

Novel treatment strategies for chronic kidney disease: insights from the animal kingdom

Peter Stenvinkel¹, Johanna Painer², Makoto Kuro-o³, Miguel Lanasa⁴, Walter Arnold², Thomas Ruf², Paul G. Shiels⁵ and Richard J. Johnson⁴

Abstract | Many of the >2 million animal species that inhabit Earth have developed survival mechanisms that aid in the prevention of obesity, kidney disease, starvation, dehydration and vascular ageing; however, some animals remain susceptible to these complications. Domestic and captive wild felids, for example, show susceptibility to chronic kidney disease (CKD), potentially linked to the high protein intake of these animals. By contrast, naked mole rats are a model of longevity and are protected from extreme environmental conditions through mechanisms that provide resistance to oxidative stress. Biomimetic studies suggest that the transcription factor nuclear factor erythroid 2-related factor 2 (NRF2) offers protection in extreme environmental conditions and promotes longevity in the animal kingdom. Similarly, during months of fasting, immobilization and anuria, hibernating bears are protected from muscle wasting, azotaemia, thrombotic complications, organ damage and osteoporosis — features that are often associated with CKD. Improved understanding of the susceptibility and protective mechanisms of these animals and others could provide insights into novel strategies to prevent and treat several human diseases, such as CKD and ageing-associated complications. An integrated collaboration between nephrologists and experts from other fields, such as veterinarians, zoologists, biologists, anthropologists and ecologists, could introduce a novel approach for improving human health and help nephrologists to find novel treatment strategies for CKD.

The evolution of species — mediated by genetic and epigenetic modifications over the past 3.8 billion years — has given rise to a wide variety of adaptations to different environments. This observation has led to the proposal that insights into adaptive mechanisms observed in nature could aid the development of therapeutic approaches for human disease¹. Comparative physiology — a subdiscipline of physiology that is based on Krogh's principle, which states “for such a large number of problems there will be some animal of choice, or a few such animals, on which it can be conveniently studied” — involves the comparison of organ systems within different taxa². Homer Smith's insightful work, for example, used a comparative physiology approach based on studies of fish and amphibians³ to form the basis of many aspects of renal physiology. Similarly, Sperber⁴ studied correlations between dietary habits, ecological distribution, urine-concentrating ability and kidney morphology in 1944. The emerging field of biomimetics explores adaptive mechanisms of a given species and imitates — or takes inspiration from

— these mechanisms to solve human problems (TABLE 1; [Supplementary information S1](#) (table)). Biomimetics is a particularly interdisciplinary field that can be used to identify new approaches to disease (FIG. 1), such as the underlying mechanisms of longevity in naked mole rats⁵, resistance to long-term renal hypoxia in seals⁶ and preserved renal function in hibernating bears^{7,8}. However, it is important to emphasize that when interpreting data from biomimetic studies, one should consider the likelihood of comparative animal data offering meaningful solutions when extrapolated to human disease. The physiological mechanisms that have evolved to enable adaptation of healthy animals to extreme environments may not necessarily be the same mechanisms that should be harnessed to avoid disease in humans.

The prevalence of chronic kidney disease (CKD) is rising worldwide. Approximately 10–15% of the global population suffers from CKD and its associated complications⁹, particularly cardiovascular disease, infectious complications, osteoporosis, muscle wasting, frailty and

Correspondence to P.S.
peter.stenvinkel@ki.se
 Division of Renal Medicine
 M99, Karolinska University
 Hospital, Karolinska
 Institutet, Hälsovägen 13,
 14157 Stockholm, Sweden.

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Key points

- Biomimetic studies of non-laboratory wild animals are useful for identifying mechanisms that protect or increase susceptibility to disease
- Domestic and captive felids are vulnerable to chronic kidney disease (CKD), supporting the hypothesis that high protein intake — particularly from red meats and in combination with dehydration — is nephrotoxic
- Extreme models of ageing, such as Hutchinson–Gilford progeria syndrome and the naked mole rat, can be used to investigate the mechanisms of vascular progeric processes in CKD
- Current evidence suggests that elevated serum phosphate levels promote ageing and cellular senescence
- The transcription factor nuclear factor erythroid 2-related factor 2 (NRF2) may offer protection against diseases in extreme environmental conditions and may promote longevity in the animal kingdom; NRF2 agonists (such as resveratrol and sulforaphane) might improve the uraemic complications of CKD
- Lipid composition of membranes has a role in seasonal acclimatization of metabolic activities in the animal kingdom
- Hibernating wild bears with anuria are protected against many of the complications observed in humans with CKD, such as muscle wasting, osteoporosis and azotaemia; future studies should investigate the mechanisms behind these protective effects

premature ageing^{10,11}. Nephrologists are faced with limited treatment options for patients with CKD, and advances in dialysis technology have not yet translated into markedly better outcomes¹⁰. As the majority of randomized controlled trials for CKD therapies have been negative¹⁰, an urgent need exists to find novel treatment options for this patient group. Here, we discuss some examples of renal biomimetics and how studies of the mechanisms by which animals adapt to hypoxia, oxidative stress, food deprivation and, conversely, to high-protein or high-phosphate diets may result in a better understanding of the uraemic phenotype (FIG. 1). Although biomimetic studies usually focus on the adaptive mechanisms that protect species from disease, changing environments (such as global warming, water availability or salinity in the oceans) can also lead to adaptations that may not offer full protection from these changes but may still shed light on disease mechanisms. Examples include the study of mechanisms that lead to the extinction of a species and the inability of that species to adapt to a changing environment.

High-protein diets and dehydration

High-protein diets that are rich in red meat accelerate the progression of both experimental and human CKD^{12,13}. The link between a high-protein diet and CKD

(FIG. 2) suggests that one can obtain mechanistic insights from studying mammals that live almost exclusively on a high-protein diet, such as the Felidae family (felids) and Desmodontinae (vampire bats). Interestingly, one group (felids) seems to be susceptible to CKD, while the other (vampire bats) seems to be protected.

CKD in felids

Felids consist of 37 species in the wild. Although they are considered among the world’s most successful carnivore families, they are particularly susceptible to kidney diseases, including polycystic kidney disease¹⁴, glomerulonephritis¹⁵, acute pyelonephritis, hypertension-associated CKD¹⁶ and nephrolithiasis¹⁷. The most common kidney pathology in domestic felids is chronic tubulointerstitial fibrosis, which is sometimes observed with glomerulosclerosis¹⁴. The prevalence of CKD in domestic cats has increased 75-fold (from 0.04% to 3%) during the past 4 decades, although this increase might be partially due to improved diagnostics^{14,18,19} and to increased non-steroidal anti-inflammatory drug (NSAID) consumption in the past decade²⁰. Even so, CKD is thought to affect 35–80% of geriatric domestic cats and is the most common cause of death in domestic cats >5 years of age¹⁸. Likewise, a necropsy study found renal lesions in 87% of large felids (mainly tigers, leopards and lions) held at zoos and safari parks in Germany²¹. In the wild, free-ranging felids experience a range of kidney diseases of differing origins, such as viral infections or amyloid deposition; however, free-ranging animals typically die from other causes before renal disease manifests or show only a mild form of disease¹⁹. Extrinsic environmental or dietary factors that might promote the development of kidney disease among felids in captivity seem to be absent among wild felids¹⁹.

As mentioned above, the most common renal pathology among captive and domestic felids is chronic tubulointerstitial fibrosis, which is associated with minimal or mild proteinuria, normal blood pressure, hypokalaemia, hyponatraemia or hypernatraemia, polydipsia and polyuria^{14,22} and an absence of diabetes mellitus²³. Hypertension, if present, is usually thought to be secondary to renal disease¹⁴. Microvascular lesions observed in chronic hypertensive renal injury are absent or only minimally present²⁴. The cause of this type of CKD remains unknown; however, it is unlikely that felids have evolved a selective susceptibility to CKD. Hence, one might hypothesize that the dramatic increase in felid CKD might reflect a new environmental exposure to which felids are particularly susceptible. Insights into the underlying mechanisms might be gained from comparisons with populations of humans and other animals that are either affected by or protected from renal disease, as discussed in further detail below. As CKD among felids has been best described in domestic cats and felids in wildlife parks, one possibility is that this disease might reflect the contamination of meat with a nephrotoxic substance. This scenario is similar to the epidemic of renal disease that occurred in vultures in India and Pakistan, which was ascribed to the practice of treating cattle with NSAIDs that contaminated the cattle meat²⁵.

Uraemic phenotype

Phenotype that includes several physical characteristics, such as vascular stiffness, sarcopenia, frailty, osteoporosis and left ventricular hypertrophy.

Chronic tubulointerstitial fibrosis

Diseases that affect the physiology of non-glomerular structures (tubules and/or the interstitium) in the kidney.

Author addresses

¹Division of Renal Medicine M99, Karolinska University Hospital, Karolinska Institutet, Hälsovägen 13, 14157 Stockholm, Sweden.

²Konrad Lorenz Institute of Ethology and Research Institute of Wildlife Ecology, Department of Integrative Biology and Evolution at the University of Veterinary Medicine, Savoyenstreet 1, 1160 Vienna, Austria.

³Division of Anti-Aging Medicine, Center for Molecular Medicine, Jichi Medical University, 3311–1 Yakushiji, Shimotsuke, Tochigi 329–0498, Japan.

⁴Division of Renal Diseases and Hypertension, 12700 East 19th Avenue, Room 7015 Mail Stop C281, University of Colorado Anschutz Medical Campus, Aurora, Colorado 80045, USA.

⁵Wolfson Wohl Translational Research Centre, University of Glasgow, Garscube Estate, Switchback Road, Bearsden, Glasgow G61 1QH, UK.

Table 1 | Selected animal models that are useful for comparative physiology studies

Species, family and/or order	Area	Mechanisms and possibilities
Naked mole rat (<i>Heterocephalus glaber</i>)	<ul style="list-style-type: none"> • Gerontology • Nephrology • Oncology • Cardiology 	These animals have developed protective mechanisms against cancer, hypoxia, cardiovascular ageing and oxidative stress (high NRF2 expression levels).
Vampire bat (<i>Desmodus rotundus</i>)	Nephrology	Blood-ingesting bats have a very high intake of proteins, which causes azotaemia (high serum urea levels). Studies of vampire bats may help to better understand how kidneys can be protected against protein overload.
Ursidae family (bears)	<ul style="list-style-type: none"> • Nephrology • Endocrinology • Cardiology • Orthopaedics • Transplantology 	Bears do not develop insulin resistance during summer despite a 25–50% accumulation in body weight (fat mass) from spring to autumn. Moreover, despite prolonged fasting, anuria and immobilization during hibernation, bears are protected from muscle wasting, pressure ulcers, thrombotic complications and osteoporosis. Studies of hibernating bears may help identify novel strategies to handle and prevent these complications as well as better ways of organ preservation.
Felidae family (cats)	Nephrology	Domestic and captive felids have a high incidence of chronic kidney disease. As members of this family are obligate carnivores, studies of felids may provide information on links between red meat consumption, gut microbiota and renal disease.
Phocidae family (seals)	Nephrology	Seals can survive prolonged asphyxia during underwater dives that last up to 120 min. Although their kidneys are subjected to prolonged vasoconstriction during diving, seals do not develop acute kidney injury.
Elephantidae family (elephants)	Oncology	The risk of elephants developing cancer is only 5% compared with 25% in humans, although they have 100 times as many cells. This protection may be due to the 20 copies of the tumour suppressor gene <i>TP53</i> , whereas humans have only one copy (two alleles).
Chimpanzee (<i>Pan troglodytes</i>)	Pharmacology	Chimpanzees have developed ways to protect themselves against pathogens by self-medicating with various plant leaves. Because one of these plants (thiarubine A) contains an antibiotic, systematic studies of these plants may help us find novel antibiotics.
Trochilidae family (hummingbirds)	Diabetology	Hummingbirds can switch their energy source from glucose to fructose, which maximizes fat storage and optimizes energy use to power their high-energy lifestyle (their heart rate can reach >1200 beats/min). Despite hyperglycaemia, they do not seem to develop diabetic complications.
Testudines order (turtles)	Neurology	Turtles have a high anoxic tolerance, and studies of these animals may help scientists to develop novel therapeutic strategies for cerebral ischaemia.
Wood frog (<i>Rana sylvatica</i>)	Physiology	Frozen wood frogs have 10–13-fold higher glucose concentrations in muscle and heart than other frog species that have been frozen in the laboratory. In addition, they have natural antifreeze glycolipids in muscle and internal organs to protect their cells. These mechanisms help them to survive over-wintering in average temperatures of –6.3 °C (minimum –18.1 °C) between October and May in the interior of Alaska. Studying wood frogs can help to understand limits to freezing tolerance.

The effect of red meat intake. Another potential mechanism underlying the high prevalence of CKD in felids might relate to their high intake of red meat (FIG. 2). To meet the high energy demands of their brain, which is relatively large in comparison with their body size²⁶, greater quantities of proteins are required, predominantly to generate glucose from amino acids through *de novo* gluconeogenesis. High-protein diets induce vasodilation of afferent renal arterioles, glomerular hypertension and hyperfiltration, which together accelerate the progression of pre-existing CKD in a variety of domestic and laboratory animals, including mice, rats and dogs²⁷. A high consumption of salt and animal proteins has also been linked to progression of CKD in humans^{12,28}, with increasing evidence indicating a greater effect of red meat consumption compared with that of other animal and vegetable protein sources^{12,13}.

Whether high-protein diets can induce *de novo* renal disease is less certain. One study reported that a commercial diet low in potassium and high in meat (40% protein) and phosphoric acid led to the development of tubulointerstitial lesions in five of nine domestic cats²⁹. In a human study, maintenance of a high-protein diet for 6 weeks increased estimated glomerular filtration rate (eGFR) by 4 ml/min/1.73 m² compared with a carbohydrate and unsaturated-fat diet in healthy individuals³⁰; however, whether long-term consumption of a high-protein diet promotes CKD is unclear. Although felids are obligate carnivores, their dietary acquisition of protein in the wild is intermittent and separated by days of fasting²⁴. By contrast, domestic cats and felids kept in zoos are often fed high-protein diets on a daily basis.

An examination of published data on 12 biochemical parameters of serum that can be used to evaluate renal

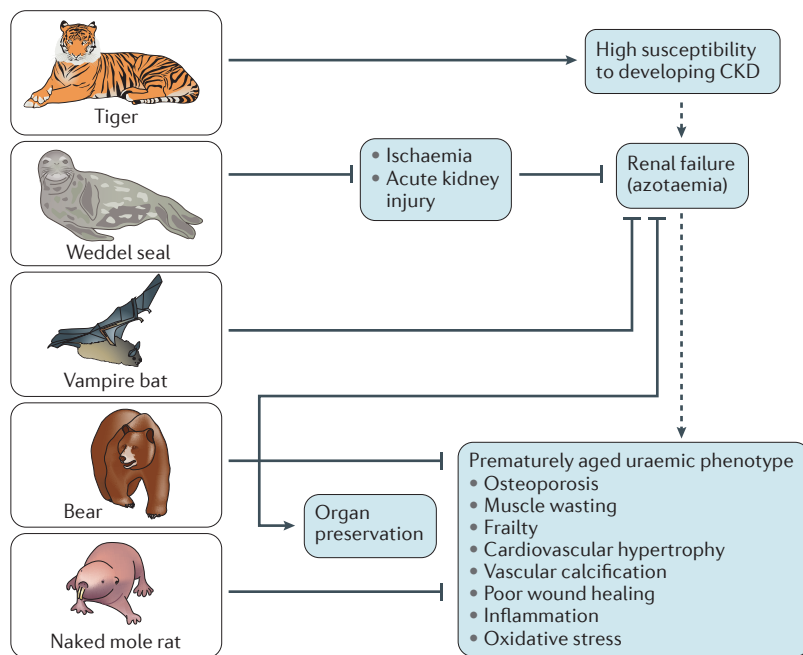


Figure 1 | Novel insight into treatment strategies of CKD from studies of wild animals. Several species in the animal kingdom have developed protective mechanisms against environmental stresses, and studying these mechanisms can provide insights into novel approaches to chronic kidney disease (CKD). For example, despite a long period of immobilization during hibernation, bears do not develop azotaemia, osteoporosis, inflammation, thrombosis, atherosclerosis and substantial muscle wasting, which could provide clues for better organ preservation. Naked mole rats (*Heterocephalus glaber*) are protected from oxidative stress, which could help to develop strategies to prevent or slow down premature ageing. As Weddell seals (*Leptonychotes weddellii*) are protected against prolonged episodes of kidney ischaemia during long periods of deep sea diving, they could provide insight to prevent acute kidney injury. Vampire bats (*Desmodus rotundus*) are protected against the consequences of high-protein intake, whereas felids (such as tigers) are particularly susceptible to CKD, most likely owing to their high intake of red meat.

functions in 97 mammalian species shows that differences in patterns of these parameters result in a clustering of species, separating carnivorous mammals from omnivorous or herbivorous mammals (Supplementary information S2 (figure)). This clustering is mainly due to higher levels of urea (+57%) and chloride (+13%), as well as reduced levels of alkaline phosphatase (−56%) among carnivores compared with mean values of the other species. This finding highlights the importance of diet, including protein source, on parameters of renal function and leads to the hypothesis that in humans, different types of diets (for example, vegetarian or highly carnivorous) might lead to similar differences in parameters of renal function.

Although the classical view is that renal injury induced by a high-protein diet is caused by changes in glomerular haemodynamics³¹ (FIG. 2), increasing evidence suggests that CKD risk is associated with protein originating from red meat and not with protein from dairy or vegetable-based sources³². For example, epidemiological studies conducted in Singapore¹² and the USA^{33,34} have shown that among different protein sources (including red meat, poultry, dairy products, fish, eggs and legumes), only red meat (beef, pork and lamb) and processed meat increased the risk of CKD. In

one study, individuals in the highest quartile of dietary red meat intake had a 1.4-fold greater risk of end-stage renal disease than those in the lowest quartile of red meat intake¹². Interestingly, diets rich in other protein sources, such as legumes and low-fat dairy products, were actually protective against CKD³³. Red and processed meat therefore seem to have direct nephrotoxic effects that increase the risk of CKD. Indirect support for differences in plant and animal proteins comes from studies in vegetarians. A study conducted in Taiwan showed that eGFR did not differ among 102 vegetarian Buddhist nuns compared with an equal number of age-matched omnivorous females³⁵. However, serum levels of sodium, glucose, urea and cholesterol, as well as blood pressure and urinary specific gravity, were lower among individuals in the vegetarian group³⁵. As vegetable proteins have different renal effects (lower GFR and renal plasma flow) than meat proteins³⁶ and as plant-based diets might protect against the development of CKD³⁷ and its complications³⁸, patients with CKD should be encouraged to consider a vegetarian diet³⁹.

In addition to CKD, red and processed meat have been linked to an increased mortality⁴⁰ and risk of other chronic diseases, such as cancer⁴¹, stroke⁴², coronary heart disease⁴³ and type 2 diabetes mellitus (T2DM)⁴⁴. Moreover, one study reported that the withdrawal of red meat from the diet of patients with T2DM reduced albuminuria and improved their serum fatty acid profile compared with their usual diet⁴⁵. Another study in patients with T2DM showed that adherence to a chicken meat-based diet for 1 year reduced urinary albumin excretion to levels comparable to those achieved by treatment with an angiotensin-converting enzyme (ACE) inhibitor⁴⁶. These findings imply that renal toxicity is generated by red meat per se and not total protein intake. Dietary management of CKD in domestic cats with a low-protein and low-phosphate (PO₄) diet was associated with increased survival compared with that of cats that did not undergo the dietary change⁴⁷. Further investigation is required to elucidate the potential differential effects of processed red meat, game meat and white meat (that is, chicken or fish) on renal function in felids.

Mechanism of red meat-induced CKD. Several factors have been proposed to be implicated in the disease-promoting effects of red meat⁴⁸ (FIG. 2). These include an associated high intake of sodium chloride (which increases blood pressure and stimulates vasopressin production and release by increasing serum osmolality), saturated fats (which drive mitochondrial oxidative stress), increased net acid production (which causes metabolic acidosis and acidic urine), the pro-oxidative effects of haem iron (which promotes oxidative stress), DNA damage caused by *N*-nitroso compounds (which leads to purine degradation and uric acid formation), the incorporation of non-human sialic acid into tissue (which promotes interaction with inflammation-provoking antibodies) and changes in the composition and/or metabolism of gut microbiota. For example, trimethylamine-*N*-oxide (TMAO), which is produced from the metabolism of red meat, eggs and fish by gut microbiota, induces

Glomerular haemodynamics
The regulation of efferent and afferent glomerular arteriolar resistance required to maintain a stable glomerular filtration rate.

Urinary specific gravity
Test that compares the density of urine to that of water.

***N*-Nitroso compounds**
Compounds found in processed meat that are formed endogenously from the intake of nitrite and nitrate.

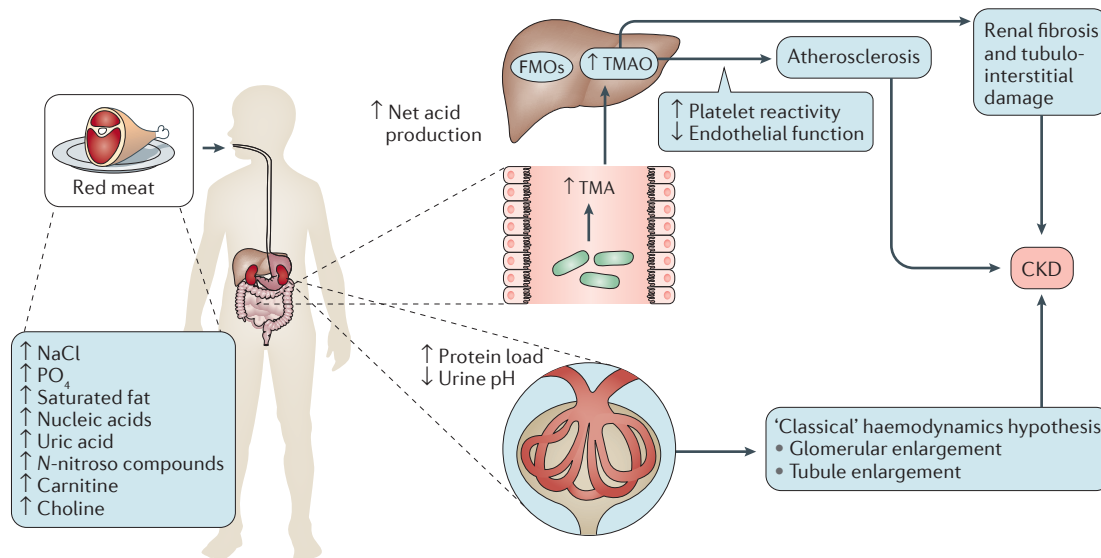


Figure 2 | Effect of red meat intake on kidney functions. Epidemiological studies suggest that red meat (but not other sources of protein) promotes chronic kidney disease (CKD). Several factors have been proposed to be implicated in the disease-promoting effects of a diet rich in red meat. In addition to hyperfiltration due to a high protein load (causing a haemodynamic insult), elevated levels of trimethylamine-*N*-oxide (TMAO) generated from gut microbiota and the metabolism of trimethylamine (TMA) in the liver via flavin-containing monooxygenases (FMOs) could contribute to CKD directly via renal fibrosis and indirectly via atherosclerosis. Additional pathophysiological mechanisms linking a high intake of red meat to cardiovascular disease, cancer and CKD have been reported. Increased intake of salt (NaCl), phosphate (PO₄), saturated fats, nucleic acids, uric acid, *N*-nitroso compounds, haeme iron, carnitine, choline and acid production with a high consumption of red meat may also contribute to the observed associations between increased red meat consumption and CKD. The intestinal microbiome represents a new potential therapeutic target for the prevention of CKD and for treatment of cardio-metabolic complications in CKD.

renal fibrosis in animal models⁴⁹ and inflammation in endothelial cells⁵⁰. Although diminished renal function impairs the ability to eliminate TMAO, it predicts outcomes in patients with CKD even after adjustment for other risk factors⁵¹. Inhibition of gut microbial trimethylamine (TMA) production prevented the development of atherosclerotic lesions in *ApoE*^{-/-} mice⁵². Moreover, exposure to carnitine (a major nutrient in red meat) in mice affects the composition of gut microbiota via the proatherogenic intermediate γ -butyrobetaine, which is converted into TMA and TMAO⁵³. Alterations in gut microbiota might also affect processes, such as haem-induced lipoperoxidation⁵⁴. Red meat consumption is also associated with increased intake of phosphate, which is associated with decreased renal function, inflammation and premature ageing⁵⁵. Moreover, phosphate activates nuclear factor- κ B (NF- κ B) signalling and promotes the generation of reactive oxygen species (ROS) in vascular smooth muscle cells (VSMCs)⁵⁶. This observation implies that the putative protective effects of antioxidative factors, such as nuclear factor erythroid 2-related factor 2 (NRF2) (BOX 1), on renal function should be investigated in the context of a diet rich in red meat.

The high content of nucleic acids in animal proteins probably also contributes to the nephrotoxic effects of red meat diets in humans and felids. Animal proteins are much more likely to raise serum uric acid levels than are proteins from vegetable and dairy sources⁵⁷. In domestic cats, a transient (up to 50-fold) increase in urine uric acid occurs following the ingestion of purine-rich animal

proteins compared with a purine-free diet, despite the presence of a uric acid-degrading enzyme (uricase)⁵⁸. In humans, consumption of animal proteins and/or purines also results in an acute rise in serum and urine uric acid levels^{59,60}, which is accompanied by a substantial acid load in urine that leads to a decrease in urine pH¹³. A urine pH of <5 is extremely common in cats with uraemic manifestations²², and a low urine pH (5.0–5.5) predicts stage 3 CKD in humans⁶¹. Although urate stones are relatively rare in cats⁶², urate crystalluria is a common problem in felids⁶³. Coupled with dehydration and heat exposure, we propose that urate crystalluria and/or uricosuria resulting from high protein intake could facilitate tubulointerstitial injury. Both soluble and crystalline forms of urate have been shown to induce inflammation in rat tubular cells *in vitro*⁶⁴. Indeed, signs of dehydration in felids with CKD are common²² and predictive for the development of CKD in domestic cats⁶⁵. In humans, although the aetiology of CKD in populations from Central America and Sri Lanka remains a subject of debate⁶⁶, we propose a role for heat stress and recurrent dehydration in the presence of high uric acid levels in disease pathophysiology⁶⁷.

Protein metabolism in vampire bats

Vampire bats (*Desmodus rotundus*) feed mostly on the blood of warm-blooded mammals. However, in contrast to humans and felids, they seem to be resistant to the detrimental metabolic effects of a high-protein intake. Their protein intake would be comparable to

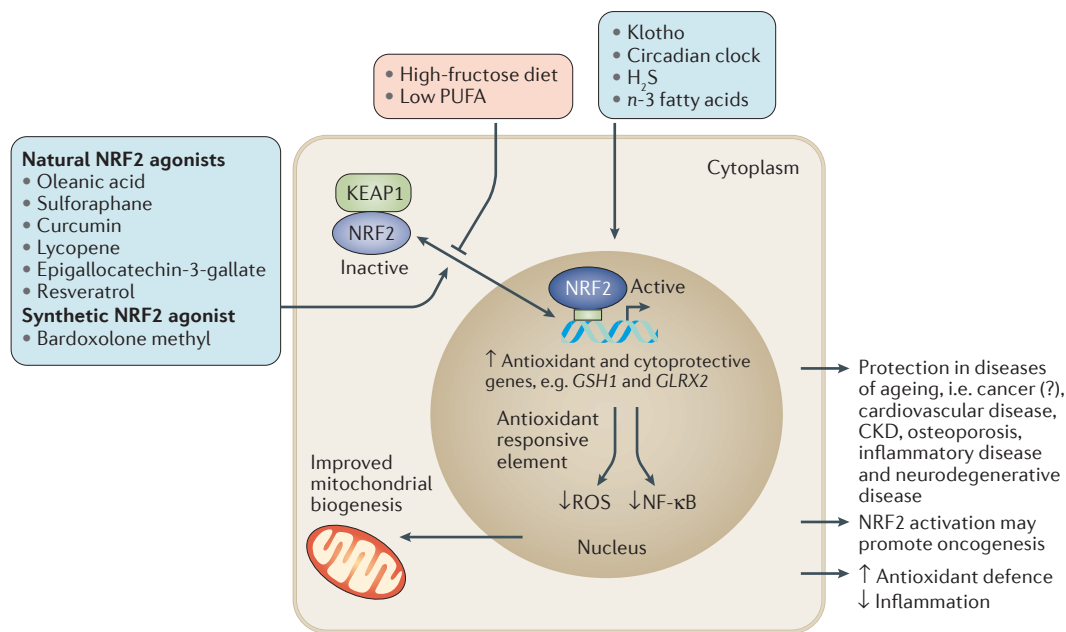
a daily intake of approximately 6 kg protein in a 70 kg man (as compared to a normal daily intake of 50–120 g in humans). As the consumption of >20 g of blood in a 20 min feed increases the body weight of vampire bats by 20–30%, they rapidly absorb the blood plasma and start urine production within 2 min of feeding⁶⁸. The blood urea concentration of vampire bats is 27–57 mmol/l (compared with 3–8 mmol/l in healthy humans), depending on the time point after feeding⁶⁹. Despite this high protein intake, the vampire bat does not have larger

kidneys than mammals of similar size⁷⁰, which suggests no difference in glomerular number and glomerular capillary surface area. Indeed, indirect allometric calculations indicate that the vampire bat's GFR is not greater than that of similarly sized mammals⁶⁹; however, to our knowledge, GFR measurements have not been performed. Of interest, mammalian blood has a lower relative purine content than does red meat⁷¹. Whether this difference accounts for the differential risk of CKD between felids and vampire bats is speculative.

Box 1 | The cytoprotective effects of the transcription factor NRF2

The transcription factor nuclear factor erythroid 2-related factor 2 (NRF2) upregulates the expression of cell-detoxifying enzymes in response to oxidative stress. Activators of NRF2 induce structural changes in Kelch-like ECH-associated protein 1 (KEAP1), which allows nuclear translocation of NRF2. In the nucleus, NRF2 initiates the transcription of >250 target genes, such as haemoxygenase, catalase and glucose-6-phosphate 1-dehydrogenase, which are important for antioxidant defences, through binding to antioxidant response elements (see the figure).

Nrf2-knockout mice have increased susceptibility to kidney damage. As impaired NRF2 activation is observed in renal fibrosis, focal segmental glomerulosclerosis and hypertensive kidney disease, NRF2-targeting therapies should be of interest for the study of chronic kidney disease (CKD) progression. Patients on haemodialysis have downregulated levels of NRF2 coupled with an upregulation of nuclear factor-κB (NF-κB)²⁷⁶ and display a phenotype characterized by persistent systemic inflammation⁷⁹ and increased oxidative stress²⁷⁷. Given the potential contribution of a repressed NRF2 system in premature ageing, both synthetic compounds, such as the potent NRF2 agonist bardoxolone methyl²⁷⁸, and natural nutrigenomic compounds, such as sulforaphane²⁷⁹, pomegranate polyphenols²⁸⁰, curcumin²⁸¹, ethanol extract of *Alisma orientale* tubers²⁸² and cinnamon polyphenols²⁸³, that restore NRF2 expression could slow progression and ageing-related CKD²⁸⁴. Indeed, as sulforaphane (found in broccoli) inhibits restenosis by suppressing inflammation and proliferation of vascular smooth muscle cells (VSMCs) in a carotid injury model²⁸⁵, it has been suggested that dietary activators of NRF2 inhibit atherogenesis²⁸⁶. Moreover, sulforaphane suppresses NRF2-mediated hepatic glucose production and attenuates exaggerated glucose intolerance by an order of magnitude similar to that of metformin in patients with type 2 diabetes mellitus (T2DM)²⁷⁹. However, forced overexpression of NRF2 might not always be enough to restore adaptive responses²⁸⁷. For example, bardoxolone methyl increased the risk of heart failure compared with placebo in a clinical trial with patients with T2DM and stage 4 CKD²⁸⁸, which highlights potential limitations of manipulating transcription factors. Although activation of NRF2 leads to improved antioxidant defences, whether this effect is independent of any influence on mitochondrial dynamics remains to be determined. Sulforaphane, for example, modulates the KEAP1–NRF2 antioxidant element response signalling pathway yet is a NRF2-independent inhibitor of mitochondrial fission²⁸⁹. Whether such an effect for bardoxolone methyl contributed to its failure owing to excess mortality²⁸⁸ remains to be proven. In future clinical trials of this compound, attention should be given to the dose-dependent effects on CKD progression²⁹⁰. Whereas too little NRF2 activity can result in loss of cytoprotection, diminished β-oxidation of fatty acids and lower antioxidant capacity, too much NRF2 activity may perturb the homeostatic balance and promote overproduction of reduced glutathione and NADPH²⁹¹. H₂S, hydrogen sulfide; PUFA; polyunsaturated fatty acids; ROS, reactive oxygen species.



Nutrigenomic compounds
Bioactive nutrients that have an effect on or interact with the genome. Nutrigenomics also encompasses the effect of genetic variations on the absorption, metabolism, elimination or biological effects of various nutrients.

Ageing and longevity

Ageing has been defined as an accumulation of deficits occurring in different individuals in different ways and with varied rates in different organs⁷². Ageing is an actively regulated process influenced by genetics, epigenetics, lifestyle, nutrition and psychosocial factors⁷³, which may act synergistically, independently or cumulatively. The ageing process is characterized by a series of hallmarks, including genomic instability, telomere attrition, epigenetic alterations, loss of proteostasis, dysregulated nutrient sensing, stem cell exhaustion, mitochondrial dysfunction, altered intercellular communication and cellular senescence⁷⁴, which are common across different taxa and affected by the uraemic milieu^{73,75,76}.

Ageing and kidney disease

In addition to the progressive loss of renal function, ageing in humans, rats and many other mammals is associated with the development of glomerulosclerosis and interstitial fibrosis^{77,78}, which are linked to impaired autoregulation of renal blood flow and impaired angiogenesis, epigenetic modifications, endothelial dysfunction, oxidative stress and inflammation⁷³. Chronic inflammation (also known as inflammageing) is an important driver of premature uraemic ageing⁷⁹ and manifests with an increased frequency of age-related complications, such as vascular stiffening, osteoporosis, muscle wasting, depression, cognitive dysfunction and frailty⁷⁹. In addition, persistent mitochondrial dysfunction with increased generation of ROS features in both normative ageing⁸⁰ and progressive CKD⁸¹.

Whether ageing-associated renal disease is inevitable⁸² or modifiable⁸³ remains controversial. The use of senolytic agents, which selectively remove senescent cells, in preclinical models suggests that ageing-associated renal disease is modifiable⁷⁵, and direct improvement of physiological function following removal of these cells indicates causality. These cells are non-proliferative, are resistant to apoptosis and are generated in response to genotoxic stress and resulting DNA damage as part of normative ageing. Loss of age-related regenerative capacity in tissues and organs occurs as a direct consequence, and a senescence-associated pro-inflammatory milieu subsequently develops. The selective removal of senescent cells from tissues and organs via immune-mediated clearance is dysregulated during normative ageing and contributes to inflammageing⁸⁴. The accumulation of these cells has been observed across a broad spectrum of non-communicable diseases.

Ageing in the animal kingdom

One way to improve our understanding of ageing processes and senescence is to study animals with unusual longevity. Long-lived animals are found across the taxonomic spectrum, such as in certain mammals, birds, sea turtles and fish. For example, extreme longevity is observed in the ocean quahog (*Arctica islandica*; >500 years)⁸⁵ and the Greenland shark (*Somniosus microcephalus*; ~400 years)⁸⁶. The study on the ocean quahog supports the notion that chronic low-grade inflammation

in the cardiovascular system is a ubiquitous feature of ageing⁸⁵. Other interesting candidates for studies of reduced senescence include the rougheye rockfish (*Sebastes aleutianus*) and the bowhead whale (*Balaena mysticetus*), both with documented lifespans of >200 years. Interestingly, ageing rockfish do not show signs of organ degeneration or a decline in liver lysosomal function⁸⁷, which are typically observed in normative ageing. By contrast, examples of exceptionally short-lived species that exhibit an accelerated expression of ageing biomarkers are found in the family of Cyprinodontidae (killifish), which have a maximal lifespan of only 13 weeks⁸⁸. Thus, a better understanding of the mechanisms by which some animals have delayed or accelerated ageing processes^{87,89} may provide insights into not only the process of ageing in humans but also ageing-related kidney disease.

Insights from the naked mole rat. Naked mole rats (*Heterocephalus glaber*) have emerged as a good model organism to study ageing and ageing-related diseases. These subterranean rodents are rarely exposed to sunlight and have no obvious dietary source of vitamin D⁹⁰. However, despite having undetectable calcifediol (25(OH)D) levels (the precursor of vitamin D), their calcium phosphate homeostasis is adequately maintained⁹¹. Although they have a small body size and are constantly exposed to hypoxia, oxidative stress and hypercapnia, they can live >30 years and maintain a healthy cardiovascular and reproductive status as well as body composition throughout their life⁵. Interestingly, the structure and function of their proteins are not affected by their substantial exposure to oxidative stress⁹², and they display high levels of autophagy and efficient removal of stress-damaged proteins throughout life⁹³. In contrast to humans and other rodents,⁹⁴ these animals preserve normal vascular and cardiac function with ageing^{95,96} and are resistant to the development of cancer⁹⁷. Moreover, their bone mineral density, articular cartilage and nitric oxide sensitivity of VSMCs are not affected by ageing⁹⁸. Whereas most nephropathologies seem to be absent in naked mole rats, cases of nephrocalcinosis have been reported⁹⁹.

NRF2-mediated antioxidant activity. Some of the molecular pathways that protect these animals from cancer have been elucidated. For example, one study reported that the fivefold higher production of high-molecular-weight hyaluronan in fibroblasts protects naked mole rats from cancer¹⁰⁰. High expression levels of the transcription factor NRF2, which stimulates intracellular antioxidant activity by regulating the expression of many target genes involved in the antioxidant response (BOX 1), may also protect the naked mole rat from cellular damage. In addition to antioxidative activities, NRF2 has other important functions, such as regulating NF- κ B activity, which may play a part in mitochondrial homeostasis¹⁰¹ and may decelerate the ageing process¹⁰². As NRF2 expression correlates positively with maximum lifespan in long-lived rodents¹⁰³, diminished NRF2 activity may be important for the ageing phenotype of organisms as diverse as worms, flies and mammals¹⁰⁴.

Telomere attrition

Telomeres are the protective endcaps of chromosomes. Attrition, or shortening, of telomeres is a form of tumour suppression and may be due to inflammation and oxidative stress as well as exposure to infectious agents, resulting in limited stem cell function, regeneration and organ maintenance during ageing.

Uraemic milieu

Toxic internal milieu in patients with uraemia that is characterized by accumulation of uraemic toxins and waste products that promote inflammation, oxidative stress, carbonylation, calcification and endothelial dysfunction.

Senescent cells

Cellular senescence is an irreversible cell cycle arrest mechanism that acts to protect against cancer. Senescent cells also have a role in complex biological processes, such as development, tissue repair and age-related disorders.

Hypercapnia

Abnormally elevated carbon dioxide (CO₂) levels in the blood.

High-molecular-weight hyaluronan

A high-molecular-weight polysaccharide found in the extracellular matrix, especially in soft connective tissues.

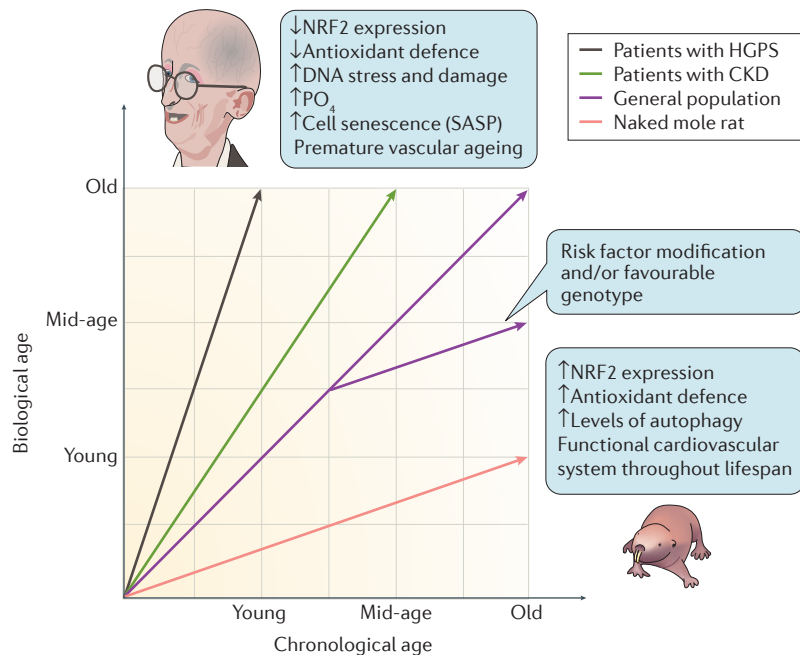


Figure 3 | Extreme models of ageing with a marked discrepancy between chronological and biological age can be used to learn more about progeric processes in CKD. Children with the rare Hutchinson–Gilford progeria syndrome (HGPS) express truncated lamin (progerin) that mediates premature ageing, especially in the cardiovascular system, resulting in premature death from stroke or myocardial infarction. On the other hand, naked mole rats undergo negligible senescence and can live for >30 years without signs of cardiovascular ageing. Integration of data from these two models of ageing can reveal detailed mechanisms of the progeric phenotype. A high biological age in chronic kidney disease (CKD) is characterized by a premature ageing phenotype, which includes vascular stiffness, frailty, osteoporosis and sarcopenia, as well as high levels of inflammation, carbonylation and oxidative stress. NRF2, nuclear factor erythroid 2-related factor 2; SASP, senescence-associated secretory phenotype.

Antagonistic pleiotropy
Scenarios in which one gene contributes to multiple traits, whereby at least one of these traits is beneficial and at least one is detrimental to the organism's health.

Phosphate appetite
A well-documented behaviour in animals that is induced by phosphate deficiency, which is especially common among herbivores.

Evidence for a role of NRF2 in ageing has also been reported in humans. For example, children with the rare Hutchinson–Gilford progeria Syndrome (HGPS), which is caused by a mutation in prelamin A/C, age extremely prematurely, are subject to increased oxidative stress and have a repressed NRF2 pathway¹⁰⁵. As reactivation of NRF2 reversed the cellular ageing defects in cells from patients with HGPS and in an animal model of HGPS repression, NRF2-mediated transcription seems to have a pathogenic role in the progeric phenotype¹⁰⁵. As HGPS shares many features common to age-associated diseases, it has been regarded as a model system to better understand ageing processes in chronic diseases¹⁰⁶. For unknown reasons at present, children with HGPS do not seem to have an increased risk of CKD despite a prematurely aged phenotype, which may reflect a feature of antagonistic pleiotropy¹⁰⁶. Despite apparent differences in the pathways underlying HGPS and CKD, we suggest that models of ageing and longevity, such as HGPS and the naked mole rat, can be used to study factors that underlie progeric processes in CKD (FIG. 3).

Vascular calcification and phosphate
Phosphate and calcification in vertebrates. The composition of sea water is similar to that of the human body in regard to the abundance of elements¹⁰⁷ (Supplementary

information S3 (table)), consistent with the view that life originated from the sea¹⁰⁸. Of the ten most abundant elements in the human body, only phosphorous is not among the ten most abundant elements present in sea water¹⁰⁹, indicating that organisms selectively accumulated phosphorus (in form of phosphate within cells) at some point in time during evolution (Supplementary information S4 (figure)). Phosphate is a major component of nucleic acids and membrane phospholipids and has a key role in numerous intracellular functions, including ATP synthesis and function and kinase-mediated signal transduction¹¹⁰. Phosphate is so fundamental to life that its deficiency can be fatal, which may be a reason for the evolution of phosphate appetite¹¹¹.

Although intracellular phosphate is essential to all forms of life, accumulation of extracellular phosphate first emerged in bony fish during the evolution of skeletons¹¹². Unlike invertebrates, which have a calcium carbonate (CaCO₃)-based exoskeleton, most skeletons of vertebrates consist of calcium and phosphate, especially in the form of calcium hydroxyapatite (Ca₁₀(PO₄)₆(OH)₂)¹¹³. This acquisition was likely required for terrestrial vertebrates to support their body weight on land. The extracellular fluid of vertebrates consists of a highly saturated solution of calcium and phosphate¹¹⁴, which enables bone formation simply by controlling where to provide a cue for nucleation of calcium phosphate, such as production and secretion of bone matrix proteins by osteoblast lineage cells. Undesired nucleation within the extrasosseous tissues is achieved by maintaining an extracellular phosphate concentration within a narrow range, in a process that is partially controlled by fibroblast growth factor 23 (FGF23) and its obligate co-receptor Klotho (encoded by *Kl*)¹¹². Interestingly, *Kl*-knockout mice have a 12-fold reduction in lifespan and a 2-fold increase in extracellular phosphate concentration compared with wild-type mice¹¹⁵. Secreted Klotho also exerts multiple functions independently of FGF23, such as inhibition of insulin-like growth factor 1 activity and upregulation of antioxidant enzyme expression levels. These additional functions may contribute to the anti-ageing properties of Klotho¹¹⁶. Furthermore, Klotho induces NRF2 expression and subsequent antioxidant defence mechanisms¹¹⁷, which links altered NRF2 expression to bone mineral metabolism and phosphate homeostasis¹¹⁸.

Phosphate in ageing and calcification. Vascular calcification is a common feature of the progeric uraemic phenotype¹¹⁹ and is linked to senescence¹²⁰. High extracellular phosphate levels, which often occur in combination with elevated calcium levels, increase the risk of calcium phosphate deposition in the vasculature and vascular calcification¹¹⁹. Cell culture studies have shown that a high phosphate concentration induces cellular senescence¹²¹ and leads to the conversion of VSMC to osteoblastic cells^{122,123}, a process that can be prevented by inhibiting calcium phosphate precipitation using pyrophosphate, phosphonoformic acid^{124,125} and phosphate binders¹²⁶. Consistent with the observation that a high phosphate concentration induces cellular senescence and that accumulation of senescent

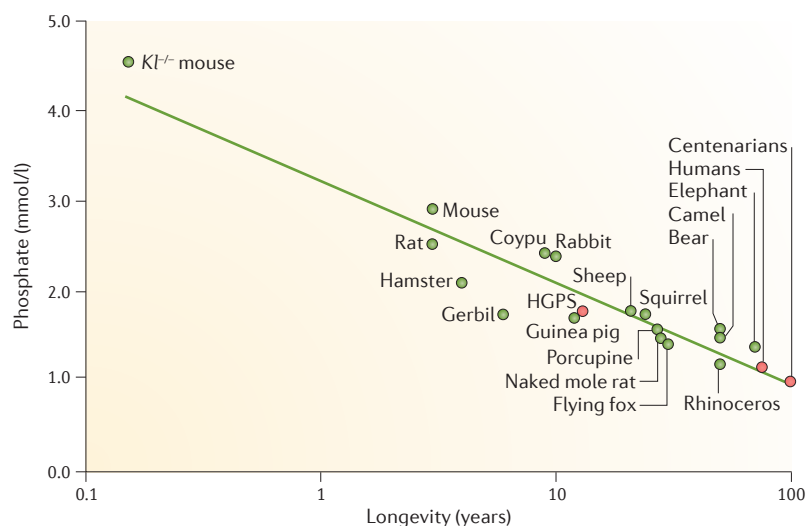


Figure 4 | The role of phosphate in ageing. Extracellular serum phosphate levels and maximum lifespan in different mammals, including humans (general population, centenarians and children with Hutchinson–Gilford progeria syndrome (HGPS); red circles), are inversely correlated. This inverse correlation indirectly suggests that high phosphate levels promote progeria and cellular senescence. α -Klotho-deficient ($Kl^{-/-}$) mice have high phosphate levels and a shortened lifespan compared to that of wild-type mice. In addition, dietary phosphate shortens the lifespan of $Kl^{-/-}$ mice further, which is likely mediated by the AKT–mTORC1 pathway (RAC- α serine/threonine-protein kinase–mechanistic target of rapamycin complex 1)¹⁴⁰. Mouse (*Mus musculus*), rat (*Rattus*), hamster (Cricetinae), coypu (or nutria) (*Myocastoridae*), rabbit (Leporidae), gerbil (Rodentia), guinea pig (*Cavia porcellus*), sheep (*Ovis aries*), squirrel (Sciuridae), porcupine (Hystricidae and Erethizontidae), naked mole rat (*Heterocephalus glaber*), flying fox (*Pteropus*), bear (Ursidae), camel (*Camelus*), rhinoceros (Rhinocerotidae), elephant (Elephantidae), human (*Homo sapiens*). Adapted with permission from REF. 128, Elsevier.

cells accelerates ageing of the organism¹²⁷, a negative correlation exists between extracellular phosphate levels and longevity across mammalian species¹²⁸ (FIG. 4). For example, children with HGPS have elevated phosphate levels, develop rapid vascular calcification and typically die of stroke or myocardial infarction as teenagers¹²⁹. Furthermore, expression of progerin (the mutated form of prelamin A associated with HGPS) in VSMCs leads to a decrease in extracellular pyrophosphate¹³⁰. As pyrophosphate protects VSMCs from calcification, high serum phosphate and low extracellular pyrophosphate may contribute to accelerated vascular calcification in HGPS. In the general population, high phosphate levels are associated with premature vascular ageing¹³¹, shortened telomere length, reduced DNA methylation content and elevated IL-6 (REF. 55), which are all biological markers of ageing.

Studies from the past few years have provided insights into the mechanisms by which high extracellular phosphate levels lead to vascular calcification. Under conditions of inflammation, oxidative stress and high extracellular phosphate levels, nanoparticle calcium phosphate precipitates can develop in the vasculature, despite the presence of multiple endogenous inhibitors of calcium phosphate deposition^{132,133} and can grow to form calciprotein particles (CPPs). CPPs are aggregates of α 2-HS-glycoprotein (AHSG) loaded with calcium phosphate precipitates and dispersed as colloids in the

blood. These CPPs may play a part in CKD progression, as recent clinical studies showed that serum CPP levels correlated with vascular calcification and/or stiffness^{132,134} and predicted mortality in patients on dialysis¹³⁵. Of interest, tigers have particularly high levels of phosphate (1.7 ± 0.3 mmol/l) and serum creatinine (265 ± 62 μ mol/l), suggesting it would be of interest to determine levels of serum CPP and FGF23 in these animals¹³⁶.

Therapeutic strategies to prevent vascular calcification in CKD. The best current approach to prevent vascular calcification in CKD is dietary phosphate restriction or chelation through the use of phosphate binders¹³⁷. However, a consequence of dietary phosphate restriction is reduced protein intake, which can lead to protein–energy wasting and inadvertently increased mortality¹³⁸. A major problem with phosphate binder therapies is patient non-adherence due to the high pill burden and gastrointestinal adverse effects¹³⁹. The high risk for gastrointestinal adverse effects imply that phosphate binders may alter the gut microbiota. Alternative treatment strategies to prevent vascular calcification could potentially be derived from comparative physiology studies. For example, agents that stimulate NRF2 (BOX 1), block mechanistic target of rapamycin complex 1 (mTORC1) signalling¹⁴⁰ or reduce phosphate absorption, such as by inducing calcium phosphate precipitation in the gut with magnesium¹⁴¹, should be tested for their ability to decrease extracellular phosphate levels.

Of interest, a diet of highly fermentable carbohydrates (for example, starch) in captive wild ruminants, such as giraffes (*Giraffa camelopardalis*) — in combination with low calcium, high phosphate and low magnesium levels in the serum — is associated with premature death in these animals¹⁴². Because introduction of a diet with lower starch content led to higher magnesium and lower phosphate levels¹⁴², one potential approach would be to use resistant starch, which is a complex carbohydrate fermented by gut microbiota that increases colonic absorption of minerals in animals. Indeed, resistant starch has been suggested to be a novel dietary method to prevent diabetic CKD¹⁴³.

Caloric restriction and ageing

Although dietary phosphate restriction is one mechanism to slow vascular calcification and ageing, a more effective approach to extend the lifespan of animals is by caloric restriction¹⁴⁴, which has demonstrated effectiveness in both short-lived species, including flies, worms, rats and mice¹⁴⁵, and more long-lived species, such as primates¹⁴⁶. Fat stores, especially those generated during fructose metabolism, result in fructose-induced oxidative stress, which is associated with increased translocation of NRF2 to the nucleus, decreases in mitochondrial DNA content and mitochondrial dysfunction, with subsequent cellular apoptosis^{147,148}. In most animals, excess fat stores are maintained as a protective mechanism for periods of food shortage¹⁴⁹. Thus, as long as food is available on a daily basis, caloric restriction would be expected to reduce mitochondrial oxidative stress and preserve mitochondrial metabolism. Other ways to

Protein–energy wasting

A process characterized by a decline in body protein mass and energy reserves, including muscle and fat wasting and loss of visceral proteins.

Protein energy wasting is often associated with inflammation and is a strong predictor of mortality.

Caloric restriction

A reduction in calorie intake without incurring malnutrition or a reduction in essential nutrients. In a variety of species, such as yeast, fish, rodents and dogs, calorie restriction has been shown to slow the biological ageing process.

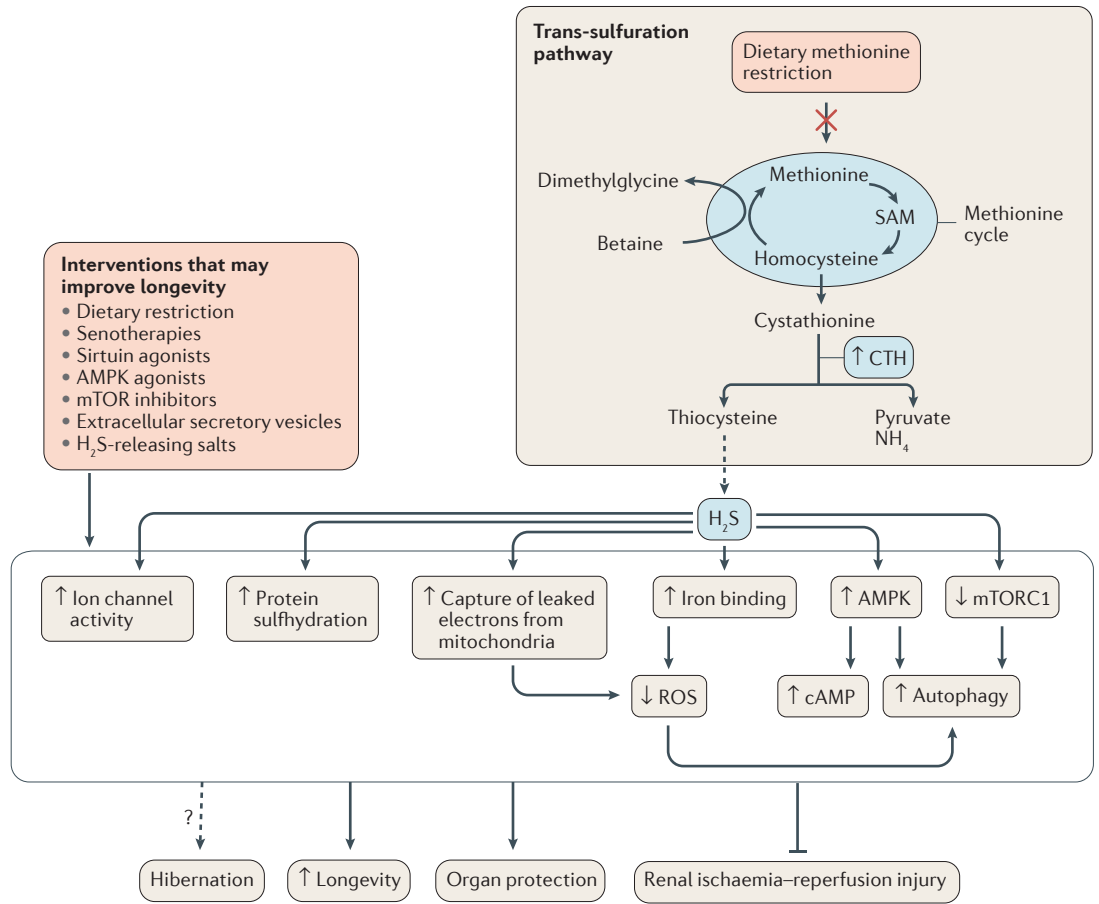


Figure 5 | Strategies to increase lifespan, protect organs and avoid renal ischaemia–reperfusion injury. Premature cardiovascular death and vascular progeria are prominent features of chronic kidney disease (CKD). On the basis of insights from long-lived animals and basic research, several treatment strategies have been identified that could be tested for their effect on longevity. Activation of the cytoprotectant functions of hydrogen sulfide (H₂S) via restriction of the sulfur-containing amino acid methionine is of major interest. Other potential treatment strategies that activate anti-inflammatory and antioxidant pathways include dietary restriction, senotherapies, sirtuin (NAD⁺-dependent protein deacetylases) agonists, 5′-AMP-activated protein kinase (AMPK) agonists, mechanistic target of rapamycin (mTOR) inhibitors, extracellular secretory vesicles and H₂S-releasing salts. Studies suggest that inhibition of mTOR and activation of nuclear factor erythroid 2-related factor 2 (NRF2) signalling by such therapies increases longevity, aids organ protection and decreases the risk of renal ischaemia and reperfusion injuries. Because hibernation shares features and pathways associated with longevity, it can be speculated that successful hibernation depends on these pathways. CTH, cystathionine γ-lyase; mTORC1, mTOR complex 1; NH₄, ammonia; ROS, reactive oxygen species; SAM, S-adenosyl-L-methionine.

Sirtuin

Sirtuins (or NAD⁺-dependent histone deacetylases) are a class of proteins that possess deacetylase activity and regulate important biological pathways and cellular processes, including ageing, inflammation, transcription and apoptosis. Sirtuin agonists include perestilbene and resveratrol.

Trans-sulfuration pathway

A metabolic pathway that involves the interconversion of homocysteine and cysteine via the intermediate cystathionine.

S-Sulfhydrylation

A post-translational modification that increases the catalytic activity of proteins. Physiological actions of sulfhydrylation include the regulation of endoplasmic reticulum stress signalling, inflammation and vascular tension.

mimic caloric restriction would be to administer agents that modulate cellular metabolism, including sirtuin agonists^{150,151} and 5′-AMP-activated protein kinase (AMPK) agonists¹⁵², the effects of which are mediated in part by the activities of the forkhead box protein (FOXO) family and the insulin signalling pathways¹⁵³. For example, resveratrol prolongs lifespan in the extremely short-lived killifish⁸⁸. We found that mice that cannot generate fructose, which are therefore protected from mitochondrial oxidative stress, were also protected from developing age-related renal disease⁸³. In theory, elevated expression of NRF2 also mimics caloric restriction, as knockdown of Kelch-like ECH-associated protein 1 (KEAP1) in mice results in accumulation of NRF2 and thus augments the activation of cellular stress responses, including fatty acid oxidation and lipogenesis¹⁵⁴.

Methionine restriction and ageing. In addition to caloric restriction, dietary restriction of proteins — especially the sulfur-containing amino acid methionine — also promotes longevity in various animal models¹⁵⁵ (FIG. 5). This effect is likely to be mediated through the cytoprotectant hydrogen sulfide (H₂S) gas and increased activation of the trans-sulfuration pathway, prevention of electron leakage from mitochondria and possible hermetic effects on the mTOR pathway and NRF2 activity¹⁵⁶. Under conditions of cellular stress, H₂S-mediated S-sulfhydrylation of KEAP1 leads to its disassociation from NRF2 and increased NRF2 nuclear translocation (BOX 1). This increases mRNA expression of NRF2-targeted downstream genes, such as GS homeobox 1 (*GSH1*; also known as *GSX1*) and glutathione reductase, and upregulates a range of cellular defences. In addition, methionine

restriction increases expression of the trans-sulfuration pathway enzyme cystathionine γ -lyase (CTH), resulting in increased H_2S production, which leads to AMPK activation and mTORC1 repression, thus reducing cellular stress and promoting physiological longevity¹⁵⁷. H_2S also binds iron and captures electrons leaked from mitochondria, which reduces mitochondrion-mediated ROS formation^{158,159}. In support of a role for H_2S activity in longevity, lower circulating methionine levels have been reported in naked mole rats than in shorter-lived laboratory rodents¹⁶⁰. The finding of low sulfide levels in naked mole rats and the inverse correlation between circulating sulfide levels and maximum longevity in six different species¹⁶¹ add to the complexity of understanding the role of H_2S in ageing. Thus, for prolonging lifespan, interconnections between methionine and caloric restriction in the context of comparative biology need to be investigated further.

Methionine restriction might be particularly important in preventing ageing and age-associated renal dysfunction⁵⁵. For example, an inverse correlation between methionine content in tissue proteins and longevity was reported in eight different species¹⁶². Although circulating methionine levels do not differ between patients with CKD and healthy controls¹⁶³, oral methionine loading in patients on haemodialysis leads to an accumulation of homocysteine and other methionine metabolites in plasma and red blood cells¹⁶⁴, indicating impairment of the trans-sulfuration pathway. High doses of vitamin B_6 and folic acid failed to mitigate this phenotype, indicating that it most probably was not due to a lack of these cofactors¹⁶⁴. Methionine restriction also increases the replicative lifespan and decelerates the accumulation of senescent cells across taxa from yeast to man¹⁶⁵. Consistent with these observations, we reported lower methionine levels in wild hibernating brown bears (*Ursus arctos*)⁸ and observed a fourfold increase in the methyl donor betaine during hibernation (P.G.S. and J.P., unpublished observations). Thus, it could be speculated that an increased production of H_2S protects the bears from ROS-mediated DNA damage. Moreover, dietary supplementation of H_2S in mice alleviates inflammation, aberrant methylation and dysfunction in a model of hypertensive kidney disease¹⁶⁶, suggesting that the use of this cytoprotective gas should be investigated further as a novel treatment strategy in CKD. A diet rich in one-carbon methyl donor units relative to calories, such as betaine (found in fruits, cereals and vegetables), can be used as an epigenetic switch and, via DNA hypermethylation and transmethylation in the methionine cycle, can promote longevity¹⁶⁷. This mechanism merits further investigation, as low betaine levels have been observed in humans with poor renal function and accelerated biological ageing (P.G.S., unpublished observations).

Hypoxia and ischaemia

Naked mole rats survive constant exposure to hypoxic conditions by generating ATP through glycolysis. This process is mediated in part by using endogenously produced fructose, which preferentially stimulates glycolysis

and lactate production¹⁶⁸. We have found that fructose metabolism commonly leads to glycogen accumulation in the liver in mice and rats (R.J.J. and M.L., unpublished observations). This metabolic mechanism might also protect the kidneys of diving marine mammals that are subject to periods of prolonged hypoxia during deep dives. Harbour seals (*Phoca vitulina*) and whales (Cetacea), for example, have large amounts of glycogen in their proximal tubules, along with high levels of glycolytic enzymes to generate ATP during hypoxia^{169,170}. The kidneys of Weddell seals (*Leptonychotes weddellii*) are protected from hypoxia despite severe renal vasoconstriction upon diving⁶. Likewise, kidneys of hibernating squirrels are protected from ischaemic injury, in a process that is probably mediated via an absence of caspase-3-like mediated activity¹⁷¹.

One potential mechanism by which glycolysis protects against hypoxia could occur through the upregulation of antioxidants. Fasting seals have high expression levels of NRF2 (REF. 172), antioxidant enzymes¹⁷³ and glutathione levels¹⁷⁴ compared with non-fasting state levels. In support of a protective role for antioxidants, hyperactivation of NRF2 prevents progression of tubular damage after renal ischaemic injury in mice¹⁷⁵. Thus, NRF2 might be a therapeutic target to prevent acute kidney injury and could activate a hypoxia survival pathway (BOX 1). By contrast, very high intracellular concentrations of dietary or endogenously produced fructose lead to rapid and transient ATP depletion, resulting in a strong pro-inflammatory response and substantial oxidative stress in human proximal tubular cells¹⁷⁶. Indeed, a high-fructose diet inhibits KEAP1–NRF2 antioxidant signalling and increases the risk of non-alcoholic hepatosteatosis in mice¹⁷⁷. Thus, high concentrations of fructose may be injurious, whereas low concentrations may carry survival functions.

Lastly, although hypoxia and ischaemia usually occur in conjunction, some species, such as the turtle, are tolerant to hypoxia but still sensitive to brain ischaemia¹⁷⁸. A better understanding of hypoxic tolerance in turtles and naked mole rats may provide novel therapeutic interventions to combat the harmful effects of cerebral, renal and cardiac ischaemia in humans.

Seasonal acclimatization and hibernation

Seasonal acclimatization of metabolic activity

Many small mammals escape food shortage during winter by hibernation or daily torpor¹⁷⁹. Other species that do not hibernate or go into daily torpor in the classical sense, such as red deer (*Cervus elaphus*) or Alpine ibex (*Capra ibex*), adopt a similar hypometabolic state during winter. The reduction in energy expenditure is, therefore, similarly to hibernators and species undergoing daily torpor, mainly accomplished by lowering endogenous heat production and increasing the tolerance to a lower body temperature. Although the 2–3 °C change in core body temperature is only moderate^{180,181}, a substantially lower body temperature down to 15 °C is present at the body's periphery¹⁸², which is indicative of a substantial reduction in the mean temperature of the entire body mass. The winter phenotype of mammals

One-carbon methyl donor units

DNA methylation influences the expression of some genes and depends upon the availability of methyl groups. Dietary methyl groups are derived from food sources that contain methionine, one-carbon units, choline or betaine (a choline metabolite).

Torpor

A state of reduced body temperature and metabolic rate in animals that enables them to survive periods of reduced food availability.

further includes a shift from an anabolic metabolism during summer to the use of body fat reserves to fuel metabolism during winter^{183,184}. As a result, many hibernators do not eat during winter¹⁸³, and non-hibernating species such as red deer reduce their food intake substantially, even when fed *ad libitum*¹⁸⁵. The endogenous nature of the seasonal cycle of appetite and its entrainment by photoperiod have been shown experimentally for many wild species¹⁸⁶. Decreased food intake during winter leads to a reduction in the size of the gut and visceral organs, such as the kidney^{185,187}, which further contributes to lower energy expenditure.

Major differences exist in the levels of serum biomarkers of microbiota metabolites between wild bears and bears in captivity (P.G.S. and J.P., unpublished observations), suggesting that nutrients and feeding patterns contribute to the metabolic changes required for hibernation. A transition in energy metabolism from carbohydrates during summer to lipids during winter is facilitated by a switch from insulin sensitivity in the summer to insulin resistance during hibernation¹⁸⁸. The observation that central administration of leptin to captive grizzly bears leads to reduced food intake in October but not in August implies that seasonal variations exist in the sensitivity of the bear brain to the anorexic effects of leptin¹⁸⁸. In addition, seasonal variations in gut microbiota might also contribute to changes in energy metabolism in hibernating bears, as transplantation of summer gut microbiota from wild bears promoted adiposity without affecting glucose tolerance in germ-free mice¹⁸⁹. Although humans do not hibernate, investigating the processes that trigger fat accumulation in the summer followed by the switch to reduced energy intake and a fat-burning state occurring immediately before animals hibernate¹⁹⁰ may help to understand the mechanisms driving obesity.

Seasonal changes in membrane composition

Seasonal variation in body temperature is preceded by changes in the composition of cellular membranes, which consists of the integration of nutritionally acquired polyunsaturated fatty acids (PUFAs) into phospholipids during periods of cold acclimatization¹⁹¹. In addition to seasonal changes, even daily rhythmic changes in the phospholipid fatty acid composition of membranes have been found in humans along with changes in body temperature¹⁹². Furthermore, physical exercise can alter the lipid composition of membranes, for example, by increasing the concentration of docosahexaenoic acid (DHA) in skeletal muscle phospholipids¹⁹³ and by enhancing insulin sensitivity, probably through increasing insulin receptor expression levels. Of interest, the composition of membrane phospholipids has also been reported to contribute to the outstanding longevity in naked mole rats¹⁹⁴.

The composition of membrane lipids influences the activity of membrane-bound enzymes, for example, sarcoplasmic/endoplasmic reticulum Ca²⁺-ATPase (SERCA) activity is increased in membranes that are rich in linoleic acid. Therefore, incorporation of linoleic acid into phospholipids of cardiac myocytes can compensate for

reduced SERCA activity owing to low temperatures and enables adequate Ca²⁺ handling in cardiac myocytes even at body temperatures close to freezing point^{191,195}. High concentrations of linoleic acid in membranes also improve muscle performance at a high body temperature, as suggested by the positive relationship between linoleic acid content of membrane phospholipids of muscle cells and maximum running speed found in a comparative study of 36 mammal species¹⁹⁶. By contrast, DHA incorporation into phospholipids decreases SERCA activity¹⁹⁵ but seems to increase the activity of key enzymes of the Krebs cycle and fatty acid β -oxidation¹⁹⁷. Accordingly, an increase of the DHA content into phospholipids during hibernation in Alpine marmots is paralleled by an increase of thermogenic capacity¹⁹¹.

As all PUFAs are of dietary origin in mammals and birds, uptake influences the fatty acid composition of membranes. However, the balance of ω -6 to ω -3 PUFA in phospholipids seems to be regulated more by deacylation–reacylation processes, that is, membrane remodelling, rather than directly by dietary intake^{185,198}. The different effects of ω -6 and ω -3 PUFA on membrane-bound enzymes hint at intriguing molecular conflicts. There probably is no optimal all-purpose PUFA composition in tissues, which creates a trade-off between costs and benefits of each fatty acid that is influenced by metabolic state, for example, fasting or fattening, and hence is subject to seasonal and even daily variations¹⁸⁵.

In rats with adenine-induced CKD, a pro-inflammatory fatty acid pattern (low PUFA and high saturated fatty acid concentrations) was associated with downregulation of NRF2 activity and increased activation of NF- κ B and its downstream cytoprotective and antioxidant proteins¹⁹⁹. As oxidation of eicosapentaenoic acid and DHA generate concentrations high enough to induce NRF2-directed gene expression²⁰⁰; this may explain the antioxidative and anti-inflammatory properties of ω -3 PUFAs. Burmese pythons (*Python molurus*) were reported to display 40% cardiac hypertrophy with increased cardiac output 48–72 hours after large meals²⁰¹. Because consumption of these meals activates expression of fatty acid transport pathways and cardio-protective enzymes, and because injection of a combination of python fatty acids found in plasma promotes physiological hypertrophy in mammalian cardiomyocytes²⁰², targeted fatty acid supplementation may be a novel strategy to modulate cardiac gene expression and function in heart failure.

Circadian clock and kidney functions

Oscillating molecules that regulate circadian clocks are common in most if not all animal species²⁰³. Disruptions of the circadian clock lead to metabolic syndrome with dyslipidaemia, hyperleptinaemia, hyperglycaemia and hepatic steatosis in *Clock*^{-/-} mice²⁰⁴. As the circadian clock activates NRF2–glutathione-mediated antioxidant defence pathways and as arrhythmic *Clock* ^{Δ 19} mice have low NRF2 expression²⁰⁵, this network might have an important role in regulating energy balance and antioxidative protection. Circadian fluctuations are also known to affect renal blood flow, glomerular filtration,

Circadian clock

The circadian clock regulates the internal and external activities of organisms, such as sleep and changes in metabolism, based on the day–night cycle.

blood pressure and water and sodium excretion²⁰⁶. Thus, whether CKD progression is affected by circadian disruption and the potential benefits of chronotherapy should be investigated²⁰⁷. In addition, further investigations are warranted into the reasons for seasonal variations in the incidence, progression and mortality of ESRD²⁰⁸.

Insights from hibernating bears

Osteoporosis, poor wound healing, vascular disease, inflammation and muscle loss, together with substantial metabolic dysfunction (FIG. 1), are common features of the uraemic phenotype¹⁰. The metabolism of bears is suppressed to approximately 25% of basal rates during hibernation¹⁹⁰. Nevertheless, hibernating bears tolerate extended periods of an extremely low heart rate (~10 beats/min)²⁰⁹ without developing congestive heart failure, atherosclerosis²¹⁰ thromboembolic events or cardiac dilation, which are common features in CKD. The protection against vascular disease may in part be mediated by changes in the coagulation pathway, in which traditionally intrinsic cascades (initiated when blood comes in contact with exposed collagen from damaged endothelial cells) are suppressed and extrinsic tissue factor pathways (initiated by vascular wall trauma) are maintained, to prevent thromboembolic events while enabling external injuries to be healed²¹¹.

The fact that hibernating bears do not develop azotaemia or uraemic complications⁸ is remarkable, considering that they have a 90% reduction in renal blood flow, anuria (70–180 ml of urine per day is reabsorbed through the urinary bladder wall²¹²), mild hypothermia (30–36 °C) and a 50–70% reduction in GFR, and experience fasting and immobilization for 5–6 months of winter sleep. It is even more intriguing that females are able to give birth to cubs and nurse them during hibernation. Although a histological study in Romanian brown bears reported signs of glomerular fibrosis after awakening from winter sleep²¹³, the reduced renal function was normalized within weeks⁸. Thus, studies of the profound metabolic changes that occur in bears from summer to winter may provide clues that point towards novel therapeutic strategies for patients with CKD⁷ (FIG. 1). Bears and marine mammals have a reniculated kidney system (renal lobulation) (Supplementary information S5 (figure)). Proximal convoluted tubules in multilobulated and reniculated kidneys are comparatively short, which decreases the resistance to intraluminal flow. The large body size in combination with the limitation of length of the proximal convoluted tubules²¹⁴ seems to be the most likely explanation for multilobulation of large terrestrial and marine mammalian kidneys. Similar to the protection from hypoxia during deep dives of seals, bears might benefit from reniculated kidneys during hibernation when their blood flow is reduced.

Applications for transplantation. Kidneys are particularly susceptible to ischaemic injury because of their high metabolic rate and oxygen consumption. Ischaemia–reperfusion injury is common in donor organs used for renal transplantation, in part owing to mitochondrial

dysfunction, oxidative stress, ATP depletion and apoptosis following rewarming of the donor kidney. Despite extensive and repetitive periods of low metabolism, starvation and low cardiac output²¹⁵, bears return from hibernation without signs of persisting organ damage. Hence, studying the molecular changes in hibernating bears may lead to novel pharmacological approaches that could mimic hibernation and limit organ damage during renal transplantation²¹⁵. As active suppression of metabolism during hibernation precedes the lowering of the body temperature, it can be speculated that lowering the basal metabolic rate may be more effective at preventing ischaemia–reperfusion injury to the donor organ than would therapeutic hypothermia¹⁹⁰. Intriguingly, the metabolic switches that occur in preparation for hibernation share features with the metabolic changes associated with longevity in the animal kingdom²¹⁶. Indeed, hibernating species have an approximately 15% higher annual survival rate than non-hibernators of similar size²¹⁷. The observation that animals initiate hibernation owing to a lack of food (or other environmental cues) and not because of a lower body temperature²¹⁸ and terminate hibernation owing to physiological factors²⁰⁹ can guide future research on the metabolic switches that induce and terminate hibernation.

On the basis of metabolic pathways that are altered in hibernation and associated with longevity, approaches that might preserve organ function during transplantation could be proposed: for example, the cytoprotective gas H₂S induces a torpor-like state in mice²¹⁹, protects against lethal hypoxia²²⁰ and, as mentioned earlier, activates anti-inflammatory and antioxidant pathways via mTOR and NRF2 (REF. 221). Thus, H₂S treatment might confer organ cytoprotection via creation of a hibernation-like environment²²². Furthermore, injection of the AMPK agonist 5'-AMP induces torpor independently of H₂S²²³, although the mechanism underpinning this observation remains to be defined. Therefore, pretreatment of donor organs with agents that inhibit inflammatory responses and activate antioxidant pathways, such as H₂S gas, sirtuin agonists, mTOR inhibitors and AMPK agonists, might prevent renal ischaemia–reperfusion injury more effectively than current approaches^{224,225}.

Applications for muscle wasting. The loss of skeletal muscle mass that can occur in patients with CKD is caused by a combination of sedentary behaviour, anorexia and the activation of catabolic pathways in the uraemic milieu. In contrast to humans, whose muscle mass and strength may be reduced by >90% during extended periods of immobilization, hibernating black bears show minor loss in skeletal muscle cell number or size²²⁶. One mechanism by which bears retain muscle strength is by *de novo* amino acid and protein synthesis from urea²²⁷, coupled with a unique ability to recycle urea during hibernation that has not yet been observed in other hibernating animals⁷. Metabolic recycling of nitrogenous waste products seems to be a conceivable mechanism to prevent loss of muscle protein (FIG. 6). In addition, the skeletal muscle of hibernating bears seems more resistant to denervation than skeletal muscle of non-hibernating

Chronotherapy

The science of timing drugs according to the circadian clock. This approach is used in various clinical conditions, such as cancer, hypertension, seasonal affective disorder and bipolar disorder.

Renal lobulation

Carnivores and most small mammals have smooth-surfaced and uni-pyramidal kidneys, whereas primates and Suidae (hogs and pigs) have a smooth-surfaced and multi-pyramidal kidney system. Large terrestrial mammals have multi-lobulated and multi-pyramidal kidneys to keep the proximal convoluted tubules short. Most marine mammals and bears have each lobe separated into renules (reniculated kidney system).

Therapeutic hypothermia

(also known as targeted temperature management). The induction of mild hypothermia (32–35 °C) after cardiac arrest for neuroprotection.

Sedentary behaviour

A type of behaviour that is characterized by an energy expenditure ≤ 1.5 metabolic equivalents while in a lying, reclining or sitting posture. Typical sedentary behaviours include watching TV, computer work, driving and reading.

Denervation

Loss of nerve supply to a part of the body, which can be due to multiple causes, such as surgery, physical injury, chemical toxicity or diseases.

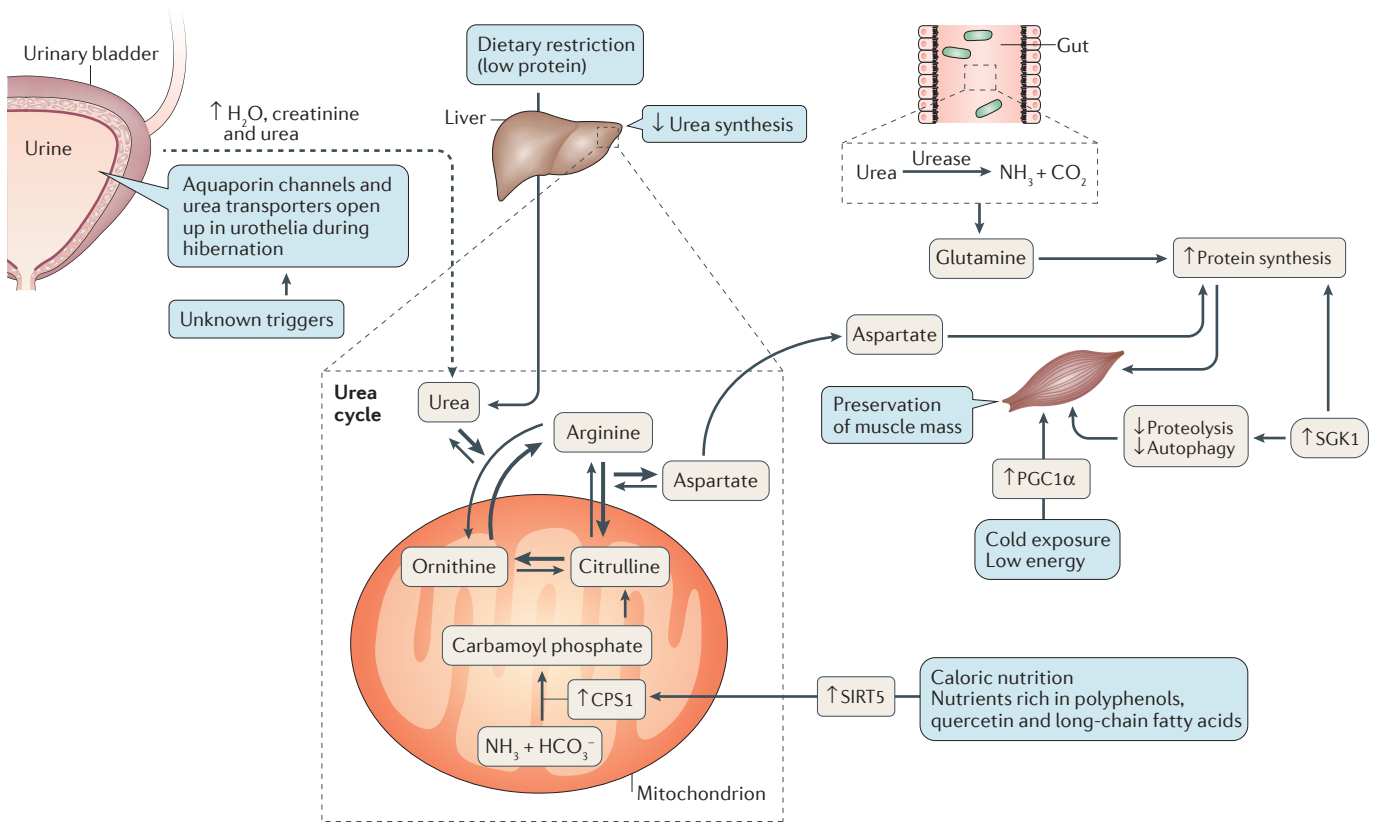


Figure 6 | Nitrogen metabolism in hibernating bears. To conserve mobility and muscle strength, hibernating bears must minimize muscle protein loss and re-utilize the vast majority of urea produced, which is mediated by microbial ureolysis and urea-N resorption. Multiple mechanisms are responsible for the reduction in serum urea levels during hibernation. Lower urea production during hibernation leads to reduced amino acid degradation. Moreover, urea is reabsorbed from urine via solute and water channels, such as urea transporters and aquaporin channels, in a leaky bladder wall. The reabsorbed urea is believed to be recycled back into skeletal muscle. Urea is also hydrolysed by urease-expressing gut bacteria into ammonia (NH_3), which is used by the host to synthesize glutamine for incorporation into proteins. Other factors that may prevent muscle loss in hibernating animals include activation of peroxisome proliferator-activated receptor γ -co-activator 1 α (PGC1 α), for example, by cold environmental temperature and the low-energy state, and serum-glucocorticoid-regulated kinase 1 (SGK1). Urea levels decrease in the autumn when food is still available, and the metabolic changes that determine urea metabolism may already occur before the bear enters hibernation. Because sirtuin (SIRT) stimulators, such as polyphenols in berries and plants, stimulate carbamoyl phosphate synthase 1 (CPS1), which is the first and rate-limiting step of the urea cycle, this may decrease urea generation and prepare the animal for low urine output during hibernation. Because urea recycling in bears has been insufficiently studied, the proposed pathways are mainly speculative. CO_2 , carbon dioxide; HCO_3^- , bicarbonate ion; H_2O , water.

Disuse atrophy

A type of muscle atrophy that occurs when a muscle is less active than usual. Disuse atrophy is a common feature in chronic debilitating diseases and immobility.

Mechanical unloading

A mechanical manoeuvre or therapy that decreases tissue growth and regeneration. Whereas mechanical loading of mammalian tissues is a potent promoter of tissue growth and regeneration, mechanical unloading in microgravity causes reduced tissue regeneration via stem cell tissue progenitors.

bears²²⁸, suggesting that hibernation is associated with changes in the neural regulation of skeletal muscle catabolic pathways and that targeting these pathways could offer novel solutions for the treatment of disuse atrophy.

The plasma of hibernating bears has an anti-proteolytic effect that inhibits wasting of isolated skeletal muscle²²⁹. Serum-glucocorticoid-regulated kinase 1 (SGK1) is activated by insulin and growth factors and helps to prevent loss of muscle mass via downregulation of proteolysis and autophagy and increased protein synthesis²³⁰. As high SGK1 expression levels have been reported in hibernating ground squirrels²³¹, mice lacking SGK1 have muscle atrophy²³⁰ and low SGK1 expression levels are found in patients with CKD²³², this serine/threonine kinase may be a novel therapeutic target to prevent uraemic muscle loss. Moreover, expression levels of peroxisome proliferator-activated receptor

γ -co-activator 1 α (PGC1 α), which activates metabolic pathways associated with endurance exercise (such as running), are induced by cold exposure²³³ and are elevated in hibernating squirrels²²². This master regulator plays a major part in renal recovery from acute kidney injury through regulation of NADH synthesis²³⁴. Hence, stimulation of PGC1 α , such as through exercise²³⁵, might also promote skeletal muscle homeostasis in CKD. Because activation of NRF2 by sulforaphane also increases endurance exercise capacity²³⁶, multiple targets and pathways to prevent uraemic muscle loss exist.

Applications for bone loss. In addition to being protected from muscle wasting, hibernating bears are protected from poor wound healing and osteoporosis. Unlike humans and other mammals, hibernating bears can withstand physical inactivity (mechanical unloading)

and nutritional deprivation for ≤ 6 months without any negative effects on bone strength²³⁷. Maintenance of calcium homeostasis is considered the most important contributing factor in bone health, but many other factors, such as growth hormones and cytokines, also have a role. Hibernating bears maintain eucalcaemia during immobilization⁸ and have decreased markers of bone resorption and formation²³⁸, which indicates precise balancing of bone remodelling activity. The suppression of bone remodelling during hibernation is likely an important mechanism to conserve energy during a long period of inactivity, decreased renal function and fasting¹⁹⁹. Other contributing factors probably include the differential regulation of gene expression and hypothalamic control of hormones involved in bone remodelling, as higher expression levels of hormones that reduce bone formation, such as cocaine- and amphetamine-regulated transcript protein (CARTPT)²³⁸. An elevated expression of anabolic genes but not bone resorption genes²³⁹ has also been reported.

Changes in vitamin D metabolism may also preserve bone mass during hibernation²⁴⁰. In contrast to humans, 25(OH)D vitamin levels do not change between seasons in bears²⁴¹, and bear kidneys continue to produce calcitriol (1,25(OH)₂D₃; the active metabolite of vitamin D) despite a marked reduction in renal function during hibernation²⁴⁰. Black bear parathyroid hormone activates cAMP, mitigates apoptosis in osteoblast cultures and increases trabecular bone volume²⁴²; hence, the anabolic effects of bear parathyroid hormone might also prevent disuse osteoporosis. In addition, NRF2 was reported to have a role in bone microarchitecture in a mouse model of osteoporosis²⁴³ and inhibited tumour necrosis factor ligand superfamily member 11 (TNFSF11)-mediated osteoclastogenesis in osteolysis-induced mice²⁴⁴. Given that increased NRF2 expression plays a major part in the antioxidant defences that are required for hibernation success in ground squirrels²⁴⁵, the potential role of NRF2 in maintaining skeletal mass in hibernating bears warrants investigation. Taken together, studies of hibernating bears can provide novel therapeutic approaches for the treatment of intracellular calcium disorders and prevention of bone loss during immobilization in humans.

Applications for wound healing. Bears also have the ability to heal wounds despite immobilization, hypothermia and anuria — conditions that are usually unfavourable for wound healing²⁴⁶. Elevated levels of δ -opioid receptor agonists and ursodeoxycholic acid have been linked to the wound-healing capabilities of hibernating bears²⁴⁶, but further insights into the underlying mechanisms involved might provide strategies to enhance wound healing. Changes in the coagulation pathways that occur during hibernation²¹¹ may also contribute to better wound healing.

Applications for azotaemia. The unique ability of hibernating bears to recycle urea back into proteins protects the bear not only from muscle wasting but also from azotaemia (FIG. 5). Because little urea is generated during hibernation²⁴⁷, minimal amounts of urine need to

be excreted²²⁷. When ¹⁴C urea and heavy water (D₂O) were administered into the bladder of hibernating bears, reabsorption of both isotopes occurred across urothelia with rapid appearance in plasma²⁴⁸. Although small quantities of solute and water transport across urinary tract urothelia are features of most mammalian species²⁴⁹, the mechanism(s) by which bears accomplish this transport during hibernation remains unknown²⁴⁹. One hypothesis is that the passage of recycled urea from the intestine contributes to *de novo* amino acid synthesis, as urease-expressing gut bacteria release ammonia that can be used by the host to synthesize glutamine for protein synthesis²⁴⁷. In contrast to bears, humans cannot recycle urea, and urea degradation in the human gut does not stimulate the conservation of nitrogen²⁵⁰. In addition, glycerol prevents azotaemia in hibernating black bears by serving as a carbon source for *de novo* amino acid synthesis²⁵¹. As urea levels decrease in the autumn before hibernation²⁵², a dietary shift may contribute to this change. The metabolic regulation of fasting is, in part, mediated by the activities of sirtuin 5 (SIRT5). SIRT5 exhibits deacetylase, desuccinylase and demalonylase activities and regulates the urea cycle enzyme carbamoyl-phosphate synthase 1 (CPS1) in liver mitochondria. Because *Sirt5*^{-/-} mice fail to upregulate CPS1 and exhibit hyperammonaemia during fasting²⁵³, this implies a role for SIRT5 in urea metabolism and the metabolic regulation of fasting. Thus, the long-term effects of sirtuin activators, such as resveratrol, on urea handling should be tested in patients with CKD.

Protective compounds in berries. Bears can ingest up to 200,000 berries per day in peak season, which occurs in late summer²⁵⁴. There is a synchronous timing of food resources that triggers the switch from salmon to berries during the summer²⁵⁵. Blueberries have potent anti-inflammatory and antioxidant properties (for example, through the actions of phenol-like antioxidants) and contain anthocyanins. Berries are also an important source of sirtuin agonists (such as pterostilbene and resveratrol), quercetin, vitamin K, vitamin C and fibres. In addition, berries contain fructose and linoleic acid that may stimulate fat storage in preparation for hibernation. Polyphenols are secondary metabolites in plants that are needed not only for plant growth but also as a defence mechanism against UVB exposure and aggression by insects and fungal pathogens²⁵⁶. In a mouse model of polygenic obesity, consumption of berries results in a shift in gut microbiota towards obligate anaerobes, which correlates with a decrease in gastrointestinal luminal oxygen and oxidative stress²⁵⁷. Potential implications on human health of the nearly anoxic conditions observed in the mouse gut lumen after berry consumption should be investigated.

In addition, resveratrol preserves bone mass in old male rats²⁵⁸, anthocyanins in berries increase serum alkaline phosphatase levels in obese male mice²⁵⁹, and the anthocyanin delphinidin inhibits excessive osteoclastogenesis in a mouse model of osteoporosis²⁶⁰. Notably, delphinidin also prevents muscle atrophy in mice²⁶¹ and lowers fasting glycaemia in individuals with prediabetes²⁶². Moreover, dietary supplementation

Eucalcaemia

The maintenance of normal and constant serum calcium levels.

Blueberries

Blueberries comprise all blue-coloured berries of the *Vaccinium* genus, of which the most common is bilberries. Blueberries have a low glycaemic index and are a rich source of fibres, vitamin K, manganese, > 15 different anthocyanins (especially delphinidin and malvidin), quercetin, myricetin and resveratrol.

Anthocyanins

Anthocyanins (> 600 molecular structures) belong to a class of molecules called flavonoids that are universal plant colourants responsible for the red, purple and blue colours in many fruits, berries, vegetables and flowers. Due to their contribution in multiple physiological activities, the consumption of these molecules is believed to have a substantial role in preventing lifestyle-related diseases.

Senolytic effects

Senolytic compounds selectively induce the death of senescent cells.

with anthocyanins isolated from roselle (*Hibiscus sabdariffa*) attenuated progression of adenine-induced CKD in rats²⁶³. Because a causal role for senescent cells in ageing-related bone loss has been demonstrated in mice²⁶⁴, nutritional compounds with senolytic effects, such as quercetin and fisetin²⁶⁵ — found in fruits, capers, vegetables and berries — may also contribute to the capacity of bears to maintain their bone mass²³⁷. Quercetin also blocks phosphate-induced apoptosis and VSMC calcification via inhibition of mitochondrial fission and oxidative stress²⁶⁶. As polyphenols that stimulate sirtuins and PGC1 α prevent muscle wasting induced by mechanical unloading²⁶⁷ or in streptozotocin-induced diabetes in rats²⁶⁸, their long-term effects should be tested in patients with CKD who experience muscle wasting. Resveratrol and grape seed proanthocyanidin extract facilitate vascular endothelial growth factor A (VEGFA) expression and angiogenesis in different wound models²⁶⁹. Hence, the long-term effects of sirtuin activators and anthocyanins on wound healing require further investigation. In healthy humans and patients with the metabolic syndrome, blueberry supplementation decreases cardiovascular risk factors²⁷⁰, increases HDL cholesterol²⁷¹ and improves insulin sensitivity²⁷². Moreover, a study based on validated food-frequency questionnaires in 93,600 women (Nurses Health Study) showed that a high intake of anthocyanins (highest versus lowest quintile) was associated with a decreased risk of myocardial infarction²⁷³. A diet rich in fermentable carbohydrates, such as berries, has been reported to have arisen in a basal ursine bear (*Protarctos abstrusus*) during the Pliocene

period (~2.6–5.3 million years ago), enabling bears to occupy cold habitats, accumulate fat and hibernate²⁷⁴. Finally, because many plants that are consumed by bears contain melatonin²⁷⁵, the effect of plants, such as tall fescue (*Festuca arundinacea*), on the metabolic changes that occur between seasons in hibernating bears should be investigated further. Taken together, the potential beneficial effects of berries and other plant nutrients on the uraemic phenotype should be assessed.

Conclusions

Species living in extreme habitats have acquired adaptive mutations through natural selection and epigenetic calibration to survive in challenging environments. Environmental factors and stressors — such as infections, toxins, starvation, climate change and psychosocial factors — have modified the epigenetic landscape throughout evolution to enable dynamically responsive changes in gene expression and associated biochemical networks to help mitigate the effects of these changes⁷³. Although humans have a common ancestry with mammals and share the same vulnerability to infections, environmental toxins and illnesses, most physicians have regarded animal diseases as different and of minor interest in the understanding of complex human diseases. In fact, almost all diseases that affect humans have an equivalent in the animal kingdom, although treatment options may differ. Thus, we propose a multidisciplinary approach to improve health care of patients with CKD by sharing new discoveries and tools from the fields of zoology, botany, ecology, veterinary medicine, anthropology and biology.

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Author contributions

P.S. and R.J.J. launched the idea of studying renal biomarkers. P.S., J.P., M.K., M.L., W.A., T.R., P.G.S. and R.J.J. researched the literature, discussed the content of the article and wrote the text. All authors reviewed or edited the article before submission.

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