) activity (Reddy and Lokesh, 1996), the elevation in which has been correlated with psoriatic activity (Heng et al., 2000).

Heng et al. (2000) investigated whether the anti-psoriatic activity of curcumin in patients is due to suppression of PhK activity. In this study, PhK activity was assayed in four groups of 10 subjects each: (i) active untreated psoriasis; (ii) resolving psoriasis treated by calcipotriol, a vitamin D3 analogue and indirect inhibitor of PhK; (iii) curcumin treatment (1% in gel); and (iv) 10 normal non-psoriatic subjects. PhK activity, from highest to lowest, was as follows: the active untreated psoriasis group, the calcipotriol-treated group, the curcumintreated group and the non-psoriatic subjects. The decrease in PhK activity in the curcumin-treated and calcipotriol-treated psoriasis groups was associated with a decrease in the expression of the keratinocyte transferrin receptor, a reduced severity of parakeratosis and a reduction in the density of epidermal CD8+ T-cells. The authors of this study concluded that drug-induced suppression of PhK activity is associated with resolution of psoriatic activity and that the antipsoriatic activity of curcumin may be achieved through its modulation of PhK.

The safety and efficacy of oral curcumin in patients with moderate to severe psoriasis has been investigated in a prospective phase II, open-label, Simon's two-stage clinical trial (Kurd et al., 2008). Twelve patients with chronic plaque psoriasis were enrolled in this study and were given a 4.5 g curcumin capsule per day for 12 weeks; this was followed by a 4 week observation period. Curcumin was well tolerated and all participants completed the study. However, the response rate was low and possibly caused by a placebo effect or the natural history of psoriasis. Nevertheless, two patients who responded to the treatment showed 83–88% improvement at 12 weeks of treatment. There were no study-related adverse events that necessitated participant withdrawal. Small sample size and the lack of a control (placebo) group were the limitations of the study.

Refractory asthma

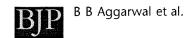
Patients' asthma is considered refractory when they experience persistent symptoms, frequent asthma attacks and/or low lung function despite taking asthma medications. Some patients with refractory asthma have to take oral steroids

such as prednisone to manage their symptoms. TNF- α has been shown to have a pathobiological role in asthma, mainly in severe refractory asthma and in chronic obstructive pulmonary disease (COPD) (Matera *et al.*, 2010). Thus, TNF- α inhibitors (infliximab, golimumab and etanercept) are now regarded as potential new medications in asthma and COPD management.

Numerous rodent studies suggest that curcumin may also have potential for treatment of asthma. Firstly, curcumin (at 20 mg·kg⁻¹ bodyweight) was reported to attenuate allergeninduced airway hyper-responsiveness in sensitized guinea pigs (Ram et al., 2003). When administered to mice, it was found to prevent ovalbumin-induced airway inflammation by regulating NO (Moon et al., 2008) and, in a more recent study, to diminish the development of allergic airway inflammation and hyper-responsiveness, possibly through inhibition of NF-κB activation in asthmatic lung tissue (Oh et al., 2011). For these studies, BALB/c mice were sensitized to ovalbumin, allowing analysis of the effects of curcumin administration (200 mg·kg-1 bodyweight per day, i.p.) on airway hyper-responsiveness, inflammatory cell number and IgE levels in bronchoalveolar lavage fluid. Ammar el et al. (2011) also examined the anti-inflammatory activity of curcumin in a murine model of asthma and showed it downmodulated the serum levels of IgE, iNOS, transforming growth factor β1 and mRNA expression of TNF-α. Secondly, curcumin has been shown to have therapeutic potential for controlling allergic responses. Animals exposed to latex showed enhanced serum IgE; latex-specific IgG1, IL-4, IL-5 and IL-13; eosinophils; and inflammation in the lungs (Kurup et al., 2007). Intragastric treatment of latex-sensitized mice with curcumin demonstrated a diminished Th2 response with a concurrent reduction in lung inflammation. Eosinophilia in the curcumin-treated mice was markedly reduced, as was the expression of the co-stimulatory molecules (CD80, CD86 and OX40L) on antigen-presenting cells, and expression of MMP-9, ornithine amino transferase and thymic stromal lymphopoietin genes was also attenuated. Thirdly, curcumin was found to reverse corticosteroid resistance in monocytes exposed to oxidants by maintaining histone deacetylase-2 activity (Meja et al., 2008). Although no clinical data are available yet, all of these pre-clinical studies suggest that curcumin has potential as a therapeutic for asthma.

Conclusions

Overall, all these studies suggest that curcumin can suppress pro-inflammatory pathways linked with most chronic diseases. It can block both the production and the action of TNF. Curcumin also binds to TNF directly. Evidence for curcumin as a TNF blocker has been obtained in both *in vitro* and *in vivo* studies. However, only a few studies have demonstrated that curcumin is effective at inhibiting TNF production in humans. Unlike most other TNF blockers, curcumin can be given orally. In addition, it is quite safe and affordable. However, more studies are needed in humans to prove that curcumin has the ability to be an effective treatment of various pro-inflammatory conditions.



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Conflict of interest

The authors declare no conflicts of interest.

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Curcumin as TNF blocker



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