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Metabolic Alterations at the Crossroad of Aging and Oncogenesis

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Abstract

Aging represents the major risk factor for cancer. Cancer and aging are characterized by a similar dysregulated metabolism consisting in upregulation of glycolysis and downmodulation of oxidative phosphorylation. In this respect, metabolic interventions can be viewed as promising strategies to promote longevity and to prevent or delay age-related disorders including cancer. In this review, we discuss the most promising metabolic approaches including chronic calorie restriction, periodic fasting/fasting-mimicking diets, and pharmacological interventions mimicking calorie restriction.

Metabolic interventions can also be viewed as adjuvant anticancer strategies to be combined to standard cancer therapy (chemotherapeutic agents, ionizing radiation, and drugs with specific molecular target), whose major limiting factors are represented by toxicity against healthy cells but also limited efficacy easily circumvented by tumor cells. In fact, conventional cancer therapy is unable to distinguish normal and cancerous cells and thus causes toxic side effects including secondary malignancies, cardiovascular and respiratory complications, endocrinopathies, and other chronic conditions, that resemble and, in some cases, accelerate the age-related disorders and profoundly affect the quality of life.

In this scenario, geroscience contributes to the understanding of the mechanisms of protection of normal cells against a cytotoxic agent and finding strategies focused on the preserving healthy cells while enhancing the efficacy of the treatment against malignant cells.

1. INTRODUCTION

Aging represents the major risk factor for cancer. In fact, the incidence of cancer increases with age due to accumulation of mutations. This association can be explained not only by the multihit or Knudson hypothesis, according to which tumor cells need to accumulate the mutations responsible for genome instability and carcinogenesis, but also by a decline of homeostasis occurring during aging (Knudson, 1971; Wu et al., 2014b). The discovery of the role of protooncogenes in accelerating aging and promoting cellular sensitization to stress

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has encouraged the investigation of metabolic interventions able to promote longevity but also useful in generating differential protection and sensitization of normal and cancer cells, respectively. In this review, we will discuss the most promising treatments including chronic calorie and protein restriction, fasting and fasting-mimicking diets (FMD), and pharmacological interventions mimicking calorie restriction (CR) relevant to both aging and cancer.

In addition to their antiaging effects, the above-mentioned strategies can also function as adjuvant approaches to be combined with standard cancer treatments in order to protect healthy cells and tissues and sensitize malignant cells to cytotoxic agents. In fact, one of the major limitations of cancer therapy, including chemotherapy, ionizing radiation, and target-specific drugs, is their inability or limited ability to distinguish normal and cancerous cells. This aspect is associated with the appearance of long-term and often serious side effects that not only impair the quality of life but may also accelerate the aging process. For this reason, geroscience, whose purpose is to understand the mechanisms of damage and protection during aging, could lead to novel interventions able to protect the host from the standard cancer treatment while enhancing its efficacy.

2. DYSREGULATION OF METABOLISM IN CANCER AND AGING

2.1 Mitochondrial Homeostasis in Cancer

Dysregulation of metabolism represents one of the hallmarks of tumorigenesis. In particular, cancer cells reprogram their carbon metabolism by upregulating glycolysis (Warburg effect) and glutaminolysis and by downregulating oxidative phosphorylation (OXPHOS). These events lead to the increase of biosynthetic intermediates such as nucleotides, amino acids, and lipids which are necessary for cell proliferation and survival (Vander Heiden et al., 2009). Moreover, disruption of mitochondrial homeostasis causes accumulation of reactive oxygen species (ROS) which can contribute to cancer progression (Hamanaka and Chandel, 2010).

In accordance with the normal function of their encoded proteins, oncogenes or tumor suppressors regulate cellular metabolism (Cairns et al., 2012). For example, Myc and phosphatidylinositol 3-kinase (PI3K) are potent inducers of glutaminolysis and glucose uptake, respectively (Elstrom et al., 2004; Wise et al., 2008), while the tumor suppressor p53 inhibits glucose transporters (GLUT), glycolysis, pentose phosphate pathway, de novo fatty acid synthesis, and enhances mitochondrial OXPHOS and tricarboxylic acid (TCA) cycle rate (Wang et al., 2014). A similar effect on the regulation of metabolism is caused by the tumor suppressor liver kinase B1 (LKB1) that is frequently mutated in Peutz-Jegher's syndrome, characterized by an increased risk of gastrointestinal cancer. LKB1 encodes for a serine-threonine kinase that activates the energetic sensor AMP-dependent kinase (AMPK) and downmodulates the regulator of growth and proliferation known as mammalian target of rapamycin (mTOR) (Shackelford and Shaw, 2009). Von Hippel Lindau disease is another example of disorders associated to cancers and caused by mutations in the *Vhl* gene, which inhibits the hypoxia-inducible factor α (HIF- α) (Kapitsinou and Haase, 2008; Maxwell et al., 1999). HIF- α is a transcriptional factor which suppresses oxidative metabolism and promotes glycolysis (Semenza, 2010).

In the last 10 years, several genetic defects affecting TCA enzymes associated to oncogenesis have been described. In particular, loss-of-function mutations or changes in the amino acid residues have been identified in the cytoplasmic and mitochondrial isoforms of the enzymes succinate dehydrogenase (SDH), isocitrate dehydrogenase (IDH), and fumarate hydratase (FH) (Cardaci and Ciriolo, 2012).

The SDH complex is an highly conserved heterotetrameric tumor suppressor, composed by two catalytic subunits (SDHA and SDHB) and two hydrophobic subunits (SDHC and SDHD) (Bardella et al., 2011), which have been found mutated in patients affected by different malignancies including hereditary paragangliomas, pheochromocytomas, gastrointestinal stromal, thyroid and renal tumors, neuroblastoma, and testicular seminoma (Astuti et al., 2001; Bardella et al., 2011; Baysal et al., 2000).

FH is a homotetrameric TCA cycle enzyme, which catalyzes the hydration of fumarate to L-malate. Heterozygous FH mutations predispose to multiple cutaneous and uterine leiomyomas, hereditary leiomyomatosis, renal cell cancer, breast, bladder, as well as Leydig cell tumors (Carvajal-Carmona et al., 2006; Lehtonen et al., 2006; Tomlinson et al., 2002).

IDH is a member of the β -decarboxylating dehydrogenase family, which catalyzes the oxidative decarboxylation of isocitrate to produce 2-oxoglutarate (α -KG). Three isoforms have been identified so far: the cytosolic IDH1, and the mitochondrial IDH2 and IDH3. Mutations associated to IDH1 and IDH2 have been identified in 70% of grade II–III gliomas, secondary glioblastomas, acute myeloid leukemia, angioimmunoblastic T-cell lymphomas, thyroid, colorectal, and prostate cancer (Abbas et al., 2010; Cairns et al., 2012; De Carli et al., 2009; Yen et al., 2010). As a result of gain-of-function mutations, IDH1 and IDH2 are unable to efficiently convert isocitrate into α -KG and acquire a neomorphic catalytic activity that allows a NADPH-dependent reduction of α -KG into the oncometabolite (*R*)-2-hydroxyglutaric acid ((*R*)-2HG) (Dang et al., 2010; Ward and Thompson, 2012). The mechanisms underlying tumorigenesis in cancers characterized by TCA cycle enzyme mutations involve the accumulation of metabolites (succinate, fumarate, and (*R*)-2-HG) that convey oncogenic signals (oncometabolites) (Yang et al., 2013). In particular, the abnormal accumulation of (*R*)-2HG mediates its potential tumorigenic effects via several mechanisms: (i) inhibition of ten-eleven translocation family (TET) of dioxygenases and histone lysine demethylase (KDM) which results into enhanced CpG island and histone methylation and the consequent remodeling of the cancer cell epigenome toward an undifferentiated and aggressive phenotype, (ii) inhibition of collagen prolyl and lysyl hydroxylases causing impaired collagen maturation and disrupted basement membrane formation, and (iii) inhibition of HIF- α prolyl hydroxylase (PHD) interactions causing a decrease of HIF-1 α degradation and an enhancement of pseudohypoxic condition. In addition, accumulation of fumarate and succinate participates in oncogenic signaling through: (i) modification of cysteine residues in proteins which confers a state of constitutive activation of the antioxidant defense pathway mediated by NF-E2-related factor (NRF2), that generates a reductive milieu and promotes cell proliferation; (ii) inhibition of the reactions involved in arginine and purine synthesis; (iii) epigenetic alterations by inhibition of TET and KDM proteins; and (iv) accumulation of HIF-1 α which in its turn promotes aerobic glycolysis and angiogenesis (Selak et al., 2005).

Furthermore, succinate has recently emerged as a key player in the promotion of inflammation which is functionally associated to cancer development and progression. In particular, proinflammatory macrophages shift their metabolism from OXPHOS to glycolysis resulting in succinate accumulation and oxidation by SDH, which drives ROS production that, finally, leads to increase of HIF-1 α and proinflammatory cytokines including IL-1 β (Mills et al., 2016).

A recent study highlighted the importance of the oncometabolite fumarate as a driver of tumorigenesis (Sciacovelli et al., 2016) by discovering that accumulation of fumarate, associated to FH loss, induces epithelial mesenchymal transition (EMT), a well-known process involved in cancer initiation, dissemination, and metastasis. Specifically, fumarate has been shown to inhibit TET-mediated demethylation of a regulatory region of the antimetastatic miRNA cluster mir-200ba429, leading to the expression of EMT-related transcription factors and enhanced migratory properties (Sciacovelli et al., 2016).

Beside the above-mentioned oncometabolites, ADP-ribose, which is synthesized by poly-ADP-ribose polymerases (PARPs) from NAD(+) and is responsible for protein posttranslational modifications, can be considered as an oncometabolite as well. A recent study demonstrated that nuclear pyruvate kinase M2 (PKM2) binds directly to ADP-ribose, and this poly-ADP-ribose-binding capability is critical for its nuclear localization. Accordingly, PARP inhibition prevents nuclear retention of PKM2 and therefore suppresses cell proliferation and tumor growth (Li et al., 2016).

The general mechanism underlying tumor metabolism dysregulation is still under investigation. The prevailing point of view is that the reprogramming of tumor metabolism (Warburg-like effect) occurs after cancer cells accumulate key mutations, promoting additional genome instability (Vander Heiden et al., 2009). According to alternative hypothesis, known as geroncogenesis, during age, normal cells undergo natural alterations in oxidative metabolism, with the consequent increased generation of ROS, which promote additional mutations and tumorigenesis. The latter hypothesis may help explain why aging is the major risk factor for most tumors (Wu et al., 2014b).

2.2 Geroncogenesis and Gerometabolites: The Pseudohypoxic Aging Side of Oncometabolites

Consistent with geroncogenesis, aging in mammals is characterized by metabolic alterations similar to those associated with cancer cells, and by a reduction of OXPHOS as a result of alteration of specific electron transport chain (ETC) complexes as well as an increase in aerobic glycolysis in different tissues (Bowling et al., 1993; Hagen et al., 1997; Trounce et al., 1989). For example, in humans, age leads to a decrease in cytochrome oxidase activity in brain and heart (Muller-Hocker, 1989; Ojaimi et al., 1999). Along this line, the activity of complex V decreases in the heart of the Fischer 344 rats and is accompanied by structural changes during aging. In contrast, complexes I and III activity remain unaltered in the heart, liver, and skeletal muscle of mice during aging (Kwong and Sohal, 1998, 2000). As a result of alterations, aging is associated with a higher production of ROS due to a decreased flux through the ETC. This event reduces the activity of upstream complexes, especially complexes I and III, enhancing “electron leak” that generates ROS (Chen and Lesnefsky,

2006). The increased production of ROS by mitochondria leads to greater oxidative damage within mitochondria, including protein sulfhydryl oxidation, lipid peroxidation, and mitochondrial (mt)DNA damage (Floyd et al., 2001; Van Remmen and Richardson, 2001). In support of the importance of mitochondrial-derived ROS to the aging process, it has been shown that interventions and mutations that prolong survival tend to decrease the production of ROS from mitochondria (Chen et al., 2007; de Cabo et al., 2014; Lagouge and Larsson, 2013).

Similar metabolic dysregulations have been observed in type 2 diabetes: an age-related disorder characterized by accumulation of HIF-1 α and a decline of OXPHOS (Petersen et al., 2004; Ptitsyn et al., 2006).

An additional possible link between aging and tumorigenesis is supported by the suggestive theory according to which the so-called “gerometabolites” (defined as small-molecule components of normal metabolism whose depletion drives aging) promote the chronic accumulation of oncometabolites which in turn are responsible for pathological metabolic reprogramming. In this scenario, nuclear NAD⁺, a central metabolic cofactor that decreases during aging (Verdin, 2015), plays a pivotal role. NAD⁺ is cosubstrate of sirtuins, a family of NAD⁺-dependent deacetylases that are regulators of metabolism in cancer and aging. Specifically, sirtuin 1 deacetylates HIF-1 α and promotes its interaction with the ubiquitin ligase VHL which leads to the degradation of HIF-1 α (Gomes et al., 2013; Haase, 2009). The decline of NAD⁺ during aging represents the causal inducer of the accumulation of HIF-1 α which promotes a pseudohypoxic state that reduces the carboxylation of α -ketoglutarate. This event leads to the production of the oncometabolite (*R*)-2HG (Gomes et al., 2013), that, together with the other oncometabolites fumarate and succinate, creates a pseudohypoxic state. The latter condition promotes the activation of c-Myc which is responsible for repressing the transcription of mitochondrial genes. As a consequence, a reprogramming of metabolism characterized by promotion of glycolysis and inhibition of OXPHOS occurs. This scenario, as depicted in Fig. 1, offers the possible connection between aging and cancer where the conjunction ring is represented by pseudohypoxia that is induced by gerometabolites. Pseudohypoxia is then associated with the generation of oncometabolites which block the differentiation and promote stemness through epigenetic mechanisms (Menendez et al., 2014). In this context, several reports indicate that hypoxia favors the increase in the cancer stem cell (CSC) pool through HIF-1 α and HIF-2 α which, in turn, amplify the CSC pool and induce additional dedifferentiation of tumor cells (Carnero and Leonart, 2016; Keith and Simon, 2007; Li et al., 2009). In particular, hypoxia and HIFs induce stemness in differentiated progenitors and non-CSCs by inducing the expression of genes such as *OCT4*, *SOX2*, and *NANOG*, or the activation of the Notch signaling pathway that regulates cell self-renewal and differentiation (Bennewith and Durand, 2004). The molecular mechanisms that underlie the impact of hypoxic conditions on the CSC compartment require either HIF-1 α expression or the inactivation of the hydroxylase activity of PHD domain-containing protein 3 (PHD3) whose loss prevents CSC differentiation and induces dedifferentiation of mature tumor cells (Iriando et al., 2015). Finally, HIF-1 α mediates telomerase transcription in hypoxic cancer cells, maintaining the immortal life span of the tumor mass. In fact, 90% of tumors show increased telomerase activity, suggesting that it is an important factor in the maintenance of CSC properties

(Blanco et al., 2007). HIF- α and pseudohypoxia have been also implicated in aging although in *Caenorhabditis elegans* this issue is a matter of debate because HIF- α has been shown to function as both a positive and negative modulator of aging (Leiser and Kaeberlein, 2010). There is also evidence that HIF- α plays a relevant role in mammalian aging. In particular, activation of HIF-1 α , which is accompanied by increased HIF-1 α DNA binding and activation of transcription of HIF-1 α -dependent genes, has been observed in aged rat liver (Kang et al., 2005). Another study examining PHD3 in rats found that PHD3 levels increase with age in liver, heart, and skeletal muscle, and this increase correlates with a decrease in HIF-1 α activity (Rohrbach et al., 2008). Finally, HIF-1 α seems to have a contradictory role in age-related disorders; in fact, studies examining A β accumulation in Alzheimer's disease suggest that increase of HIF-1 α tends to be protective in disease-free states, but that increased HIF-1 α levels are also a sign of advanced disease progression (Ogunshola and Antoniou, 2009). These seemingly conflicting results might be due to the fact that in stress response HIF-1 α upregulation can be either protective or reactive in nature (Ogunshola and Antoniou, 2009).

2.3 Sirtuins: Regulators of Metabolism of Cancer and Aging

Sirtuins are mono-ADP-ribosyltransferase and beta-nicotinamide adenine dinucleotide (NAD⁺)-dependent lysine deacetylase enzymes that play key roles in the regulation of metabolism, inflammation, and DNA repair and are critical regulators at the crossroads between cancer and aging (Chalkiadaki and Guarente, 2015; Saunders and Verdin, 2007). The mechanisms underlying the protective effect of sirtuins in aging and cancer include: (1) protection against DNA damage and oxidative stress and (2) protection against accumulation of mutations and genomic instability (Saunders and Verdin, 2007). In fact, the loss of sirtuin expression, activity, or regulation allows cell division to proceed without the proper repair of DNA, resulting in accumulation of mutations and genomic instability which can lead to tumor development. More recently, sirtuin proteins have also been found to finely regulate energy metabolism. In particular, sirtuin 3 facilitates TCA cycle, OXPHOS, and fatty acid metabolism (Hirschey et al., 2010). For example, deletion of *Sirtuin 3* causes spontaneous formation of mammary tumors and a metabolic reprogramming characterized by increased glucose uptake and decreased ATP generation (Kim et al., 2010). This metabolic switch is accompanied by an enhanced production of ROS that stabilize HIF-1 α , which in its turn promotes glycolysis and reduces OXPHOS (Bell et al., 2011; Finley et al., 2011). The role of sirtuin 3 as a tumor suppressor is however controversial since other studies indicated that sirtuin 3 is overexpressed in breast cancer compared to normal tissues (Alhazzazi et al., 2011; Ashraf et al., 2006).

Another sirtuin protein involved in tumor metabolism regulation is sirtuin 6, whose deletion is associated with an increase in HIF-1 α and c-Myc, leading to activation of glycolysis (Sebastian et al., 2012; Zhong et al., 2010). Similarly, sirtuin 1, which was the first family member identified as a tumor suppressor that delays lymphoma and protects mice against carcinogen-mediated hepatocellular carcinoma (Herranz et al., 2010; Oberdoerffer et al., 2008), regulates the transcriptional activity of HIF-1 α and activates LKB1 through deacetylation (Lan et al., 2008; Lim et al., 2010). Sirtuin 2 is another tumor suppressor involved in maintaining genome stability and suppressing tumorigenesis by deacetylating

and activating Cadherin-1 (CDH1), a protein that limits glycolysis (Almeida et al., 2010; Kim et al., 2011). More recently, sirtuin 4 and sirtuin 7 have also been implicated in tumorigenesis with an effect of sirtuin 4 in repressing the mitochondrial glutamine metabolism, and a role of sirtuin 7 in negatively regulating HIF-1 α and HIF-2 α (Hubbi et al., 2013; Jeong et al., 2013).

Although all the above studies indicate that sirtuins can function as tumor suppressors, their role in tumorigenesis is still controversial since different reports indicated that sirtuins, including sirtuin 1, sirtuin 2, and sirtuin 3, also promote cancer (Bosch-Presegue and Vaquero, 2011). In this regard, sirtuin 1 has been shown to inhibit p53 and to stabilize c-Myc, which may explain why its overexpression can increase tumor growth (Menssen et al., 2012; Suh et al., 2011; Villeda et al., 2011).

2.4 Nutrient-Sensing Pathways: A Common Signaling in Aging and Cancer

Several studies in yeast, worms, flies, and mice indicate that conserved nutrient-sensing pathways governed by insulin/insulin growth factor (IGF), glucose, and amino acids regulate aging and affect genomic integrity, DNA repair, ROS generation, and cellular apoptosis, highlighting their role in the promotion of cancer (Fontana et al., 2010; Longo and Fontana, 2010). As depicted in Fig. 2, a signaling pathway is triggered when a specific ligand binds to its cognate receptor. These ligands are mainly represented by glucose and amino acids in yeasts; Ins/IGF-1-like peptides in worms and flies; and insulin, IGF-1, and IGF-2 in mammals. In yeast, Ras is involved in the activation of adenylate cyclase (AC) which in its turn activates protein kinase A (PKA), leading to the phosphorylation and inactivation of transcription factors including MSN2/4 and GIS1. The cognate receptors include the insulin-like receptor DAF-2 in worms, InR in flies and IGF-1R, IR-A, and IR-B in mice. Upon the interaction between ligands and their receptors, the signal is transduced through adaptor proteins such as Ras in yeasts and mice, IST-1 in worms, CHICO in flies, and insulin receptor substrate 1 (IRS1–4) in mammals. These proteins are involved in the activation of PI3K which generates phosphatidyl inositol (3,4,5)-trisphosphate (PIP3). In animals, PIP3 activates AKT/PKB which phosphorylates and inactivates specific antiaging transcription factors (DAF-16 in worms, and Forkhead box protein O (FOXO) in flies and mice). This event causes the downregulation of protective stress resistance genes including superoxide dismutase (SOD), catalase, and heat shock proteins (HSPs) which contribute to the protection against oxidative damage contributing to aging as well as cancer. An additional pathway involved in the promotion of aging is controlled by mTOR and the serine kinase 6 (S6K). Dampening these nutrient-sensing pathways by specific mutations, as it occurs in the yeast *sch9* null mutants, CHICO mutant flies, Ames and Snell dwarf mice lacking growth hormone (GH) and IGF-1 as well as “Laron dwarf mice” produced by targeted disruption of the GH receptor/GH-binding protein gene (GHR-KO mice), significantly increases the life span (Longo and Finch, 2003).

Similar effects can be obtained by strategies based on dietary restriction, which not only extend longevity but also reduce or delay age-related disorders, particularly spontaneous tumors. The effects of these mutations on aging may be explained in part by the finding that fibroblasts from long-lived adult Ames or Snell dwarf mice and GH receptor knockout mice

are better protected against oxidative stress, suggesting that IGF-1 has a key role in regulating protection against toxins (Fontana et al., 2010). In agreement with the latter observation, individuals who carry mutations in the GH receptor gene that lead to severe GHR and IGF-1 deficiencies (Laron population) are protected against cancer and diabetes (Guevara-Aguirre et al., 2011). The explanation for these results is provided in part by in vitro studies in which mammary epithelial cells incubated with serum from the Laron individuals were shown to be more protected from DNA damage caused by hydrogen peroxide but are also more likely to die as a consequence of damage compared to those incubated with serum from unaffected relatives.

Nutrient signaling pathways play also a relevant role in the regulation of cell metabolism. Upon binding to its receptor, IGF-1/2 can activate Ras which, in turn, can mediate the activation of the PI3K/AKT and MAPK pathways, both of which increase glycolysis through different mechanisms. AKT increases the expression and membrane localization of the glucose transporter GLUT1, which stimulates phosphofructokinase activity and induces the translocation of hexokinases 1 and 2 into the mitochondria where they drive the first reaction of glycolysis (Barthel et al., 1999; DeBerardinis et al., 2008; Robey and Hay, 2006). Downstream of AKT, mTOR stimulates the switch to the glycolytic pathway through upregulation of HIF-1 α , protein translation, and lipogenesis by inducing the SRBPE-dependent transcription of different enzymes such as ATP citrate lyase and fatty acid synthase (Krycer et al., 2010; Yang et al., 2002). Interestingly, in an HIF-1 α - and Myc-dependent manner, mTOR upregulates the activity of PKM2, a crucial glycolytic enzyme involved in promoting the pro-Warburg effect (Sun et al., 2011). PKM2 interacts with HIF-1 α and enhances the expression of HIF-1 α target genes as well as it upregulates c-Myc transcription (Luo et al., 2011). In addition, mTOR as well as MAPK activate Myc whose overexpression causes the conversion of pyruvate into lactate thereby contributing to the Warburg effect (Dang et al., 2009). Myc is also involved in glutaminolysis induction through the upregulation of glutamate synthesis (Gao et al., 2009).

2.5 Inflammation and Cancer

One of the most typical feature and also a probable cause of aging are represented by a chronic and low level state of systemic inflammation in the absence of any infection that has been referred to as inflammaging (Franceschi and Campisi, 2014). Chronic inflammation increases cancer risk and promotes all cancer stages, including cancer initiation, progression, and metastatic diffusion (Mantovani et al., 2008). Thus, inflammation is likely to contribute to age-dependent cancer.

Inflammation can be derived from different sources including: (i) damaged macromolecules, organelles, and cells (self-debris) that accumulate with age and behave as “damage”-associated molecular targets that activate innate immunity and the Nlrp3 inflammasome; (ii) products derived from the microbial constituents of the human body, such as oral or gut microbiota, that elicit an inflammatory response (Biagi et al., 2011); (iii) senescent cells which accumulate with age and secrete proinflammatory cytokines; (iv) the coagulation system which is activated and increases during aging; (v) changes in immunosenescence; and (vi) defective or inappropriate regulation of the complement pathway that can lead to

local inflammatory reactions (Franceschi and Campisi, 2014). All of these stimuli can result in the activation of nuclear transcription factor NF- κ B which is considered as a hub in carcinogenesis, linking inflammation, cellular senescence, and cancer (Ostan et al., 2015).

3. METABOLIC INTERVENTIONS WITH EFFECTS ON AGING AND CANCER

Since nutrient-sensing pathways have been demonstrated to regulate longevity and to be modulated by dietary interventions, in the following sections, we will discuss different metabolic approaches demonstrated to have pro-longevity effects and to prevent or delay multiple age-related diseases and improve health span. These include: (1) CR, (2) periodic fasting (PF), (3) protein restriction, and (4) pharmacological interventions mimicking CR such as inhibitors of mTOR/S6K signaling, of glycolysis, and of GH or IGF-1 signaling, activators of sirtuins, and pharmacological inhibition of inflammation (Longo et al., 2015).

3.1 Calorie Restriction

CR is defined as a 20%–40% reduction in caloric intake without causing malnutrition, and while, in most cases, maintaining meal frequency. CR represents the most potent physiological intervention able to delay and prevent aging and age-associated diseases in single-celled, invertebrate, and vertebrate animals (Fontana and Partridge, 2015). In yeast, starvation obtained by switching the cells from a medium containing nutrients to one containing only water extends the chronological life span (Wei et al., 2008). Similarly, life span extension has been also observed in worms and flies under CR (Fontana et al., 2010). More than 80 years ago, different studies demonstrated that mice and rats under CR live longer and healthier in part by delaying the occurrence of several chronic diseases such as cancer, diabetes, atherosclerosis, and cardiomyopathy (Fontana et al., 2010). In term of cancer prevention, CR has been shown to inhibit the occurrence and growth of spontaneous, chemically induced, and radiation-induced tumors in various animal models (Cheney et al., 1983; Tannenbaum and Silverstone, 1949). Moreover, CR in mice greatly increases insulin sensitivity (Patel et al., 2005). Accordingly, young adult Rhesus monkeys under CR present a significant reduction of age-related pathologies including type 2 diabetes, cardiovascular disease, sarcopenia, and cancer (Colman et al., 2014; Mattison et al., 2012). In addition, immunosenescence and atrophy of the brain's gray matter are attenuated (Colman et al., 2014). In humans, long-term CR causes metabolic, molecular, and cellular adaptations which are responsible for its protective effect against type 2 diabetes, hypertension, cardiovascular disease, dementia, and cancer (Cava and Fontana, 2013). In particular, overweight women and elderly subjects under 3–4 months of CR present significantly improved cognitive function (Kretsch et al., 1997). The mechanism underlying the antiaging effect of CR include three levels of adaptations: (1) metabolic adaptations associated to a decrease of insulin, IGF-1, sex hormones, oxidative stress, inflammation, and to an increase of cortisol and adiponectin; (2) molecular adaptations which result in downregulation of PI3K/AKT/S6K, mTOR, RAS/MAPK pathways and upregulate NRF2, sirtuins, AMPK, FOXO, and PTEN; and (3) cellular adaptations including a decrease in cellular proliferation, oxidative stress, and enhancement of apoptosis, autophagy, DNA repair, genome stability, and immunosurveillance (Longo and Fontana, 2010). For example, CR-mediated inhibition of AKT activates FOXOs transcription factors that, in turn, activate protective systems

controlling cell proliferation, autophagy, stress resistance, and DNA repair (Webb and Brunet, 2014). Inhibition of mTORC1 increases autophagy and enhances stem cell function (Johnson et al., 2013; Kapahi et al., 2010). Moreover, genome stability and antioxidant defenses can be increased by the overexpression of certain sirtuins such as sirtuins 1, 3, and 6 and by the activation of heat shock factor 1 (HSF1) and of the NRF2 transcription factor (Akerfelt et al., 2010; Martin-Montalvo et al., 2011).

3.2 Protein Restriction

CR, describes the reduction of calories, rather than referring to both the quantity and composition of the diet, which research is now showing to be the major factor responsible of its health span effects in rodents. In fact, it has been observed that neither carbohydrate nor lipid restriction seems to have major effects on longevity alone. In contrast, several studies performed in yeast, worms, flies, and rodents point to protein restriction without a decrease of calorie as a nutritional modification able to increase the life and health span.

There is a clear evidence that the reduction of specific amino acids including serine, valine, threonine, asparagine, glutamate, or methionine extends the life span in yeast *Saccharomyces cerevisiae*, *Drosophila melanogaster*, and rodents (Ables et al., 2012; Miller et al., 2005; Richie et al., 1994; Wu et al., 2013). Particular interest has been focused on methionine restriction that exerts healthy effects by decreasing mitochondrial oxidative stress and consequently oxidative damage to mitochondrial DNA (Sanchez-Roman and Barja, 2013). The mechanisms underlying the effect of amino acids on aging are associated to their effect on the activation of mTOR and the control of the nonderepressible 2 (GCN2) gene. mTOR is modulated by different essential amino acids, mainly leucine, in a tissue-specific manner, while GCN2 is a serine/threonine-protein kinase that, once activated by amino acid deficiency, stabilizes the transcription factor ATF4 which is essential for the integrated stress response (Li et al., 2014). In yeast, methionine restriction promotes longevity in a GNC2-dependent manner (Wu et al., 2013), but other mechanisms including induction of autophagy (Ruckenstuhl et al., 2014) and mitochondrial retrograde response (Johnson and Johnson, 2014) have been proposed. The importance of other amino acids such as serine, valine, and threonine in promoting aging in yeast has been recently demonstrated (Mirisola et al., 2014). Specifically, threonine and valine promote cellular sensitization and aging primarily by activating the TOR/S6K pathway, while serine induces sensitization via phosphoinositide-dependent protein kinase 1 (PDK1) orthologs Pkh1/2. These events cause intracellular relocation and transcriptional inhibition of the stress resistance protein kinase Rim15 (Mirisola et al., 2014).

The extension of the health span in *D. melanogaster* is attributable to a restriction of protein-containing yeasts or sugar (Mair et al., 2005). According to these results, in mice, a low-protein and high-carbohydrate diet exerts life-extending effects through downregulation of the hepatic mTOR pathway (Solon-Biet et al., 2014).

Protein restriction has been also associated to decrease the prevalence and the severity of age-related diseases. For example, in mice, a low-protein and high-carbohydrate diet reduces the blood pressure, low-density lipoproteins, and triglycerides; improves glucose tolerance; and increases the levels of high-density lipoprotein (Solon-Biet et al., 2014). These results

are in agreement with data performed on human subjects in which high-protein and low-carbohydrate diets are associated with an increase of cardiovascular diseases (Floegel and Pischon, 2012; Lagiou et al., 2012). Thus, the balance of protein to carbohydrate, rather than energy intake, may be the driver of a healthy cardiometabolic profile.

The serum IGF-1 reduction and mTOR downregulation by an isocaloric restriction of protein has been also associated to a marked inhibition of prostate and breast cancer growth in experimental models (Fontana et al., 2013; Levine et al., 2014). In humans little is known about the effects of protein restricted diets on cancer. However, an interesting study reported that low-protein intake during middle age (between 50 and 65 years old) is associated with decreased risk of cancer and diabetes mortality as well as overall mortality. Conversely, low-protein intake is associated with increased risk of cancer and overall mortality in respondents over 65 (Levine et al., 2014). In agreement with the observation that an increased protein intake and the resulting increase in IGF-1 may prove beneficial for the skeletal muscle metabolism of older adults (Heaney et al., 1999), a very low-protein diet may be detrimental for older adults in whom weight begins to decline thus making them more susceptible to protein malnourishment (Levine et al., 2014).

3.3 Fasting and Fasting-Mimicking Diet

Fasting represents the most extreme version of CR where nutrients are totally eliminated. Two major forms of fasting can be practiced: (i) an intermittent fasting (IF) also known as alternate day fasting that involves a 24-h fast followed by a 24-h nonfasting period and (ii) prolonged and PF in which absence of food lasts two or more consecutive days every 2 or more weeks (Longo and Mattson, 2014). The first experiments performed on the yeast *S. cerevisiae* demonstrated that the shift from standard growth medium to only water causes a significant increase of life span as well as the resistance to multiple stress (Longo et al., 1997). This intervention is associated to downregulation of mTOR/S6K (Sch9) and RAS-AC-PKA pathways resulting in increased transcription of: (i) stress resistance genes such as SOD, catalase, and HSPs (Madia et al., 2009) and (ii) DNA repair genes including Rev1 (Wei et al., 2008) (Fig. 2). As a result, fasting exerts a protective effect against DNA damage and promotes longevity (Longo and Fontana, 2010). Similar observations have been made in the nematode *C. elegans* and *D. melanogaster* in which food deprivation increases the life span through the downmodulation of mTOR/S6K and IGF-1-like/PI3K/AKT pathways resulting in the activation of transcription factors DAF16 and FOXO (Fontana et al., 2010; Greer et al., 2007; Piper and Partridge, 2007) (Fig. 2).

In mice, IF reduces the incidence of spontaneous tumors including lymphoma and sarcoma (Berrigan et al., 2002; Descamps et al., 2005), while PF can delay cancer progression as effectively as chemotherapy (Lee et al., 2012). In addition, IF ameliorates age-related behavioral deficits in the triple-transgenic mouse model of Alzheimer's disease (Halagappa et al., 2007), reduces degeneration of dopaminergic neurons in models of Parkinson's disease (Duan and Mattson, 1999), and slows disease progression in huntingtin mutant mice by normalizing glucose metabolism and brain-derived neurotrophic factor levels (Duan et al., 2003). In agreement with the latter results, a recent study demonstrated that IF, when imposed in the middle age, delays or prevents the age-associated impairment of brain

functions and promotes healthy aging by improving the motor coordination and learning response recovery (Singh et al., 2012, 2015). Furthermore, IF protects the heart against ischemic injury in myocardial infarction models (Ahmet et al., 2005) and reduces the age-induced inflammation and fibrosis by inhibiting oxidative damage and NF- κ B activation (Castello et al., 2010). IF also decreases the diabetes incidence in BB rats (Pedersen et al., 1999) and prevents/reverses different aspects associated to metabolic syndrome by increasing insulin and leptin sensitivity, suppressing inflammation, and stimulating autophagy (Wan et al., 2010). Interestingly, it has been shown that the effects of IF on life span depend on genotype and age of initiation (Goodrick et al., 1990; Kendrick, 1973).

In contrast, the use of a FMD for 4 days every 2 weeks started at middle age extends health span, reduces tumor incidence, decreases inflammation, and delays the age-dependent cognitive decline as well as bone loss decline (Brandhorst et al., 2015). In this respect, severe dietary restriction or fasting have been hypothesized to exert beneficial effects in young and middle-age mice, but to be detrimental in older animals that begin to lose weight, according to what has been observed in humans (Brandhorst et al., 2015; Levine et al., 2014).

In humans, fasting has been shown to affect certain factors implicated in aging such as insulin and IGF-1, and to cause a fivefold increase of IGF-1 binding protein 1 (IGFBP1) which sequesters IGF-1 from the circulation thus reducing its bioavailability (Fontana et al., 2010; Harvie et al., 2011; Thissen et al., 1994). Notably, the effect of fasting on IGF-1 is due to the combination of protein and CR, but protein intake represents the major regulating factor (Thissen et al., 1994). One of the most evident effects of fasting in humans has been observed in patients affected by rheumatoid arthritis who have clinically significant beneficial long-term effects in term of reduction of pain and inflammation (Muller et al., 2001). However, when normal diet is resumed, most patients relapse unless fasting is followed by a vegetarian diet (Kjeldsen-Kragh et al., 1991). The beneficial effects of fasting have also been observed in subjects affected by hypertension with an improvement of systolic blood pressure below 120 after 13 days of water-only fasting (Goldhamer et al., 2002). Interestingly, a recent case report revealed that a medically supervised 21 days of water-only fasting followed by a diet of minimally processed plant foods free of added sugar, oil, and salt causes tumor regression in a woman affected by lymphoma (Goldhamer et al., 2015). Since fasting is often challenging for individuals affected by pathology, Brandhorst et al. developed a FMD to be administered periodically that is low in calories, protein, and sugar (Brandhorst et al., 2015) and which mimics the effects of fasting on markers associated with the stress resistance or longevity, including low levels of glucose and IGF-1 and high levels of ketone bodies and IGFBP-1 (Longo and Mattson, 2014). Bimonthly FMD cycles started at middle age extend the longevity of mice by lowering visceral fat, reducing cancer incidence and skin lesions, rejuvenating the immune system, and retarding bone mineral density loss. In old mice, FMD cycles promoted hippocampal neurogenesis, lowered IGF-1 levels and PKA activity, elevated NeuroD1, and improved cognitive performance. In a pilot clinical trial performed in generally healthy adults, three FMD cycles decreased risk factors/biomarkers for aging, diabetes, cardiovascular disease, and cancer without major adverse effects, providing support for the use of FMD to promote health span (Brandhorst et al., 2015). Moreover, preliminary data from a randomized pilot clinical trial conducted to assess the safety and feasibility of the FMD on patients affected

by multiple sclerosis showed that the FMD is safe, feasible, and potentially effective by reducing the number of autoimmune lymphocytes (Choi et al., 2016). Among fasting mimetic compounds, polyamine spermidine deserves particular attention since it has been demonstrated to: (i) markedly extend the life span of yeast, flies and worms, and human immune cells; (ii) decrease the oxidative stress; and (iii) exert cardioprotective effects by reducing cardiac hypertrophy and by preserving diastolic function in old mice (Eisenberg et al., 2009, 2016). Interestingly, autophagy, which is able to minimize the functional decline of aged cardiomyocytes by degrading and recycling long-lived damaged proteins as well as dysfunctional mitochondria (Nakai et al., 2007; Taneike et al., 2010), has been shown to be required for spermidine-mediated cardioprotection (Eisenberg et al., 2016). In agreement with the latter results, the dietary consumption of spermidine in humans correlates with reduced blood pressure and a lower incidence of cardiovascular disease (Eisenberg et al., 2016).

Of note, spermidine has been also implicated in cancer prevention since it is able to reduce tumor incidence through inhibition of age-associated alteration in DNA methylation status (Soda et al., 2013). Moreover, an elegant recent study demonstrated that spermidine as well as hydroxycitrate, both defined as CR mimetics, improves the inhibition of in vivo tumor growth by chemotherapy (Pietrocola et al., 2016). This effect was only observed for autophagy-competent tumors, depended on the presence of T lymphocytes, and was accompanied by the depletion of regulatory T cells from the tumor bed (Pietrocola et al., 2016).

3.4 Pharmacological Interventions Mimicking CR

3.4.1 Inhibitors of Nutrient-Sensing Pathways—Reduced nutrient-sensing pathways including mTOR/S6K, PI3K/AKT, Ras, or AC/PKA signaling, through genetic or pharmacological interventions lead to life span extension in yeast, worm, flies, and mice (Johnson et al., 2013). mTOR kinase is formed by two subunits called TORC1 and TORC2, where the former induces activation of S6K and 4E-BP1. The core components of mTORC1 consist of mTOR, mammalian lethal with sec-13 protein 8 (mLST8), and regulatory-associated protein of TOR (raptor), to which additional proteins including a DEP-domain-containing mTOR-interacting protein (DEPTOR) and proline-rich Akt substrate 40 kDa (PRAS40) are associated. The mTOR complex 2 (mTORC2) core is composed of mTOR, the rapamycin insensitive companion of mTOR (rictor), stress-activated protein kinase-interacting protein 1 (mSIN1), and mLST8. One of the best-known mTOR inhibitors is rapamycin, a pharmacological agent which was initially discovered as an antifungal metabolite and subsequently found to be immunosuppressive by inhibiting S6K1 activation (Chung et al., 1992). In the context of mTOR complex, only mTORC1, which integrates different signals governed by nutrients, growth factors, oxygen, and energy and activates anabolic processes required for cell proliferation and growth, is acutely sensitive to inhibition by rapamycin. Rapamycin binds to the intracellular protein FKBP12 to generate a complex that binds to and destabilized mTORC1 at all times after drug addition, consistent with the capacity of FKBP12-rapamycin to bind to it and weaken the raptor–mTOR interaction (Kim et al., 2002). In contrast, the modulation of mTORC2 by rapamycin is more complex and controversial. Since acute treatment with rapamycin does not perturb mTORC2

signaling and FKBP12-rapamycin cannot bind to intact mTORC2, this complex was originally thought to be rapamycin insensitive (Jacinto et al., 2004; Sarbassov et al., 2004). However, prolonged treatment with rapamycin has been shown to reduce mTORC2 signaling in some, but not all cell types and does so by suppressing mTORC2 assembly (Laplante and Sabatini, 2012; Sarbassov et al., 2006).

Rapamycin extends the life span in various model organisms including mice (Harrison et al., 2009; Miller et al., 2005). However, its serious side effects such as metabolic dysregulation (i.e., hyperglycemia, hyperinsulinemia, and insulin resistance) and blockade of hematopoietic cell lineage proliferation hampered its clinical use as an antiaging therapeutic (Soefje et al., 2011). Current clinical trials on healthy older subjects will help to understand whether inhibition of mTOR could be a safe and feasible antiaging intervention.

3.4.2 Inhibitors of Glycolysis—Consistent with the metabolic dysregulation associated with aging, i.e., reduction of OXPHOS and increase of glycolysis, there is currently great interest in applying pharmacological interventions aimed at inhibiting glucose catabolism and generation. Among these drugs, specific attention has been focused on 2-deoxy-D-glucose (2DG), a glucose analog which, once it is phosphorylated by hexokinase, cannot be further metabolized and consequently blocks glycolysis. Administration of 2DG to rats improves glucose and insulin regulation and increases recovery from stress (Minor et al., 2010). Another compound is an avocado extract called mannoheptulose that inhibits hexokinase and increases the life span in nematodes and mice (Minor et al., 2010). Acarbose, which limits glucose supply to cells by inhibiting α -glucosidase in the intestine, increases the life span of mice (Harrison et al., 2014), and exerts cardioprotective effects in patients affected by diabetes type 2 (Chiasson et al., 2002).

3.4.3 Inhibitors of the GH/IGF-1 Axis—Several reports indicate that downregulation of GH/IGF-1 pathway can extend life span in different species (Longo and Finch, 2003). For example, human IGF-1 receptor gene mutations have been found in centenarians (Suh et al., 2008) and decreased serum levels of IGF-1 predict survival in humans with exceptional longevity (Milman et al., 2014). Accordingly, GHR/IGF-1-deficient mice present a lower incidence and delayed occurrence of fatal neoplastic lesions compared with their wild-type littermates (Ikeno et al., 2009; Zhou et al., 1997). Similarly to what it was observed in (GHR/BP) knockout mice, patients affected by GHR-deficient Laron syndrome are characterized by a significant decrease of cancer and diabetes risk (Guevara-Aguirre et al., 2011; Ikeno et al., 2009; Shevah and Laron, 2007; Zhou et al., 1997).

According to these results, pharmacological interventions aimed at inhibiting GH/IGF-1 such as monoclonal antibodies and drugs directed against IGF-1R have been used in cancer patients (Carboni et al., 2009). Moreover, several classes of compounds that inhibit GH/IGF-1 axis have been used in patients affected by acromegaly (Giustina et al., 2014). Among the latter drugs, particular attention has been given to somatostatin analogs which lower serum GH and IGF-1 but, unfortunately also reduce other endocrine hormones and can cause serious side effects such as anorexia, diarrhea, and gallstones. For these reasons, the clinical antiaging use of somatostatin analogs is currently poorly understood and pursued. A GHR antagonist currently used in patients affected by acromegaly is pegvisomant which

specifically binds to and inhibits GHR (Kopchick et al., 2002; Trainer et al., 2000; van der Lely et al., 2001b). Interestingly, this compound lowers serum IGF-1 levels and blocks the diabetogenic action of GH (Trainer et al., 2000; van der Lely et al., 2001a). However, the very high cost and the need for frequent injections (van der Lely et al., 2012) provide a very tall obstacle for its application as a longevity and health span intervention. Finally, an interesting strategy to reduce IGF-1 action is to decrease its availability. In this respect, loss of the protease pregnancy-associated plasma protein-A (PAPP-A) reduces IGF-1 signaling and extends life span in mice while also reducing age-related diseases (Conover, 2013).

3.4.4 Activators of Sirtuins—Given their beneficial effects in promoting longevity, sirtuin family proteins are a very interesting drug target. Sirtuin-activating compounds (STACs) including plant-derived metabolites and the well-known resveratrol, represent the first and most potent sirtuin activator and have been shown to extend life span in various organisms (Hubbard and Sinclair, 2014; Pearson et al., 2008). Synthetic activators such as SRT1720 and SRT2104 improve the metabolic profile and extend life span and health span of mice under a high-fat and normal diet (Mitchell et al., 2014; Minor et al., 2011). Interestingly, SRT1720 improves insulin sensitivity, lowers plasma glucose, and increases mitochondrial capacity in experimental diabetes models, thus representing a promising new therapeutic approach for treating age-related diseases such as type 2 diabetes (Milne et al., 2007). Moreover, in rhesus monkeys, under a high-fat and high-sugar diet, resveratrol exerts antiinflammatory effects in visceral white adipose tissue (Jimenez-Gomez et al., 2013). In mice and nonhuman primates fed a high-fat diet, resveratrol protects against the effects of obesity and age-related metabolic decline, increases insulin sensitivity and mitochondrial functions, and prevents liver steatosis (Baur et al., 2006; Fiori et al., 2013; Jimenez-Gomez et al., 2013). In addition, resveratrol delays the onset of neurodegeneration and improves learning and memory in aged mice (Graff et al., 2013; Zhao et al., 2013). The promising results in preclinical models led the clinicians to test resveratrol in humans where it exerts beneficial effects on elderly and obese subjects (Timmers et al., 2011). Furthermore, resveratrol provides positive effects for systolic blood pressure, hemoglobin A1c, and creatinine in patients affected by type 2 diabetes (Hausenblas et al., 2015). Synthetic STACS have also been demonstrated to exert beneficial cardiovascular effects on healthy cigarettes smokers (Venkatasubramanian et al., 2013). The mechanism underlying the beneficial effects of resveratrol is controversial since it has been proposed that the direct activation of sirtuin 1 by resveratrol is an *in vitro* artifact (Behr et al., 2009; Kaeberlein et al., 2005; Pacholec et al., 2010) and that resveratrol works primarily by activating AMPK (Canto et al., 2009), potentially by inhibition of phosphodiesterases (PDE). AMPK may then activate sirtuin 1 indirectly by elevating intracellular levels of its cosubstrate NAD⁺ (Canto et al., 2009). Alternatively, resveratrol may first activate sirtuin 1 *in vivo*, leading to AMPK activation via deacetylation and activation of the AMPK kinase LKB1 (Hou et al., 2008; Lan et al., 2008).

3.4.5 Activators of AMPK Pathway—AMPK is a serine/threonine kinase which, upon activation by low cellular energy levels, exerts insulin-sensitizing effects resulting in increased glucose uptake in skeletal muscle and fatty acid oxidation in several tissues as well as decreased hepatic glucose production (Ruderman et al., 2013). Different AMPK activators

have been developed including biguanides, thiazolidinediones, agonists of glucagon-like peptide-1 receptor, salicylates, and resveratrol (Coughlan et al., 2014). Among biguanides, metformin is a drug used in the first line therapy for type 2 diabetes (Rena et al., 2013) that reduces the risk of cancer and overall mortality (Evans et al., 2005; Franciosi et al., 2013), cardiovascular disease, and possibly cognitive decline (Ng et al., 2014; No author, 1998; Wu et al., 2014a). However, before introducing metformin as an antiaging agent in generally healthy people, further studies to understand the mechanism of action of this compound are necessary. In fact, metformin is also able to inhibit gluconeogenesis, suggesting that, for people under diet-restricted or ketogenic diet, who depend on gluconeogenesis, it might be toxic. According to its prolongevity effects, metformin has been demonstrated to extend the life span of worms and rodents by targeting the folate cycle and methionine metabolism (Anisimov, 2010; Cabreiro et al., 2013; Martin-Montalvo et al., 2013).

3.4.6 Inhibitors of Inflammation Pathways—Strategies aimed at decreasing inflammatory pathways can be viewed as candidate target to combat and prevent aging-associated diseases. Various factors contribute to the generation of inflammatory conditions. Some of them are exogenous such as persistent cytomegalovirus infection (Sansoni et al., 2014), while some others are endogenously produced. The latter include circulating mitochondrial DNA (mtDNA) that is a potent inflammatory stimulus increasing with aging (Pinti et al., 2014), galactosylated *N*-glycans and proinflammatory micro-RNA (inflammaMIR), all of them potent inflammatory stimuli which increase in circulation with age (Dall’Olio et al., 2013; Olivieri et al., 2013). Approaches targeting one of the pathways that promote age-dependent inflammation, including the de-activation of inflammasomes (Youm et al., 2013), elimination of senescent cells (Tchkonia et al., 2013), and specific dietary restriction regimens have been developed (Berendsen et al., 2013). Initial evidence for the potential of antiinflammatory drugs in retarding aging comes from the effect of nordihydroguaiaretic drugs such as aspirin on life span increase in mice (Strong et al., 2008).

4. GEROSCIENCE AS A STRATEGY TO OPTIMIZE CANCER THERAPY

Conventional cancer therapy is mainly based on the use of chemotherapeutics, ionizing radiation, and novel drugs with specific molecular targets, whose major limitation is represented by the toxicity toward healthy cells. Thus, normal cells are damaged and acquire phenotypes similar to those observed during aging. For example, chemotherapy causes mutations and DNA alterations and induces oxidative stress in agreement with what occurs during aging. Moreover, epidemiologic studies have also observed that long-term cancer survivors treated with chemotherapy or radiotherapy have good response in term of tumor regression but often suffer of delayed effects related to cancer treatment including secondary tumors, cardiac pathologies, respiratory complications, infertility, and other chronic disorders. Thus, the discovery of strategies able to generate differential stress resistance (DSR) and sensitization conditions in which tumor cells are sensitized while normal cells are protected against the cytotoxicity of a chemotherapeutic drug is of central importance, particularly as many therapies include multiple chemotherapy drugs or the combination of chemotherapy and non-chemotherapy drugs which can also have adverse effects.

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Geroscience is a multidisciplinary field whose purpose is the understanding of the association between aging and age-related diseases and the elucidation of the mechanisms underlying cellular damage, protection, and death. Thus, geroscience offers the opportunity to find strategies focused on the protection of healthy cells without interfering with the efficacy of the treatment against malignant cells. In this respect, geroscience and cancer therapy could be considered complementary to each other since the former aims to protect the patient while the second to kill tumor cells (Fig. 3). In the following sections, we will discuss some of the geroscience-based approaches used to generate DSR and stress sensitization conditions in normal and malignant cells, in order to improve the killing of cancer cells, but also to preserve the patient's healthspan posttreatment.

4.1 Fasting and Fasting-Mimicking Diet

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Multiple cycles of fasting in tumor-bearing mice can selectively sensitize cancer cells to chemotherapy while protecting normal cells from the associated toxicity (Lee et al., 2012; Raffaghello et al., 2008). These phenomena are known as differential stress sensitization and DSR. Specifically, in healthy cells, fasting was found to activate protective metabolic pathways that confer resistance to a variety of chemotherapeutics (Raffaghello et al., 2008). In contrast, starved cancer cells were found to be unable to turn on such a protective response due to uncontrolled activation of growth promoting signaling cascades by oncogenes and to become even more sensitive to DNA damaging agents (Lee et al., 2012; Safdie et al., 2009, 2012; Shi et al., 2012). The protection of normal cells against chemotherapy-dependent damage was found to be associated to a reduction of IGF-1 and downregulation of protooncogene signals (Lee et al., 2010).

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In addition to protecting hematopoietic cells from chemotoxicity, multiple cycles of fasting promote hematopoietic stem cell self-renewal to alleviate or reverse the immunosuppression or immunosenescence caused by chemotherapy treatment and aging (Cheng et al., 2014). The latter effect was associated to inhibition of IGF-1 and PKA signaling.

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More recently, fasting has also been shown to enhance the therapeutic index of chemotreatments by exerting an anti-Warburg effect. In this way, cancer cells are shifted from a glycolytic mode into an uncoupled OXPHOS which promotes increased ROS generation and apoptosis (Bianchi et al., 2015). Preliminary clinical data indicate that fasting in cancer patients is not associated with major adverse effects and may reduce several of the toxic effects of chemotherapy (de Groot et al., 2015; Safdie et al., 2009) including a decreased toxicity to lymphocytes (Cheng et al., 2014). These preliminary clinical data have been collected as part of different pilot Phase I and Phase II clinical trials ([NCT01304251](#), [NCT01175837](#), [NCT00936364](#), and [NCT01175837](#)) that demonstrated the safety and efficacy of fasting cycles in combination with chemotherapy in adult oncologic patients.

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A recent study demonstrated that a FMD is as effective as fasting in killing different cancer cell types (Di Biase et al., 2016). In particular, the combination of chemotherapy and FMD increases the levels of lymphoid progenitor cells and cytotoxic CD8⁺ tumor-infiltrating lymphocytes, leading to a major delay in breast cancer and melanoma progression. In breast tumors, the mechanism underlying the latter effect involves the downregulation of the stress-protecting enzyme hemeoxygenase-1 (HO-1). These data indicate that cycles of FMD in

combination with chemotherapy can enhance T cell-dependent killing of cancer cells through the stimulation of the hematopoietic system and the enhancement of CD8⁺-dependent tumor cytotoxicity (Di Biase et al., 2016), although it is likely that the initial damage of cancer cells caused by the FMD (Shim et al., 2015) is a key step in the activation of the T cell-dependent immune response. According to these results, back-to-back studies reported that fasting enhances chemotherapy-induced immunosurveillance (Pietrocola et al., 2016).

Fasting has also been associated to tyrosine kinase inhibitors (TKI), which represent the most broadly applied cancer therapeutics. However, a major limitation of these agents is that their efficacy is short lived since the great majority of patients unavoidably relapse (Gridelli et al., 2014a,b). Thus, strategies that help increase the efficacy of these agents making them more powerful and capable of effectively eradicating cancer cells are warranted. In this respect, fasting has been shown to potentiate the in vivo anticancer effects of various TKI including erlotinib, gefitinib, lapatinib, crizotinib, and regorafenib through inhibition of MAPK kinase signaling and E2F factor transcription (Caffa et al., 2015).

4.2 Glycolysis Blockade

As mentioned in the previous chapter, 2-DG is a glucose analog that is phosphorylated by hexokinase to 2-DG-phosphate, which causes a glycolysis blockade. While early studies demonstrated that 2-DG exerts promising anticancer effects in experimental models (Maher et al., 2004), more recent reports show that the mechanisms underlying the activity of 2-DG is heterogeneous and sometimes 2-DG was shown to activate prosurvival pathways in tumor cells (Zhong et al., 2009). Furthermore, the clinical use of 2-DG has been limited by its toxicity upon long-term application (Dwarakanath and Jain, 2009; Landau et al., 1958; Vijayaraghavan et al., 2006). A more promising approach is to combine 2-DG with cytotoxic agents such as chemotherapeutic drugs and ionizing radiation in order to enhance their efficacy (Maschek et al., 2004). In this respect, 2-DG increases the therapeutic index of chemotherapeutic agents in mice-bearing human osteosarcoma and nonsmall cell lung cancer (Maschek et al., 2004). Moreover 2-DG synergizes with etoposide through the induction of immunogenic cell death and in this way increases the life span of immunocompetent tumor-bearing mice (Beneteau et al., 2012). Clearly, the combination of 2-DG with chemotherapy will result in a even higher degree of toxicity, which may be life threatening.

4.3 Nutrient-Sensing Pathway Interventions

As described in a previous section, one of the most relevant nutrient-sensing pathway involved in longevity is governed by GH/IGF-1 and glucose which trigger the activation of downstream mTOR/S6K, PI3K/AKT, Ras, and AC/PKA axes, most of them highly conserved from yeast to humans (Fontana et al., 2010). Rapamycin and its analogs (rapalogs) including temsirolimus and everolimus are mTOR inhibitors that have been approved by the FDA for treating different cancers. In addition to their anticancer effects as single agents, rapamycin and rapalogs have been shown to potentiate the efficacy of various cytotoxic agents such chemotherapeutic drugs, ionizing radiation, and proteasome inhibitors against malignant cells through the activation of caspase-dependent apoptosis (Mondesire et

al., 2004). In addition, rapamycin and rapalogs prevent epithelial stem cell senescence and protect mice from ionizing radiation-induced side effects (Iglesias-Bartolome et al., 2012). Accordingly, in vitro data indicate that also the combination of rapamycin and metformin protects normal fibroblasts and epithelial cells against the toxicity of cell-cycle specific chemotherapeutic agents such as mitotic inhibitors (Apontes et al., 2011). However, in vivo rapamycin and other m-TOR inhibitors also cause hyperglycemia which may potentially promote the progression of certain cancers and the sensitization of normal cells to chemotherapy.

Metformin has recently emerged as an adjuvant anticancer drug to be used in combination with chemotherapy in order to increase its efficacy and lower the doses. Specifically, metformin has been shown to increase the therapeutic index of various standard chemotherapeutic agents such as paclitaxel, carboplatin, and doxorubicin. This combinatorial effect includes tumor regression and prevention of relapse (Hirsch et al., 2009; Iliopoulos et al., 2011). In agreement with these results, a retrospective analysis of esophageal adenocarcinoma patients under metformin treatment indicated a better response to chemotherapy in subjects treated with metformin compared to those treated with chemotherapy alone (Skinner et al., 2013). The proposed mechanism underlying the chemosensitizing effect of metformin against tumor cells is based on downregulation of mTOR pathway and CSC genes (Honjo et al., 2014). In addition to enhancing the efficacy of chemotherapy, metformin has been shown to protect the host against the chemotoxicity by reducing the chemo-induced peripheral neuropathy and sensory deficits (Mao-Ying et al., 2014).

5. CONCLUSIONS

Cancer and aging are closely related to each other since they are both associated with damage to DNA and loss of function, but also because aging is a major risk factor for tumorigenesis. In turn, many cancer treatments are contributors to the aging process. From this perspective, geroscience-based approaches appear to be urgent since they provide a potential solution not only for preventing or delaying age-related disorders but also for the optimization and enhancement of conventional cancer treatments. In this context, there is a need to educate patients of the long-term side effects caused by standard cancer therapy and to inform them about the strategies used to prevent or reduce such toxicity. In this respect, FMD but also a few drugs including metformin targeting antiaging pathways, or the combination of dietary and pharmacological interventions represent one of the most promising approaches that could be incorporated into therapeutic protocols for oncologic patients in order to ameliorate the clinical outcome for these patients with an overall impact on the costs of medical care.

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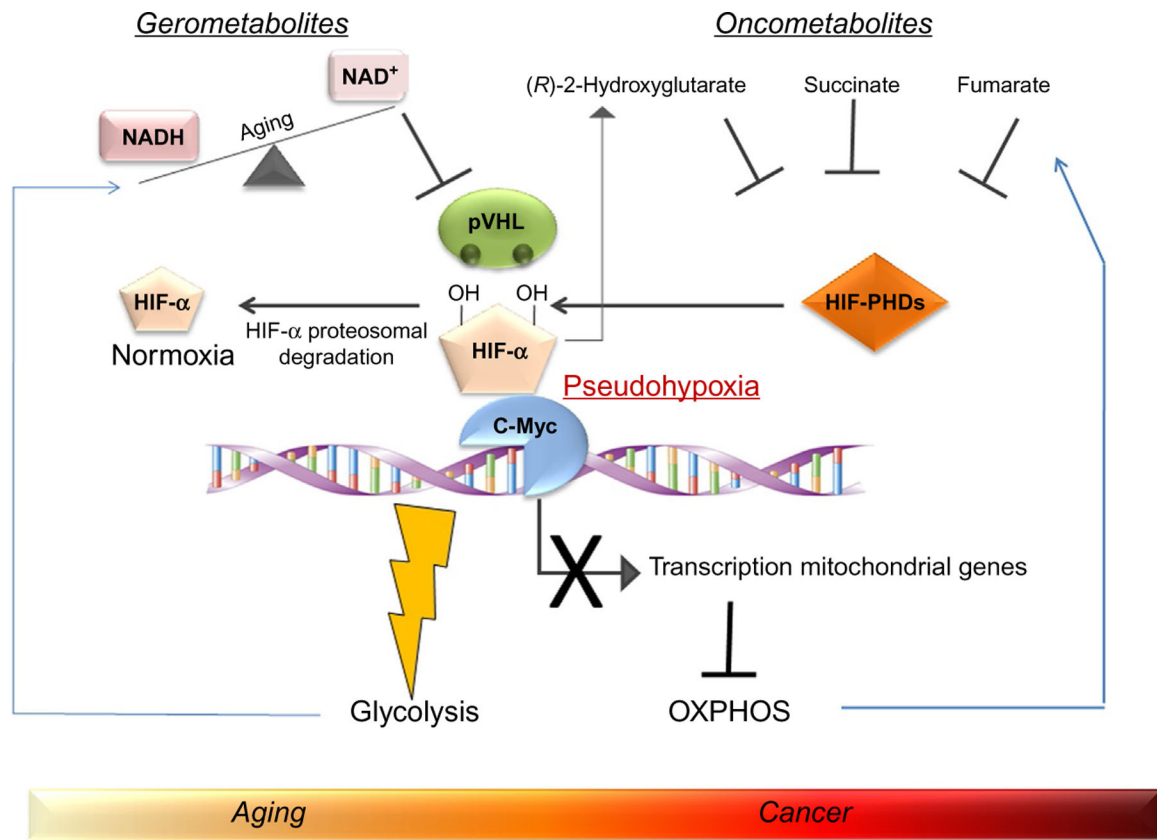
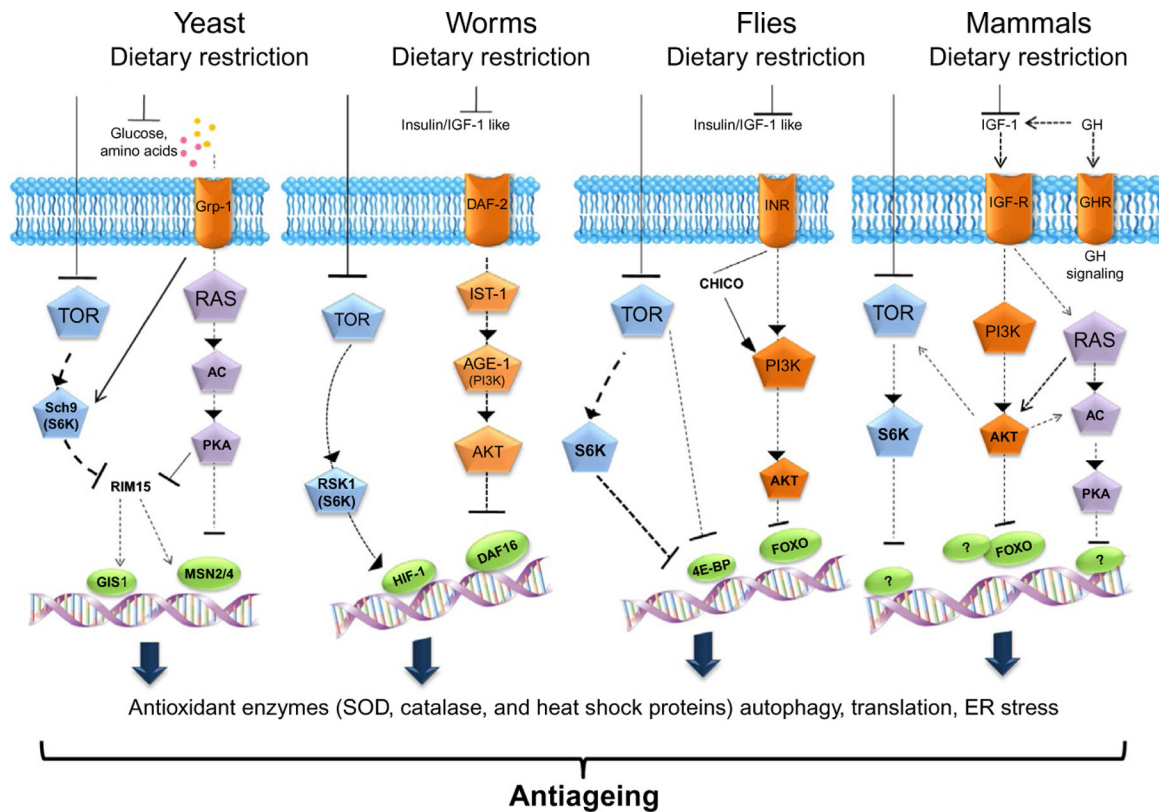
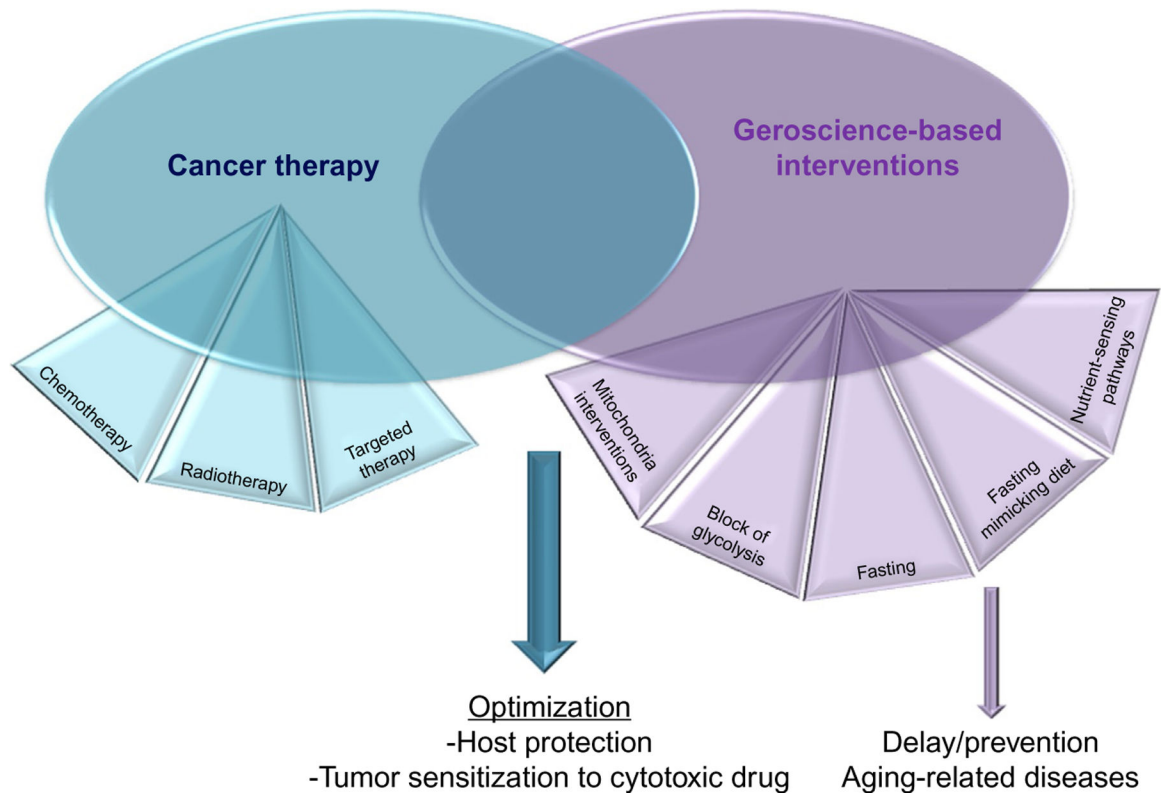


Fig. 1. The link between metabolites and oncometabolites in aging and cancer. Metabolites, such as NAD^+ , which decline as we age, cause the accumulation of oncometabolites through the induction of a pseudohypoxia condition (Gomes et al., 2013). This latter event occurs as a consequence of an increased HIF-1 α stabilization due to the ubiquitin ligase pVHL inhibition. Oncometabolites inhibit the interaction between HIF-1 α and prolyl hydroxylases (PHDs) resulting in a decreased HIF-1 α degradation. As a consequence, oncometabolites generate a pseudohypoxia condition as other metabolites do. This event converges on c-Myc activation, which represses the transcription of mitochondrial genes resulting in glycolysis promotion and oxidative phosphorylation (OXPHOS) inhibition.

**Fig. 2.**

The nutrient-sensing pathways in aging. Certain nutrients sensing pathways are evolutionarily conserved in different organisms including yeasts, worms, flies, and mammals. These pathways are triggered by ligands (glucose and amino acids in yeast, insulin and insulin growth factor-1 like (IGF-1) in worms and flies, IGF-1 in mammals) which activate a receptor (G protein-coupled receptor 1 (Gpr1) in yeasts; DAF-2 in worms; insulin-like receptor (InR) in flies; and IGF-1R, IR-A, and IR-B in mammals). The signal is then transduced through adaptor proteins which include Ras in yeasts, IST-1 in worms, CHICO in flies, and insulin receptor substrate 1 (IRS1–4) and Ras in mice. In yeast, Ras activates adenylate cyclase and protein kinase A (PKA). In worms, flies, and mice, the adaptor proteins mediate the activation of phosphatidylinositol-4,5-bisphosphate 3-kinase (PI3K) (AGE-1 in worms) which generates phosphatidylinositol (3,4,5)-trisphosphate (PIP3). PIP3 activates AKT/PKB, or PKA in yeast, phosphorylates, and inactivates specific stress resistance transcription factors (MSN2/4 and GIS1 in yeasts, DAF-16 in worms, Forkhead box protein O (FOXO) in flies and mice) involved in the upregulation of protective stress resistance genes such as superoxide dismutase (SOD), catalase, and heat shock proteins (HSPs). A parallel pathway involved in the inactivation of the above-mentioned transcription factors is governed by the target of rapamycin (TOR) and its substrate S6 kinase (S6K). The nutrient-sensing pathways can be down-modulated by diet restriction, which reduces the levels of glucose and IGF-1.

**Fig. 3.**

Geroscience and cancer therapy. Cancer therapy is mainly based on the use of chemotherapeutics, ionizing radiation, and novel drugs with specific molecular targets, whose major limitation is represented by the toxicity toward healthy cells. This is mainly due to their inability to distinguish between healthy and cancer cells. Geroscience, defined as the science of healthy aging, can provide not only strategies able to prevent or delay aging and age-related disorders but also to optimize conventional cancer therapy by protecting the host and sensitizing the cancer cells to the cytotoxic agent. These approaches include fasting and fasting-mimicking diet, mitochondrial interventions, glycolysis blockers, and inhibitors of the nutrient-sensing pathways.