



The role of leucine in the activation of cellular metabolism: a large integrative review

Amir Salomão Gebrin^{1*}, Idiberto José Zotarelli-Filho²

¹ USP – University of São Paulo, São Paulo, Brazil.
 ² FACERES – Faculty of Medicine of São José do Rio Preto, São Paulo, Brazil.

Corresponding Author: Dr. Amir Salomão Gebrin, USP - University of São Paulo, São Paulo, Brazil. Email address: amir.gebrin@gmail.com DOI: https://doi.org/10.54448/ijn22S201 Received: 08-11-2022; Revised: 10-28-2022; Accepted: 11-11-2022; Published: 11-23-2022; IJN-id: e22S201

Abstract

This review addressed the signaling of cellular activation by leucine, discussed the risks of excessive signaling by proteins in the Western diet, and explored the potential of leucine stimulation in tissue regeneration. As result, amino acids are, in addition to building blocks of macromolecules, cellular activation signals. Essential amino acids are not produced by animals and leucine appears to be the main signaling amino acid. Mammals adjusted the cell activation and growth rate of their young by the leucine concentration of the milk produced. Several studies demonstrate the benefits of leucine supplementation in preventing sarcopenia, improving muscle and liver performance, as well as a possible neuroprotective role in head trauma and dementia. However, its excess, so common in the Western diet, is related to obesity, type II diabetes, neurodegenerative diseases, and cancer. The mTORC1 kinase integrates cellular activation stimuli from macro protein synthesis to epigenetic regulation. Controlling mTORC1 activity by consuming leucine can prevent, treat, or cause disease. A greater understanding of the regulatory effects of leucine and mTOR in unstable tissues such as tumors or fragile tissues such as the CNS are areas of great relevance and with extensive fields still to be explored.

Keywords: Leucine. Cell metabolism. Signaling. Tissue regeneration.

Introduction

Amino acids are essential elements used as "building blocks", energy substrates in certain metabolic situations, cell activation signals, and buffering agents (glutamate in the CNS, for example) **[1-6]**. Insulin and

growth factors are incapable of cell activation in the absence of amino acids such as leucine **[7-13]**.

Plants and bacteria synthesize their amino acids, but animals are unable to synthesize half of the twenty existing amino acids, thus making their ingestion necessary, hence called essential amino acids, the main ones being branched chain (leucine, isoleucine, and valine), known as BCAAs (Branched-Chain Amino Acids) [13].

Therefore, this review addressed the signaling of cellular activation by leucine, discussed the risks of excessive signaling by proteins in the Western diet, and explored the potential of leucine stimulation in tissue regeneration.

Milk, leucine and mTOR complex

Milk and its derivatives are the main natural source of leucine. According to Melnik (2012) **[14]**, the action of milk on the growth rate of the calf has a direct relationship between growth and leucine concentration in the species' milk. Rat milk has 7.9 mg of leucine/mL and their pups double in weight every 4 days; calves double in weight after 40 days (cow's milk has 3.3 mg leucine/mL) and human milk has the lowest concentration of all mammals: 1.0 mg/mL of leucine, which slows growth for greater learning acquisition. The human baby doubles in weight only after six months of life.

Furthermore, Melnik describes breastfeeding not as a simple source of nutrition, but as an evolution of cell activation signaling **[1,14-16]**. The leucineactivated mTOR (mammalian/mechanistic Target Of Rapamycin) complex integrates nutritional, hormonal, and environmental stimuli in the control of cellular metabolism **[17-19]**.

In the 1970s Sabatini et al. **[18]** investigating the lack of fungi in the soil of Easter Island discovered a

bacterial strain that produced a potent antifungal, named rapamycin (named after the island Rapa Nui, in the local language). Rapamycin is a potent cellular inhibitor by blocking the mTOR protein **[20-22]** and is being used in clinical practice even today as an immunosuppressant, antistenotic, and in the treatment of cancer **[23-25]**.

The discovery of the mTOR complex began with the discovery of its inhibitor, rapamycin in 1975, and

with the cloning of its genes in 1993. It was discovered that mTOR forms two distinct complexes when associating with other proteins: mTORC1 when binding to RAPTOR (Regulatory-Associated Protein of mTOR) and mTORC2 when associated with RIcTOR (Rapamicyn-Insensitive Companion of mTOR) (**Figure 1**) [26-28].

Figure 1. Structure mTORC1 (A) and mTORC2 (B) and their main functions in cell activation.



The mTORC1 complex consists of the union of mTOR, Raptor, mLST8 (mammalian Lethal with Sec13 protein T8), Deptor, and DAS40 protein. Activated initiates cellular anabolism with lipid and protein synthesis. Deactivated, it causes autophagy by activating the kinase glycogen synthetase GSK3 β [29,30]. The mTORC2 complex organizes the actin cytoskeleton, cell migration, and survival.

Also, mTORC1 integrates cellular activation stimuli from fungi to mammals **[31-35]**. Several studies indicate mTORC1 as a central regulator of gene transcription, ribosomal translation, mRNA transcription, suppressor of autophagy, and activator of ribosomal and mitochondrial neogenesis **[36-41]**. The human mTORC1 framework is a dimer of mTOR, Raptor, and mLST8 **[30]**. The mTORC1 complex is extremely sensitive to amino acid exposure and leucine appears to be one of its main activators **[42-45]**. The absence of leucine prevents mTORC1 activation even in the presence of all other amino acids, growth factors, glucose, and insulin **[46-48]**. The mTORC2 complex appears to be insensitive to amino acids, responding primarily to trophic factors **[48-50]**.



Actions of leucine on muscle, liver, and CNS

Tissue changes from trauma, disease or aging disrupt cell physiology and regenerative responses are necessary for homeostasis. Mammals have little regenerative capacity in vital organs such as the heart and central nervous system (CNS) but the liver, skin, intestine, and muscles have the great regenerative capacity in adult mammals **[51-53]**.

Understanding the mechanisms of tissue regeneration is essential for therapeutic interventions and the mTORC1 system seems to be its main pathway. In the CNS (minimum regenerative capacity), the mTORC1 pathway is stimulated in the inactivation of PTEN (phosphatase and tensin homologous) and TSC1 (tuberous sclerosis complex 1) kinases, allowing axonal regenerative expansion **[54-57]**.

Leucine is well known for its action on muscle hypertrophy, obesity, metabolic disorders, liver disease, immune activation, and cancer **[14]**. The correlation between obesity and an excessive supply of amino acids is observed in obese individuals due to mTOR hyperactivation, which may justify greater viral replication in these people and a worse prognosis in cases of COVID-19 **[58]**. Interestingly, the coronavirus has a 5'-end mRNA and uses the same cellular protein translation machinery, which is hyperactivated in the obese **[59,60]**.

Amino acid signaling has been extensively studied in recent decades for therapeutic use in multiple trauma, severe burns, and senile sarcopenia, where the clinical benefit has been demonstrated in increasing the supply of BCAAs **[61-65]**.

Muscle

Leucine increases muscle performance during exercise and its intake reduces fat mass and prevents both senile and inadequate diet obesity, as well as type 2 diabetes **[66-70]**. Leucine is the most important amino acid for protein synthesis **[71]** and can be administered as a nutraceutical agent in the prevention of sarcopenia.

Leucine has been known to induce muscle anabolism since the 1970s. In skeletal muscle, 20% of leucine is metabolized for energy production (reduction of glutamate and keto acids). The remaining leucine (80%) activates protein synthesis (via mTORC1) and satellite cell expansion **[72-75]**. Maltais et al. (2016) **[75]** randomized 26 overweight sarcopenic men into resistance training for 4 months, offering post-exercise dairy or rice milk as a control group. Resistance training increased lean body mass (DEXA) in both groups, but the dairy group decreased their body fat rate more and gained more muscle mass.

Another similar study randomized 26 sarcopenic obese women to receive hydrolyzed whey protein (whey protein) or placebo for 3 months under resistance training, resulting in a greater increase in lean mass in the group receiving the milk protein **[76]**. HMB (hydroxy-methyl butyrate) is a leucine metabolite used in an attempt to prevent muscle breakdown or lean mass gain, but it has controversial results in human and animal studies **[77-79]** performed a meta-analysis of 11 randomized trials of HMB in training. of resistance and concluded that there was no significant effect on lean mass gain, fat mass loss, or strength increase with HMB, even advising its use as a nutraceutical supplementation.

Gran and Cameron-Smith (2011) **[80]** studying human muscle cultures demonstrated the activation of mTORC1 by leucine at physiological doses. Chronic stimulation resulted in increased eIF4G activity (ribosomal transcription) at two peaks: 3h and 24h after leucine/insulin introduction. Continuous stimulation by leucine and hormones can generate a persistent cellular anabolic state **[81]**.

Deldique et al (2008) [82] studying muscle

cultures observed a 50% increase in mTORC1 activation soon after the addition of 5mM leucine, falling at 30 minutes. Atherton et al (2010) **[83]** reported lower doses (2 Mm) also activated mTORC1. They demonstrated that 5 mM leucine increases 70-S6K activation by 10-fold and 2 mM increases p70-SK6 activity by 5.9-fold in muscle cultures.

In vivo tissue perfusion models are also used in studies of anabolic stimulation, protein synthesis, and/or cellular edema studies. Some studies have found that increasing the concentration of amino acids by up to ten times its plasma value does not result in tissue edema **[84,85]**.

Bolster et al (2004) **[86]** cannulated and perfused the hind legs of rats with 1x or 10x concentrations of serum leucine and monitored mTOR, p70-SK6, eIF4E, and 4E-BP1 activation. They observed a 66% increase in protein synthesis in the gastrocnemius muscle and a 70% increase in the soleus muscle of the paws perfused with a 10x leucine solution. The effects of tissue infusion supplementation with 10x leucine did not cause muscle edema in the rats.

Several studies of muscle culture use leucine in concentration five or ten times the physiological one, being able to increase protein synthesis. Omitting leucine from the solution, even increasing the concentration of the other amino acids tenfold, does not increase protein synthesis in muscle **[87-89]**. Peyrollier et al (2000) **[90]** adding 2 mM leucine (normal concentration) to amino acid-free muscle cultures obtained rapid anabolic activation, doubling the protein synthesis marker p70-S6K and increasing five-fold activation of the PI3K pathway. The addition of leucine also increased the cellular uptake of amino acids by 50%.

MAP4K3 (Mitogen-Activating Protein Kinase-Kinase-Kinase-Kinase-3) activity is also regulated by amino acids, but not by insulin. MAP4K3 kinase activates satellite cell myogenesis in rat muscle, however complete mTORC1 activation occurs only in the presence of leucine **[91]**.

Supplementation with BCAAs improves oxidative respiration and prevents mitochondrial dysfunction. Studies show mitochondrial biogenesis in the cardiac and skeletal muscles of patients with Barth myopathy **[92]** and skeletal muscle of elderly rodents **[93]**. Supplementation with BCAAs increased mitochondrial survival in cardiomyocytes from animals intoxicated by doxorubicin **[94]**.

Leucine signaling increases the number and size of mitochondria in the muscle fiber. It is an energy substrate, potentiates the oxidation of fatty acids, and increases glucose absorption **[95-97]**. Leucine supplementation prevents mitochondrial dysfunction in nervous tissue, muscle, and liver, and is indicated in aging, neurodegenerative and cardiovascular diseases, obesity, and diabetes **[98-104]**.

Amino acid supplementation increases cellular respiration by inducing mitochondrial biogenesis **[105-108]**. Mitochondria undergo cycles of fusion and fission that either unite with each other (respirosome) or attach to lysosomes and the endoplasmic reticulum **[109-111]**. Mitochondria show ancestral symbiosis with the host cell, as they have their own (circular) DNA, but depend on proteins transcribed from the host cell's DNA **[112,113]**.

There is evidence that leucine enhances mitochondrial biogenesis via PGC-1a and SIRT1 factors **[114-118]**. In animals, supplementation with BCAAs resulted in mitochondrial biogenesis with increased SIRT1 factor in skeletal muscle **[93,107]**.

Liver

The literature is rich in evidence of the benefits of BCAA supplementation in hepatic failure encephalopathies **[119]**. Pavlov (1893) made the first description of hepatic encephalopathy with ataxia and convulsions after portocaval anastomosis in meatfed dogs and the reversal of encephalopathy by switching to a milk-only diet in these same dogs **[120]**.

Muting and Wortmann in 1956 described a decrease in the concentration of BCAAS in cirrhotic patients and an increase in aromatic amino acids **[120]**. This condition became known as Fischer's ratio: the lower the concentration of BCAAs, the greater the brain intoxication **[121]**. Supplementation with leucine/BCAAs is indicated in liver disease in patients with cirrhosis **[122-124]**.

Supplementation with BCAAs is a nutracenic factor in chronic liver disease **[123]**. Decreased serum BCAA levels are credited with muscle uptake and glutamine synthesis by the cirrhotic patient's muscle. In vitro experiments have shown that high levels of ammonia cause leucine oxidation in muscle, consuming nitrate. This sequestration of ammonia in situations of azotemia produces glutamine in muscle tissue **[125]**.

Supplementation with BCAAs stimulates mitochondrial biogenesis in the liver, preventing alcohol steatosis in animal models [126]. Increased postregeneration hemihepatectomy has also been demonstrated with leucine supplementation [127]. Jefferson & Korner (1967) [128] perfusing the liver of rats with amino acids at a physiological dose of growth hormone (GH) did not obtain an increase in protein production, but with three times the dose of amino acids (the same dose of GH), they obtained increased protein

synthesis by the liver.

Krause et al (2002) [129] studying the effects of leucine, glutamine, and insulin on rat liver cells found that isolated insulin does not increase anabolism in the liver, but isolated leucine can activate p70-S6K and ACC (synthesis markers). protein and lipid, respectively), concluding that the action of insulin on the hepatocyte depends on the joint presence of leucine. Dennis et al (2011) **[130]** perfused the liver of rats with amino acids with and without insulin and found that at normal or doubled insulin concentrations they obtained maximal PI3K/Akt stimulation, but no increase in protein synthesis. Perfusion without insulin and at four times the concentration of amino acids produced moderate protein synthesis but a combination of insulin and four times more amino acids caused maximum protein synthesis, suggesting that stimulation by amino acids in conjunction with insulin is necessary for mTORC1 activation.

CNS

The central nervous system (CNS) depends on oxygen and glucose, but adaptive responses allow the use of ketone bodies in metabolic crises, where amino acids are an energy source that does not require mitochondrial respiration **[131-134]**. Under CNS stress conditions, BCAAs (leucine, isoleucine, and valine) are metabolized to glutamate and keto-acids. Glutamate can yield glutamine or alanine and, in a second reduction, acetyl-CoA to the Krebs cycle **[135,136]**.

BCAAs are reduced to glutamate and ketone bodies in the CNS by glial mitochondrial aminotransferases (astrocytes) and by cytoplasmic aminotransferases in neurons. The presence of leucine stimulates the reduction of glutamate in glutamine and acetyl-CoA, an energy source for the Krebs cycle **[137,138]**.

The action of BCAAs in Neuronal Metabolic Crisis

After traumatic brain injury (TBI) there is a halt in mitochondrial glucose metabolism in the CNS, a situation known as a "metabolic crisis". Some studies suggest that supplementation with BCAAs promotes neuroprotection, minimizing neural damage and improving clinical recovery **[139-141]**. Jeter et al (2013) **[142]** studied the metabolic alterations of human TBI, measuring the serum level of BCAAs in the first 24h in moderate TBI (EG: Glasgow scale > 12), severe TBI (EG <8), orthopedic traumatic injuries and in healthy individuals. They observed little reduction in BCAAs after moderate TBI and in orthopedic injuries, but very low values in severe TBI. Low serum BCAA levels in TBI are also a prognostic factor for intracranial hypertension (ICP≥25 mm Hg).

Mitochondria are the cellular energy machinery par excellence and use glucose and oxygen as their primary substrate. But in the absence of glucose or oxygen - or the presence of any other mitochondrial dysfunction ketone bodies and acetyl-CoA are the energy source of the CNS. Differentiating between ischemia and metabolic crisis is fundamental in the management after TBI. Studies suggest that supplementation with intravenous BCAAs minimizes neurological damage to the CNS after the reversal of ischemia **[143,144]**.

Ketone bodies can reach 70% of the energy matrix in the CNS of mammals during fasting, trauma, or prolonged exercise. Blood-brain barrier (BBB) permeability for ketone bodies also increases during lactation and fasting. Recall that mammalian infant neuronal development is dictated by milk, which is rich in BCAAs, glucose, and fats **[145-150]**.

Leucine, glutamate, and glutamine metabolism in the CNS

Astrocytes are responsible for brain energy stability through the production of ketone bodies from amino acids and fatty acids in metabolic crises by mitochondrial aminotransferases (BCTAm) **[151,152]**. Leucine (typical BCAA) is the amino acid that crosses the BBB the fastest **[151-154]**, being metabolized in astrocytes as soon as they enter the BBB. It is estimated that 30% to 50% of CNS glutamate and glutamine derive from leucine absorbed in the BBB **[155-158]**.

Nissin et al (1987) **[159]** were the first to suggest buffering in the CNS by glutamate and ketoacids in leucine transamination. These keto acids (KICs) are taken up by neurons and re-aminated into leucine, with the consumption of glutamate in the synaptic cleft (Figure 1). Excess glutamate in the synaptic cleft is responsible for secondary injury to the TBI/SCI, tissue ischemia, and sequential neuronal death after CNS trauma [160-163]. Glutamate needs to be quickly removed from the synaptic cleft, being taken up by astrocytes and reduced to keto acids (mitochondrial aminotransferases: BCTAm) - glutamate-glutamine cycle - or reduced to keto acids by neurons (cytosolic aminotransferases: BCTAc) - leucine-glutamate cycle (Figure 2a). Glutamate can be transformed into glutamine (glutamine synthetase - GS) in astrocytes and exchanged for leucine in the BBB (Figure 2b) or taken up by the neuron, returning to glutamate via glutaminase (GlnAse) [164-166].

Figure 2. (A) Glutamate is removed from the synaptic cleft by the neuron itself and reduced to keto acid + leucine by cytosolic aminotransferases (leucine-glutamate cycle) or taken up by astrocytes and reduced to glutamine + keto acid by mitochondrial aminotransferases (leucine-glutamate cycle). **(B)** Glutamine is exchanged in the BBB for leucine in counterflow pumps or taken up by the neuron (returning to glutamate via glutaminase). The astrocyte can also take up glutamate and convert it to glutamine by the enzyme glutamine synthetase.



Figure 2. (C) Equilibrium of reanimation of keto acids generating glutamate (in the neuron) or leucine (in the astrocyte).



Source: Own authorship.

Leucine is the amino acid that most easily crosses the BBB, via transporters' counter-exchange for glutamine. This leucine is taken up by the astrocyte and mBCAT metabolizes leucine into glutamate and keto isocaproic acid (KIC). This is reanimated in the neuron into leucine and the ketoacid a-ketoglutarate (aKG), with the consumption of glutamate (**Figure 2c**).

The CNS is the only tissue that has both aminotransferases and BCAAs are fundamental to biochemical regulation in the CNS, because, in addition to being a raw material in the synthesis of glutamate, they buffer its excess in the CNS, avoiding toxic levels. /leucine and ketoacid/glutamate and observed three times more leucine formation than glutamate **[165,166]**. The participation of astrocytes in the synapse resulted in the concept of the tripartite synapse **[167]** and the presence of a dense extracellular matrix, "sealing off" some synapses and preventing the extravasation of glutamate from the cleft resulted in the concept of the quadripartite synapse **[168,169]**. Schafer et al (2013) **[170]** also describe the participation of microglia in the regulation of some synapses (**Figure 3**).

Figure 3. Neurons and astrocytes modulate the presence of glutamate in the synaptic cleft in the tripartite synapse model. A dense extracellular matrix "seals" some synapses preventing the diffusion of neurotransmitters in the quadripartite model.



Metabolic alterations of BCAAs and their effects on the CNS

Aminoacidopathies are errors in amino acid metabolism. Maple syrup urine disease (MSUD), or human leucosis, is a congenital deficiency of the ketoacid dehydrogenase that metabolizes BCAAs resulting in their progressive accumulation. It is a rare and serious genetic disease with encephalopathy crises, lethargy, lowering of consciousness, convulsions, and death due to an excessive increase in serum leucine (normal: 100 +-60 μ ml/L; MSUD up to 60,000 μ mol/L). Complete elimination of BCAAs from the diet or urgent dialysis during decompensations reverses the neurological picture [171-178].

Studying the intelligence coefficient (IQ) of children with MSUD under 6 years, Hoffman et al (2006) observed IQ 1.2 times higher in children with a serum leucine level below 200 μ mol/L (189 \pm 82 μ mol/L) compared to children with higher plasma leucine levels (572 \pm 217 μ mol/L). Leukemias of up to 1000 μ mol/L may even be asymptomatic, but negatively affect intelligence scores **[174]**.

Several studies indicate alterations in BCAA metabolism in neurodegenerative diseases such as Alzheimer's, Parkinson's, and Hungtinton **[179,180]**. Low levels of valine are linked to accelerated cognitive decline. On the other hand, high levels of valine reduce the risk of Alzheimer's. BCAA-related "metabolic signatures" have been identified in other diseases such as senile obesity, type II diabetes, and atherosclerosis **[181,182]**.

Recent studies in epidemiology have shown that protein intake is vital for brain function in the elderly population. Shang et al (2021) **[182]** in a 9-year cohort study linking protein consumption and dementia found that higher protein intake was associated with less cognitive decline in the elderly. Sato et al (2021) **[183]** observed lower neuroinflammation in protein-deficient mice after supplementation with essential amino acids.

Besides, mTOR activation in brain tissue optimizes memory and learning **[184,185]**. During aging, there is a decrease in neuronal mTOR activity **[186-188]**. It has been shown that in retinal and cerebral cortex neurons, mTOR signaling decreases with age and alters the capacity for axonal regeneration and dendritic remodeling, underscoring the importance of the consumption of essential amino acids **[189,190]**. In this context, Suzuki et al (2020) **[191]** conducted a randomized double-blind study of cognitive assessment in adults aged 55 years and over with supplemental essential amino acid intake. Daily intake of 3 g or 6 g of amino acids for 12 weeks resulted in better attention, cognition, and psychosocial functioning compared to the pre-supplementation state and the placebo group.

Elderly people with dementia have a lower protein intake than healthy elderly people. Tynkkynen et al (2018) [192], in a cohort study, demonstrated low serum levels of BCAAs associated with the development of Alzheimer's disease. Fernando et al (2018) [193], in another cohort study, observed that the more protein was consumed, the lower the presence of β -amyloid in the brain of the population studied. These results highlight the protective impact of protein intake on the brains of older adults. Leucine is a BCAA known to activate the mTOR pathway. Glycogen synthetase 3β kinase (GSK-3 β) is a counter-regulatory and potent inhibitor of mTORC1. GSK-3^β is hyperactivated in neurodegenerative diseases [194-198] and the accumulation of β -amyloid is responsible for neurotoxicity in tauopathies [198-201].

Cellular uptake of leucine

Despite the recognition of the importance of amino acids in cell activation since 1955 (Hosios et al, 2016) **[202]**, only in the last decades has there been a better understanding of the mechanisms of cell activation by amino acids, their transporters, metabolizing enzymes and correlation with diagnosis and treatment of tissue disease and dysfunction **[203-206]**. The uptake of amino acids depends on specific channels, but also nonselective endocytosis processes. There are at least 17 amino acid transport channels, the main ones being the L (leucine-preferred), A (alanine-preferred), and ASCT2 (alanine-serine-cysteine 2) transporter **[207-209]**.

Free amino acids are rapidly taken up by cells via membrane channels and activate mTORC1 within minutes (Nicklin et al, 2009) **[210]** whereas albumin or larger proteins activate mTOR only after 2 hours, peaking at 4 hours. Therefore proteins taken up by endocytosis activate mTORC1 much more slowly than free amino acids **[211]**.

Selective uptake of amino acids

Amino acids such as leucine and glutamine are as essential to cellular metabolism as oxygen and glucose **[212-215]**. The L channel is the main transporter of essential amino acids and is highly expressed in the brain, gonads, pancreatic islets, and placenta, but also tumors such as lung, prostate, and breast. Blockade of L channels, in particular LAT1, results in apoptosis and is a therapeutic target in the treatment of acute lymphoblastic leukemia, osteosarcoma, and cholangiocarcinoma **[216-220]**. The LAT1 channel is the main transporter of essential amino acids to the CNS, T lymphocytes, and skeletal muscle, with LAT1 increasing in skeletal muscle 1 and 3 hours after ingestion of essential amino acids. The LAT1 transporter is of great importance in amino acid uptake and cell signaling **[221-223]**. Leucine

transport by L channels (SLC1A5, SLC7A5, and SLC3A2) exchanges intracellular glutamine for extracellular leucine. Glutamate, metabolized to glutamine, also enhances leucine absorption in Lexchange channels **[224,225]** (**Figure 4**).







The LAT1 transporter is the main controller of free amino acid entry and subsequent mTORC1 activation in the brain, muscle, and immune system. In the brain, the LAT1 channel is essential for the development of the nervous system and its exclusion in KO (Knock Out) mice is lethal. In humans, LAT1 mutations are related to autism spectrum disorders and disorders such as microcephaly and seizures **[226]**. Inhibiting the uptake of BCAAs may be beneficial in neurological tumors with high LAT1 expression such as in gliomas **[227]**.

LAT1-dependent amino acids influence the bone skeleton, a tissue that undergoes constant remodeling dependent on the activation of osteoclasts and osteoblasts **[225]**. Ozaki et al (2019) **[226]** identified the LAT1 transporter in osteoclasts and its reduction in post-ovariectomy mice. Decreased leucine uptake inhibits mTORC1 activation in osteoclasts resulting in osteoporosis. mTOR activation promoted bone loss recovery in LAT1 knock-out mice. There is a strong dependence of leucine levels on the growth of bone sarcomas. Biopsies show an increase in BCAA enzymes in these tumors and the use of a leucine analog, N-Acetyl-Leucine Amide (NALA) blocks leucine uptake, dramatically decreasing the activity of bone sarcomas **[228]**.

Leucine is an amino acid transported by LAT1 channels and its blockade is a target in cancer control, a channel overexpressed in malignant tumors **[227-231]**. High LAT1 expression in cancer biopsies is even a poor prognostic factor. The activated mTORC1 system increases the expression of amino acid transporters

[232-233]. Lymphocyte activation in the immune response depends on metabolic reprogramming, with increased expression of glucose and amino acid channels for rapid proliferative expansion [234]. Leucine is a potent mTORC1 stimulator and depleting leucine or blocking its transporter prevents T lymphocyte activation to the same extent as total amino acid deprivation. Inhibition of leucine entry into T cells prevents their proliferation, allowing control over allergies and lymphomas for example [235-244].

Non-selective uptake of amino acids

Mammalian cells utilize glucose and free amino acids as an energy source, even in protein- or albuminrich environments such as plasma **[245-247]**. In addition to specific transporters, cells have developed alternative uptake of amino acids in the presence of growth factors or ischemic and deficient conditions. Although mTORC1 is activated only by free amino acids such as leucine, these are the smallest fraction in circulating plasma. Even with an increase in selective transporters, the macropinocytosis system is the main uptake mechanism during cell activation **[248]**.

The first cellular change that occurs after activation of the membrane receptor by growth factor is surface cytoskeleton remodeling **[246-248]**, which forms pseudopods that "embrace" large amounts of extracellular solute. Even in tumor cells, which are independent of growth factors, mTORC1 activation depends on the uptake of free amino acids **[249-255]** (**Figure 5**). **Figure 5. A)** Entry of amino acids and glucose into the cell occurs through selective channels (left) or, more intensely and nonspecifically, via macropinocytosis activated by growth factors (right) - adapted from Yoshida et al, 2009. **B)** Graph showing a large amount of albumin, other proteins, etc. and the minimum fraction of free amino acids in the blood plasma (smallest slice) highlighted - adapted from Palm et al, 2015.



B



Also, macropinosomes are growth factor signal transducers **[256]**. Macrophages exhibit immediate macropinocytosis after exposure to the growth factor MCSF **[257-259]**. The protein synthesis marker S6K increases 5 minutes after the addition of MCSF to the culture medium, as well as the anabolic activation markers MAPK, ERK, PI3K, and mTORC2. Cultures in amino acid-rich media have greater mTORC1 activity than amino acid-poor media with the same MCSF concentration **[260]**.

Growth factors activate mTORC1 by macropinocytosis with the rapid uptake of free amino acids. mTORC1 activation appears to be proportional to

leucine uptake **[261]**. With the same PDGF trophic factor concentration, but different leucine concentrations (0.4 mM and 4 mM), there was an increase in mTORC1 activity in cultures with higher leucine concentrations.Macropinocytosis was first demonstrated by Lewis in 1931, calling his description "pinocytosis" ("cell engulfing") **[255]**. Macropinosomes form macro protein vesicles that bind to the Golgi Complex and lysosomes hydrolyze macro proteins into the free amino acids required for mTORC1 activation **[262-264]**.

Electron microscopy studies have demonstrated macropinocytosis in the CNS and its decrease appears

to be related to amyloid accumulation and neurodegenerative diseases such as Alzheimer's **[265,266].** Macropinocytosis also occurs at the regenerating ends of axons; the growth cones. In vitro and in vivo analyses demonstrated that these terminations form membrane extensions with high molecular weight (10 KDa) vesicles, suggesting macropinocytosis in the axon and synaptic connections **[267-270]**.

By verifying the amount of leucine captured by transporters and by macropinocytosis, the PDGF-dependent mTORC1 activation was measured by the presence of the dipeptide Ala-Leu (does not pass through the transporters) in culture media. There was an increase in S6KF only 30 minutes after the introduction of the AlaLeu dipeptide, indicating that mTORC1 activation occurs only after the hydrolysis of AlaLeu into free leucine. The presence of free leucine activates mTORC1 between 2 and 3 minutes [**261**].

In tumor cells, macropinocytosis occurs without the need for growth factors (RAS-mutant cells). These have enormous energy demand that is met by glutamine. Glutaminolysis generates the NADPHs and fatty acids needed for growth without the need for mitochondrial respiration. This is known as the Warburg Effect and allows cell growth even under ischemic conditions [271-276]. The Warburg effect is the dissociation of mitochondrial metabolism generating a cellular growth phenotype even in hypoxia, highlighting the importance of amino acids in situations of metabolic stress [277,278]. BCAAs are reduced to keto acids, glutamate, and glutamine, substrates for the Krebs cycle during the Warburg effect. Glutamine is the fuel used by most cancer cells through macropinocytosis (Ras-mutant cells) [279-283].

The presence of amino acids allows cellular anabolism without the need for mitochondrial respiration (even in the presence of oxygen and glucose) in tumor and nontumor cells, as in ischemic wound healing [284-286]. Activated T cells use macropinocytosis for rapid amino acid uptake and immediate lymphoproliferative response, both immune and tumor, without the need for mitochondrial respiration [287-288]. There is a strong dependence between the levels of BCAAs and the growth of osteo and chondrosarcomas. Tumor biopsies in patients reveal overexpression of BCAAmetabolizing cytosolic aminotransferases [288]. These are prognostic markers as they are increased in more aggressive tumors [284-287]. BCAT1 (cytosolic aminotransferase for BCAAs) is overactive in chronic myeloid leukemia (CML) and overexpressed in chronic myeloid leukemia [282-284].

Physiological and pathological signaling mTORC1

mTORC1 controls cell activation in the presence of nutrients and trophic factors. Several studies indicate mTORC1 as the center of convergence of the various anabolic activation signals **[289-298]**. mTORC1 regulates gene transcription, and ribosomal translation suppresses autophagy and activates the mitochondrial machinery and the protein and lipid synthesis machinery **[299-304]**. mTORC1 is activated by the Rag (Ras adenosine guanidine) and Rheb (Ras homolog enriched in the brain) GTPases, each controlled by a pathway. Rag fixes mTORC1 on the lysosome surface in the presence of some amino acids (Raptor binding) **[305-309]** and Rheb, present in the lysosome, activates mTORC1 through trophic factors and glucose **[310]**.

Initial activation: the RAG pathway

Sancak et al (2010) [305] demonstrated that, in the presence of amino acids, free mTORC1 binds to the lysosome surface (via RAG) and is only later activated (via Rheb) by trophic factors. Mammals have four Rags (A, B, C, and D) that form A/B and C/D dimers. Raptor binding (from mTORC1) to RagD (present in lysosome) is activated by leucine and arginine [311-315]. Leucine mTOR activation is a consensus and the mechanisms are detailed in several reviews [316-320]. Recently Meng et al (2020) [316] reassessed the mTORC1 activation capacity of each amino acid and found that 10 amino acids (alanine, arginine, asparagine, glutamine, histidine, leucine, methionine, serine, threonine, and valine) are capable of binding to mTORC1 to the lysosome, but leucine, arginine, and methionine are the most potent, increasing S6K1 (a marker of ribosomal activity) in just 15 minutes, while glutamine, asparagine, and methionine take, for example, over an hour [317,318]. Activation of mTORC1 by glutamine is slower because it is RAGindependent [319]. There is also an exchange of glutamine for leucine by antitransport channels [320,321]. The discovery of Rags improved the understanding of mTORC1 regulation by amino acids [319,322].

In this sense, Han et al identified the LRS (leucyltRNA synthetase) sensor, which binds leucine to RAG-D, binding free cytosolic mTORC1 to the lysosome. LRS/RagD – mTORC1 associations are observed only in the presence of leucine. LRS-deficient mice are unable to attach mTORC1 to the lysosomal surface, even in the presence of leucine **[323]**. Other studies have also confirmed the role of the LRS sensor in mTORC1 preactivation **[324-329]**.

In addition to LRS, Sestrin 2 is another leucine

sensor, which inhibits the mTORC1 inhibitor kinase GATOR1 [329]. The presence of leucine disrupts the Sestrin 2 - Gator2 bond, releasing the GATOR 2 kinase that blocks GATOR1 and activates mTORC1. Wolfson et al tested the effect of amino acids on Sestrin 2 and found that only leucine (and not arginine) produces Sestrin 2-GATOR 2 dissociation with GATOR 1 blockade and mTORC1 activation [330-332]. Further to leucine, arginine and methionine are mTORC1 activators. Arginine uses the CASTOR1 pathway and methionine inhibits SAMTOR and GATOR 1/2 kinases. Both activate mTORC1 by inhibiting GATOR1 [333-338]. These sensors are present in lysosomes, mitochondria, rough endoplasmic reticulum, and Golgi complex, attaching mTORC1 to their surfaces in the presence of these amino acids [339-341].

In cell cultures, the removal of leucine or arginine prevents S6K activation, suggesting that both, in addition to methionine, are the main regulatory amino acids of mTORC1 activity. Wolfson et al (2016) [332] examined Sestrin-2/GATOR2 binding and observed that a lack of leucine, but not arginine, blocks mTORC1 activation. Thus, GATOR 2 acts more as an amino acid sensor than a mTORC1 activator [334]. There are other indirect mechanisms of mTORC1 activation by amino acids, such as the increase of intracellular calcium, by the mobilization of stores of the endoplasmic reticulum by SHP-2, activated by amino acids [342,343]. The acetyl-Coa metabolite enhances the RagD-Raptor interaction, binding mTORC1 to the lysosome [340]. Very recent studies have identified dysfunction of GATOR kinases 1 and 2 in the origin of some epilepsy, diseases called GATORpathies [344-350]. GATOR dysfunction alters mTORC1 activation, being part of the group of mTORC1 hyperactivity disorders (mTORpathies) **[351-354]**. mTORpathies occur in obesity, cancer, neurodegenerative diseases, and type II diabetes **[355-363]**, which will be discussed below.

Final activation: the Rheb pathway

With mTORC1 attached to the lysosome surface in the presence of amino acids, it binds to Rheb, activated by trophic factors and glucose **[364]**. Lysosomes are the ideal site of mTORC1 activation due to the high concentration of amino acids **[365-366]**. Other organelles, such as mitochondria and the Golgi complex, also attach mTORC1 to their surfaces in the presence of amino acids **[360]**. Recently, a fusion of the Golgi complex to lysosomes in the presence of amino acids has been described to transfer Rheb from the Golgi complex to lysosomes, increasing Rheb on the lysosomal surface to maximize mTORC1 activation in lysosomes **[362-365]**.

Rheb is the final pathway of mTORC1 activation that occurs in the presence of trophic factors and glucose. Trophic factors dissociate TSC-1 from TSC-2 (Tuberous Sclero Complex 1 and 2) **[319,320]**. During fasting, cytosolic TSC-2 is shifted to the lysosome, inactivating Rheb and turning off mTORC1 activity **[321-324]**. Tuberous sclerosis is the prototypical disease of TSC1/2 dysfunction, with consequent mTORC1 hyperactivation (**Figure 6**). It is a rare autosomal dominant with tumor formation in kidneys, heart, lungs, eyes, skin, and brain. About 80 to 90% of these people have epilepsy, developmental delay or mental retardation, behavioral disorders, and autism **[325-329]**.

Figure 6. Growth factors and amino acids activate mTORC1 via different pathways: amino acids (right) bind mTORC1 to the lysosome via RAG and via RHEB (left) it is the ultimate activator of mTORC1 by growth factors and glucose.



Source: Own authorship.

Therapeutic Interventions on mTORC1

Congenital mTOR hyperactivity disorders are often associated with difficult-tocontrol epilepsies [367-370] and mTOR inhibitors such as rapamycin, everolimus, and other rapanalogues are studied in the treatment of these epilepsies. As in tuberous sclerosis, autism, dementia, traumatic brain injury and stroke, and cancer [371-379]. The tuberous sclerotic complex TSC1/2 is inhibited in most tumors, with mTORC1 hyperactivity and rapid tumor growth [379,380]. Kaposi's sarcoma, for example, is a highly vascularized tumor and angiogenesis and growth are blocked with the use of rapamycin [381-386].

Mantle cell lymphomas (MCL) respond to rapamycin [387-390]. Rapanalogues are used in the treatment leukemias block of to the PI3K/AKT/TSC1/2/mTORC1 pathway, which is overactivated [387-392]. mTORC1 inhibitors are also used as immunosuppressants in certain kidney transplants [393], in the treatment of gliomas [394-399], and other cancers [400-405]. Recent data suggest imbalances of mTOR activity in diseases such as Parkinson's, Huntington's, Alzheimer's, frontotemporal dementia, and amyotrophic lateral sclerosis [406-409]. Rapanalogues and newer mTOR inhibitors can slow these neurodegenerative changes [410-413].

Alzheimer's disease (AD) is the most common form of dementia [414], characterized by the accumulation of proteins (β amyloid and tau) in brain tissue resulting in cognitive decline. Post-mortem studies of human AD brains indicate mTOR hyperactivation and overproduction of β amyloid and tau proteins [415-421]. Down syndrome is the most frequent chromosomal abnormality and it is associated with a congenital intellectual deficit. There is also an accumulation of tau and β amyloid proteins in brain tissue by mTOR hyperactivation [422-426]. Several studies of mTORC inhibition with rapamycin and rapanalogues suggest strong clinical potential in slowing the progression of cognitive impairment in Alzheimer's disease and Down syndrome [427-430].

Not only mTOR blockade has therapeutic action, but its activation is also a clinical target, as discussed in topic 3. mTORC1 activation is necessary for tissue regeneration such as muscle, liver, bone, intestine, etc. and its non-activation is a crucial factor. from the lack of CNS regeneration **[431-438]**. Neurons are particularly distinct, highly polarized cells with unique morphology and axon processes that can be thousands of times in length relative to the size of their cell bodies. The accumulation of cytoplasmic organelles along the axon forms small regional centers of synthesis, with some independence from the distant nucleus, which allows the synthesis of proteins necessary for reparative axonal budding or creation of new synapses in the CNS [438-441].

Also, mTOR activation results in axonal regeneration of optic nerve injury in animal models stimulated by growth factors **[442,443]**. mTORC1 activation occurs at the axon tip, on the surface of lysosomes present there, producing the macromolecules necessary for the formation of the growth cone and neoaxon expansion **[443,444]**.

Besides, mTORC1 activation may, however, be contrary to the recovery of the injured spinal cord, as activated astrocytes form extensive glial scars and block local axonal attempts at regeneration **[444]**. TRM models of hemisection and mTORC1 hyperactivation provoked by injection of interleukin 6 (PIP3/AKT/mTOR pathway activator) or by PTEN kinase blockade (PIP3/AKT/mTOR pathway inhibitor) provoked regenerative growth of the corticospinal tract in mice **[445-447]**.

Therefore, the current understanding of the molecular mechanisms of cell activation indicates that axonal organelles are fundamental organizational centers for growth cone progress and mTORC1 stimulation of axonal lysosomes may be one of the missing keys to CNS repair.

Conclusion

Amino acids are, in addition to building blocks of macromolecules, cellular activation signals. Essential amino acids are not produced by animals and leucine appears to be the main signaling amino acid. Mammals adjusted the cell activation and growth rate of their young by the leucine concentration of the milk produced. Several studies demonstrate the benefits of leucine supplementation in preventing sarcopenia, improving muscle and liver performance, as well as a possible neuroprotective role in head trauma and dementia. However, its excess, so common in the Western diet, is related to obesity, type II diabetes, neurodegenerative diseases, and cancer. The mTORC1 kinase integrates cellular activation stimuli from macro protein synthesis to epigenetic regulation. Controlling mTORC1 activity by consuming leucine can prevent, treat, or cause disease. A greater understanding of the regulatory effects of leucine and mTOR in unstable tissues such as tumors or fragile tissues such as the CNS are areas of great relevance and with extensive fields still to be explored.

Acknowledgement

Not applicable.

Funding

Not applicable.

Ethics approval

Not applicable.

Informed consent

Not applicable.

Data sharing statement

No additional data are available.

Conflict of interest

The authors declare no conflict of interest.

Similarity check

It was applied by Ithenticate@.

About the license

© The author(s) 2022. The text of this article is open access and licensed under a Creative Commons Attribution 4.0 International License.

References

- Jewell JL, Russell RC, Guan KL. Amino acid signalling upstream of mTOR. Nat Rev Mol Cell Biol 133-139, 2013.
- Zheng L, Zhang W, Zhou Y et al. (2016) Recent advances in understanding amino acid sensing mechanisms that regulate mTORC1. Int J Mol Sci. doi:10.3390/ijms17101636
- Zhang S, Zeng X, Ren M et al. Novel metabolic and physiological functions of branched chain amino acids: a review. J Animal Sci Biotechnol, (2017) doi: 10.1186/s40104-016-0139-z
- **4.** Siddik MAB, Shin AC. Recent progress on branched-chain amino acids in obesity, diabetes, and beyond. Endocrinol Metab):234-246, 2019.
- **5.** Sivanand S, Vander Heiden MG. Emerging roles for branched-chain amino acid metabolism in cancer. Cancer Cell. 2020:147-156.
- Wei Z, Liu X, Cheng C, Yu W, Yi P. Metabolism of amino acids in cancer. Front Cell Dev Biol. (2021). doi: 10.3389/fcell.2020.603837
- Peyrollier K, Hajduch E, Blair AS et al: L-leucine availability regulates phosphatidylinositol 3kinase, p70 S6 kinase and glycogen synthase

kinase-3 activity in L6 muscle cells: evidence for the involvement of the mammalian target of rapamycin (mTOR) pathway in the L-leucineinduced up-regulation of system A amino acid transport. Biochem J. 2000, 361-368.

- Krause U, Bertrand L, Maisin L et al: Signalling pathways and combinatory effects of insulin and amino acids in isolated rat hepatocytes. Eur J Biochem. 2002, 3742-3750.
- Lynch CJ, Halle B, Fujii H et al. Potential role of leucine metabolism in the leucine-signaling pathway involving mTOR. Am J Physiol Endocrinol Metab. 2003, 854-863.
- Bolster DR, Vary TC, Kimball SR, Jefferson LS: Leucine regulates translation initiation in rat skeletal muscle via enhanced eIF4G phosphorylation. J Nutr. 2004, 1704-1710.
- Vianna D, Teodoro GFR, Torres-Leal, FL, Tirapegui J. Protein synthesis regulation by leucine. Braz. J. Pharm. Sci., 2010, 29-36.
- Dodd KM, Tee AR. Leucine and mTORC1: a complex relationship. Am J Physiol Endocrinol Metab, 2012, 1329-342.
- Bröer S, BröerA. Amino acid homeostasis and signalling in mammalian cells and organisms. Biochem, 2017, J 1935–1963.
- Melnik, BC. Excessive leucine-mtorc1-signalling of cow milk-based infant formula: the missing link to understand early childhood obesity. Journ Obes. (2012) doi: 197653. 10.1155/2012/197653
- **15.** Melnik BC: Milk—A nutrient system of mammalian evolution promoting mtorc1-dependent translation. Int. J. Mol. Sci., 2015, 7048-17087.
- Melick CH, Jewell JL. Regulation of mTORC1 by upstream stimuli. Genes (2020). doi:10.3390/genes11090989
- Sengupta S, Peterson TR, Sabatini DM. Regulation of the mTOR complex 1 pathway by nutrients, growth factors, and stress. Mol Cell. 2010, 310-322.
- Sabatini DM. Twenty-five years obsessing over mTOR. PNAS, 2017, 1181811825.
- Melnik BC. Lifetime impact of cow's milk on overactivation of mTORC1: from fetal to childhood overgrowth, acne, diabetes, cancers and neurodegeneration. Biomolecules. (2021). doi:10.3390/biom11030404.
- Vezina C, Kudelski A, Sehgal S N. Rapamycin (AY-22,989), a new antifungal antibiotic I. Taxonomy of the producing streptomycete and isolation of the active principle. J. Antibiot. 1975, 721–726.
- **21.** Martel R R, Klicius J, Galet S. Inhibition of the immune response by rapamycin, a new antifungal

Vol 15 Suppl 2 Year 2022 International Journal of Nutrology (Official Journal of He ABRAN - Brazilian Association of Nutrology

antibiotic. Can J. Physiol. Pharmacol. 1977, 48–51.

- Eng CP, Sehgal SN, Vezina C. Activity of rapamycin (AY-22,989) against transplanted tumors. J. Antibiot. 1984, 1231–1237.
- **23.** Heitman J, Movva NR, Hall MN. Targets for cell cycle arrest by the immunosuppressant rapamycin in yeast. Science. 1991, 905-909.
- **24.** Zoncu R, Efeyan A, Sabatini DM. mTOR: from growth signal integration to cancer, diabetes and ageing. Nat Rev Mol Cell Biol. 2011, 21-35.
- Qian J, Su S, Liu P. Experimental approaches in delineating mTOR signaling. Genes. (2020). doi: 10.3390/genes11070738.
- **26.** Wullschleger S, Loewith R, Hall MN.TOR signaling in growth and metabolism. Cell, 2006, 471-484.
- 27. Sancak Y, Peterson TR, Shaul YD et al: The Rag GTPases bind raptor and mediate amino acid signaling to mTORC1. Science 2008, 1496–1501.
- 28. Tan VP, Miyamoto S. Nutrient-sensing mTORC1: integration of metabolic and autophagic signals. J. Mol. Cell Cardiol. 2016, 31–41.
- **29.** Dunlop EA; Tee AR. mTOR and autophagy: a dynamic relationship governed by nutrients and energy. Semin Cell Dev. 2014, 121–129.
- **30.** Aylett CH, Sauer E, Imseng S et al: Architecture of human mTOR complex. Science. 2016, 48-52.
- **31.** Efeyan A, Zoncu R, Sabatini DM. Amino acids and mTORC1: from lysosomes to disease. Trends Mol Med. 2012, 524-533.
- **32.** Efeyan, A.; Sabatini, D.M. Nutrients and growth factors in mTORC1 activation. Biochem. Soc. Trans. 902–905, 2013.
- **33.** Saxton RA, Sabatini DM. mTOR Signaling in Growth, Metabolism, and Disease. Cell. 960-976, 2017.
- **34.** Kim J, Guan KL. mTOR as a central hub of nutrient signalling and cell growth. Nat Cell Biol. 63-71, 2019.
- **35.** Jhanwar-Uniyal M, Wainwright JV, Mohan AL et al: Diverse signaling mechanisms of mTOR complexes: mTORC1 and mTORC2 in forming a formidable relationship. Adv Biol Regul. 51-62, 2019.
- **36.** Proud CG. A new link in the chain from amino acids to mTORC1 activation. Molecular Cell.7–8, 2011.
- **37.** Kim J, Guan KL: Amino acid signaling in TOR activation. Annu Rev Biochem., 2011, 1001-32.
- **38.** Dibble CC, Manning BD. Signal integration by mTORC1 coordinates nutrient input with biosynthetic output. Nat Cell Biol :555-556, 2013.
- **39.** Chantranupong L, Wolfson RL, Sabatini DM: Nutrient-sensing mechanisms across evolution.

Cell. 2015, 67-83.

- Biswas D, Duffley L, Pulinilkunnil T. Role of branched chain amino acid– catabolizing enzymes in intertissue signaling, metabolic remodeling, and energy homeostasis. FASEB J. 2019, 8711– 8731.
- Takahara T, Amemiya Y, Sugiyama R et al. Amino acid-dependent control of mTORC1 signaling: a variety of regulatory modes. J Biomed Sci. (2020); doi: 10.1186/s 12929-020-00679-2.
- **42.** Han JM, Jeong SJ, Park MC et al. Leucyl-tRNA synthetase is an intracellular leucine sensor for the mTORC1-signaling pathway. Cell. 410–424, 2012.
- **43.** Bar-Peled L, Sabatini DM. Regulation of mTORC1 by amino acids. Trends Cell Biol. 2014, 400–406.
- **44.** Jewell JL, Kim YC, Russell RC et al: Differential regulation of mTORC1 by leucine and glutamine. Science. 2015, 194–198.
- Wolfson RL, Sabatini DM. The Dawn of the age of amino acid sensors for the mTORC1 pathway. Cell Metab. 2017, 301–309.
- **46.** Son SM, Park SJ, Lee H et al. Leucine signals to mTORC1 via its metabolite acetyl-coenzyme A. Cell Metab. 2019, 192–201.
- Son SM, Park SJ, Stamatakou E et al. Leucine regulates autophagy via acetylation of the mTORC1 component raptor. Nat Commun. (2020). doi: 10.1038/s41467020-16886.
- Luo Y, Xu W, Li G, Cui W. Weighing in on mTOR complex 2 signaling: the expanding role in cell metabolismo. Oxidative Medicine and Cellular Longevity (2018) doi.org/10.1155/2018/7838647.
- Smith SF, Collins SE, Charest PG. Ras, PI3K and mTORC2 - three's a crowd? J Cell Sci. (2020). doi: 10.1242/jcs.234930.
- 50.Fu W, Hall MN. Regulation of mTORC2 signaling.
Genes (Basel).2020 doi:
10.3390/genes11091045.
- Baddour JA, Sousounis K, Tsonis PA. Organ repair and regeneration: An overview. Birth Defects Res. C Embryo Todday, 1–29, 2012.
- Cruz, B., Oliveira, A., Ventrucci, G et al. A leucinerich diet modulates the mTOR cell signalling pathway in the gastrocnemius muscle under different Walker-256 tumour growth conditions. BMC Cancer. (2019).doi.org/10.1186/s12885-0195448-0
- Wei X, Luo L, Chen J. Roles of mTOR signaling in tissue regeneration. Cells. (2019). doi:10.3390/cells8091075
- 54. Choi YJ, Di Nardo A, Kramvis I et al. Tuberous

sclerosis complex proteins control axon formation. Genes Dev. 2008;22(18):2485-2495. doi:10.1101/gad.1685008.

- **55.** 55. Han JM, Sahi M. TSC1/TSC2 signaling in the CNS. FEBBS Letters, 973-980, 2011.
- **56.** Ohtake Y, Hayat U, Li S. PTEN inhibition and axon regeneration and neural repair. Neural Regen Res. 1363-1368, 2015.
- 57. Zhang J, Yang D, Huang H et al: Coordination of necessary and permissive signals by PTEN inhibition for CNS axon regeneration. Front Neurosci (2018). doi.org/10.3389/fnins.2018.00558.
- Philips AM, Khan N. Amino acid sensing pathway: A major check point in the pathogenesis of obesity and COVID-19. Obes Rev. (2021) doi:10.1111/obr.13221.
- **59.** Wang CH, Chung FT, Lin SM et al. Adjuvant treatment with a mammalian target of rapamycin inhibitor, sirolimus, and steroids improves outcomes in patients with severe H1N1 pneumonia and acute respiratory failure. Crit Care Med. 313 321, 2014.
- **60.** Siddik MAB, Shin AC. Recent progress on branched-chain amino acids in obesity, diabetes, and beyond. Endocrinol Metab 234-246, 2019.
- **61.** Leenders M, van Loon LJ. Leucine as a pharmaconutrient to prevent and treat sarcopenia and type 2 diabetes. Nutrition Reviews, 675-689, 2011.
- **62.** Dodd KM, Tee AR. Leucine and mTORC1: a complex relationship. Am J Physiol Endocrinol Metab. 1329-1342, 2012.
- **63.** Martínez-Arnau FM, Fonfría-Vivas R, Cauli O. Beneficial effects of leucine supplementation on criteria for sarcopenia: A systematic review. Nutrients. (2019). doi: 10.3390/nu11102504.
- 64. Yoshimura Y, Bise T, Shimazu S et al. Effects of a leucine-enriched amino acid supplement on muscle mass, muscle strength, and physical function in post-stroke patients with sarcopenia: A randomized controlled trial. Nutrition. 1-6, 2019.
- **65.** Akan B. Influence of sarcopenia focused on critically ill patients. Acute Crit Care.2021, 15-21.
- **66.** Zhang Y, Guo K, LeBlanc RE et al: Increasing dietary leucine intake reduces dietinduced obesity and improves glucose and cholesterol metabolism in mice via multimechanisms. Diabetes.1647-1654, 2007.
- **67.** Binder E, Bermúdez-Silva FJ, André C et al: Leucine supplementation protects from insulin resistance by regulating adiposity levels. (2013).

doi.org/10.1371/journal.pone.0074705.

- **68.** Bloomgarden Z. Diabetes and branched-chain amino acids: What is the link? J Diabetes. (2018). doi: 10.1111/1753-0407.12645.
- 69. Almeida AP, Fortes FS, Silveira BKS et al. (2020). Branched-Chain amino acids intake is negatively related to body adiposity in individuals at cardiometabolic risk. Revista de Nutrição,doi.org/10.1590/1678-9865202033e190208.
- 70. Ye Z, Wang S, Zhang C, Zhao Y. Coordinated modulation of energy metabolism and inflammation by branched-chain amino acids and fatty acids. Front Endocrinol. 2020. doi: 10.3389/fendo.2020.00617.
- Duan Y, Li F, Li Y et al: The role of leucine and its metabolites in protein and energy metabolism. Amino Acids. (2016). doi: 10.1007/s00726-015-2067-1
- 72. Harris RA, Joshi M, Jeoung NH, Obayashi M. Overview of the molecular and biochemical basis of branched-chain amino acid catabolism. The Journal of nutrition. 2005:1527-30.
- **73.** Rogero MM, Tirapegui J. Aspectos atuais sobre aminoácidos de cadeia ramificada e exercício físico. Rev Bras Cienc Farm. 563-575, 2008.
- 74. Soares JD, Howell SL, Teixeira FJ, Pimentel GD. Dietary amino acids and immunonutrition supplementation in cancer-induced skeletal muscle mass depletion: A mini-review. Curr Pharm Des. 970-978, 2020.
- **75.** Maltais ML, Perreault K, Courchesne-Loyer A et al. Effect of resistance training and various sources of protein supplementation on body fat mass and metabolic profile in sarcopenic overweight older adult men: a pilot study. Int J Sport Nutr Exerc Metab. 71–77, 2016.
- **76.** Nabuco HCG, Tomeleri CM, Fernandes RR et al. Effect of whey protein supplementation combined with resistance training on body composition, muscular strength, functional capacity, and plasma-metabolism biomarkers in older women with sarcopenic obesity: a randomized, doubleblind, placebocontrolled trial. Clin Nutr ESPEN. 88–95, 2019.
- Mirzoev TM. Skeletal muscle recovery from disuse atrophy: protein turnover signaling and strategies for accelerating muscle regrowth. Int J Mol Sci. (2020). doi: 10.3390/ijms21217940.
- **78.** Bennett BT, Mohamed JS, Alway SE. The effects of calcium-β-hydroxy-βmethylbutyrate on aging-associated apoptotic signaling and muscle mass and function in unloaded but nonatrophied

extensor digitorum longus muscles of aged rats. Oxid Med Cell Longev. (2020). doi: 10.1155/2020/3938672.

- 79. Jakubowski JS, Nunes EA, Teixeira FJ et al: Supplementation with the leucine metabolite βhydroxy-β-methylbutyrate (hmb) does not improve resistance exercise-induced changes in body composition or strength in young subjects: a systematic review and meta-analysis. Nutrients. (2020). doi: 10.3390/nu12051523.
- 80. Gran P, Cameron-Smith D. The actions of exogenous leucine on mTOR signalling and amino acid transporters in human myotubes. BMC physiology.2011 Published online 10.1186/1472-6793-11-10.
- Liu H, Liu R, Xiong Y et al. Leucine facilitates the insulin-stimulated glucose uptake and insulin signaling in skeletal muscle cells: involving mTORC1 and mTORC2. Amino Acids. 2014, 1971-1979.
- Deldique L, Sanchez-Canedo C, Horman S et al: Antagonistic effects of leucine and glutamine on the mTOR pathway in myogenic C2C12 cells. Amino Acids. 147-155, 2008.
- Atherton PJ, Smith K, Etheridge T et al: Distinct anabolic signalling responses to amino acids in C2C12 skeletal muscle cells. Amino Acids. 2010, 1533-1539.
- **84.** Meijer A, Baquet A, Gustafson L et al. Mechanism of activation of liver glycogen synthase by swelling. J. Biol. Chem. 5823–5828, 1992.
- **85.** Vary T C, Jefferson S, Kimball SR: Amino acidinduced stimulation of translation initiation in rat skeletal muscle.Am. J. Physiol, 1077-1086, 1999.
- 86. Bolster DR, Vary TC, Kimball SR, Jefferson LS. Leucine regulates translation initiation in rat skeletal muscle via enhanced eIF4G phosphorylation. J Nutr. 1704-1710, 2004.
- **87.** Anthony JC, Anthony TG, Kimball SR et al: Orally administered leucine stimulates protein synthesis in skeletal muscle of postabsorptive rats in association with increased eIF4F formation. J. Nutr. 2000, 139-145.
- **88.** Proud C. mTOR-mediated regulation of translation factors by amino acids. Biochem Biophys Res Commun 2004, 429–436.
- **89.** Stipanuk MH: Leucine and protein synthesis: mTOR and beyond. Nutr. Rev. 122129, 2007.
- **90.** Peyrollier K, Hajduch E, Blair AS et al: L-leucine availability regulates phosphatidylinositol 3kinase, p70 S6 kinase and glycogen synthase kinase-3 activity in L6 muscle cells: evidence for the involvement of the mammalian target of

rapamycin (mTOR) pathway in the L-leucineinduced up-regulation of system A amino acid transport. Biochem J. 361-368, 2000.

- **91.** Guo CY, Yu MX, Dai JM et al: Roles of mitogenactivating protein kinase kinase kinase kinase-3 (MAP4k3) in preterm skeletal muscle satellite cell myogenesis and mammalian target of rapamycin complex 1 (mTORC1) activation regulation. Med Sci Monit. 3562-3570. 2017.
- **92.** Antunes D, Chowdhury A, Aich A et al. Overexpression of branched-chain amino acid aminotransferases rescues the growth defects of cells lacking the Barth syndrome-related gene TAZ1. J Mol Med (2019). doi: 10.1007/s00109-018-1728-4.
- D'Antona G, Ragni M, Cardile A et al. Branchedchain amino acid supplementation promotes survival and supports cardiac and skeletal muscle mitochondrial biogenesis in middle-aged mice. Cell Metab. (2010). doi: 10.1016/j.cmet.2010.08.016.
- **94.** Tedesco L, Rossi F, Ragni M et al. A special amino-acid formula tailored to boosting cell respiration prevents mitochondrial dysfunction and oxidative stress caused by doxorubicin in mouse cardiomyocytes. Nutrients. (2020). doi: 10.3390/nu12020282.
- **95.** Duan Y, Li F, Liu H et al. Nutritional and regulatory roles of leucine in muscle growth and fat reduction. Front Biosci :796-813, 2015.
- **96.** Duan Y, Li F, Li Y et al. The role of leucine and its metabolites in protein and energy metabolism. Amino Acids. 41-51, 2016.
- 97. Kamei Y, Hatazawa Y, Uchitomi R et al: S. Regulation of skeletal muscle function by amino acids. Nutrients. (2020) doi: 10.3390/nu12010261.
- **98.** Zhang Y, Guo K, LeBlanc RE et al: Increasing dietary leucine intake reduces dietinduced obesity and improves glucose and cholesterol metabolism in mice via multimechanisms. Diabetes.1647-1654, 2007.
- **99.** Valerio A, D'Antona G, Nisoli E. Branched-chain amino acids, mitochondrial biogenesis, and healthspan: an evolutionary perspective. Aging. 464-478, 2011.
- 100. Binder E, Bermúdez-Silva FJ, André C et al: Leucine supplementation protects from insulin resistance by regulating adiposity levels. (2013). doi.org/10.1371/journal.pone.0074705
- **101.** Liu R, Li H, Fan W et al. Leucine supplementation differently modulates branched-chain amino acid catabolism, mitochondrial function and metabolic

profiles at the different stage of insulin resistance in rats on high-fat diet. Nutrients. (2017). doi: 10.3390/nu9060565.

- **102.** Bloomgarden Z. Diabetes and branched-chain amino acids: What is the link? J Diabetes. (2018). doi: 10.1111/1753-0407.12645.
- **103.** Zhang L Li F, Guo Q et al: Leucine supplementation: a novel strategy for modulating lipid metabolism and energy homeostasis. Nutrients. (2020). doi: 10.3390/nu1205129
- **104.** Ye Z, Wang S, Zhang C, Zhao Y. Coordinated modulation of energy metabolism and inflammation by branched-chain amino acids and fatty acids. Front Endocrinol. 2020. doi: 10.3389/fendo.2020.00617.
- 105.ValerioT.Mitochondrialbiogenesis:pharmacologicalapproaches.CurrPharmDes.(2014).doi:10.2174/13816128203514091114211.
- **106.** Stancliffe RA: Role of beta-hydroxy-betamethylbutyrate (hmb) in leucine stimulation of mitochondrial biogenesis and fatty acid oxidation. Masters Theses University of Tennessee, 2012.
- 107. Liang C, Curry BJ, Brown PL, Zemel MB. Leucine Modulates Mitochondrial Biogenesis and SIRT1-AMPK Signaling in C2C12 Myotubes. J Nutr Metab. (2014). doi: 10.1155/2014/239750.
- 108. Sergi D, Naumovski N, Heilbronn LK et al. Mitochondrial (dys)function and insulin resistance: from pathophysiological molecular mechanisms to the impact of diet. Front Physiol. (2019).doi: 10.3389/fphys.2019.00532.
- **109.** Yu SB, Pekkurnaz G. Mechanisms orchestrating mitochondrial dynamics for energy homeostasis. J Mol Biol. 3922-3941, 2018.
- **110.** Tubbs E, Chanon S, Robert M et al. Disruption of mitochondria-associated endoplasmic reticulum membrane (mam) integrity contributes to muscle insulin resistance in mice and humans. Diabetes. 636-650, 2018.
- **111.** Giacomello M, Pyakurel A, Glytsou C. et al. The cell biology of mitochondrial membrane dynamics. Nat Rev Mol Cell Biol, 204–224, 2020.
- Baker M J, Frazier A E, Gulbis J M, Ryan M T. Mitochondrial proteinimport machinery: Correlating structure with function. Trends in Cell Biology , 456-464, 2007.
- 113. Wenz LS, Opaliński L, Wiedemann L, Becker T. Cooperation of protein machineries in mitochondrial protein sorting. Molecular Cell Research 11191129, 2015.
- **114.** Sun X, Zemel MB. Leucine modulation of mitochondrial mass and oxygen consumption in

skeletal muscle cells and adipocytes. Nutr Metab (2009). doi.org/10.1186/1743-7075-6-26.

- **115.** Craig DM, Ashcroft SP, Belew MY et al. Utilizing small nutrient compounds as enhancers of exercise-induced mitochondrial biogenesis. Front Physiol. 2015 doi: 10.3389/fphys.2015.00296.
- **116.** Skinner DM. The effect of leucine supplementation on mitochondrial biogenesis and mitochondrial protein synthesis in rats fed a high-fat diet. Honors Theses. University of Arkansas, 2015.
- **117.** Almeida AP, Fortes FS, Silveira BKS et al. (2020). Branched-Chain amino acids intake is negatively related to body adiposity in individuals at cardiometabolic risk. Revista de Nutrição,doi.org/10.1590/16789865202033e190 208.
- 118. Myers MJ, Shepherd DL, Durr AJ et al. The role of SIRT1 in skeletal muscle function and repair of older mice. J Cachexia Sarcopenia Muscle.929-949, 2019.
- **119.** Dam G, Aamann L, Vistrup H, Gluud LL. The role of Branched Chain Amino Acids in the treatment of hepatic Encephalopathy. J Clin Exp Hepatol:448-451, 2018.
- Muting D, Wortmann V. Amino acid metabolism in liver diseases. Dtsch Med Wochenschr. 1853– 1856,1956;
- **121.** Fischer J.E., Rosen H.M., Ebeid A.M. The effect of normalization of plasma amino acids on hepatic encephalopathy in man. Surgery. 77–91,1976.
- **122.** Román E, Torrades MT, Nadal MJ et al. Randomized pilot study: effects of an exercise programme and leucine supplementation in patients with cirrhosis. Digestive Diseases and Sciences. 1966-1975, 2014.
- 123. Gluud L, Dam G, Les I et al: Branched-chain amino acids for people with hepatic encephalopathy. Cochrane Database Syst Rev. 2017 18;5:CD001939
- 124. Yang YJ, Kim DJ. An overview of the molecular mechanisms contributing to musculoskeletal disorders in chronic liver disease: osteoporosis, sarcopenia, and osteoporotic sarcopenia. International Journal of Molecular Sciences. (2021). DOI: 10.3390/ijms22052604
- **125.** Holecek M, Kandar R, Sispera L, Kovarik M. Acute hyperammonemia activates branched-chain amino acid catabolism and decreases their extracellular concentrations: different sensitivity of red and white muscle. Amino Acids. 575– 584, 2011.
- **126.** Tedesco L, Corsetti G, Ruocco C et al. A specific

amino acid formula prevents alcoholic liver disease in rodents. Am J Physiol Gastrointest Liver Physiol. (2018). doi: 10.1152/ajpgi.00231.2017.

- **127.** Wei X, Luo L, Chen J. Roles of mTOR signaling in tissue regeneration. Cells. (2019). doi:10.3390/cells8091075
- **128.** Jefferson LS, Korner A: A direct effect of growth hormone on the incorporation of precursors into proteins and nucleic acids of perfused rat live Biochem. J. 1967, 826-832.
- **129.** Krause U, Bertrand L, Maisin L et al: Signalling pathways and combinatory effects of insulin and amino acids in isolated rat hepatocytes. Eur J Biochem. 2002, 3742-3750.
- **130.** Dennis MD, Baum JI, Kimball SR, Jefferson LS: Mechanisms involved in the coordinate regulation of mTORC1 by insulin and amino acids. J Biol Chem. 2011, 8287-8296.
- **131.** Prins ML. Glucose metabolism in pediatric traumatic brain injury. Childs Nerv Syst. 2017, 1711-1718.
- **132.** Bowman CE, Scafidi J, Scafidi S. Metabolic perturbations after pediatric TBI: It's not just about glucose. Exp Neurol, 2019, 74-84.
- **133.** Bernini A, Masoodi M, Solari D et al. Modulation of cerebral ketone metabolism following traumatic brain injury in humans. J Cereb Blood Flow Metab. 2020, 177-186.
- **134.** Hewton KG, Johal AS, Parker SJ. Transporters at the Interface between cytosolic and mitochondrial amino acid metabolism. Metabolites. (2021) doi:10.3390/metabo11020112.
- **135.** Rogero MM, Tirapegui J. Current aspects of branched chain amino acid and exercise. Revista Brasileira De Ciências Farmacêuticas, 563-575, 2008.
- **136.** Sperringer JE, Addington A, Hutson SM. Branched-chain amino acids and brain metabolism. Neurochem. 2017, 1697–1709.
- **137.** Spinelli JB, Haigis MC. The multifaceted contributions of mitochondria to cellular metabolism. Nat Cell Biol. 2018, 745-754.
- Niklison-Chirou MV, Agostini M, Amelio I, Melino G. Regulation of adult neurogenesis in mammalian brain. Int J Mol Sci. (2020). doi: 10.3390/ijms21144869.
- **139.** Aquilani R, Iadarola P, Contardi A et al. Branchedchain amino acids enhance the cognitive recovery of patients with severe traumatic brain injury. Arch Phys Med Rehabil. 1729-1735, 2005.
- **140.** Sharma B, Lawrence DW, Hutchison MG. Branched chain amino acids (bcaas) and traumatic brain injury: a systematic review. J

Head Trauma Rehabil. 33-45, 2018.

- **141.** Bowman CE, Scafidi J, Scafidi S. Metabolic perturbations after pediatric TBI: It's not just about glucose. Experimental Neurology, 74-84, 2019.
- **142.** Jeter CB, Hergenroeder GW, Ward NH et al. Human mild traumatic brain injury decreases circulating branched-chain amino acids and their metabolite levels. J Neurotrauma. 671-629, 2013.
- 143. Vespa P, Bergneider M, Hattori N et al. Metabolic crisis without brain ischemia is common after traumatic brain injury: a combined microdialysis and positron emission tomography study. J Cereb Blood Flow Metab. 763–774, 2005.
- 144. Carre E, Ogier M, Boret H et al. Metabolic crisis in severely head-injured patients: is ischemia just the tip of the iceberg? Front Neurol. (2013). doi: 10.3389/fneur.2013.00146.
- **145.** Agostini M, Romeo F, Inoue S et al. Metabolic reprogramming during neuronal differentiation. Cell Death Differ, 1502–1514, 2016.
- **146.** Koppel SJ, Swerdlow RH. Neuroketotherapeutics: A modern review of a century-old therapy. Neurochem Int. 114-125, 2018.
- 147. Maffezzini C, Calvo-Garrido J, Wredenberg A, Freyer C. Metabolic regulation of neurodifferentiation in the adult brain. Cell Mol Life Sci. 2020, 2483-2496.
- 148. Guzmán M, Blázquez C. Ketone body synthesis in the brain: possible neuroprotective effects. Prostaglandins Leukot Essent Fatty Acids. 287-92, 2004.
- **149.** White H, Venkatesh B. Clinical review: Ketones and brain injury. Crit Care (2011). doi.org/10.1186/cc10020.
- **150.** Barber CN, Raben DM. Lipid metabolism crosstalk in the brain: glia and neurons. Front Cell Neurosci. (2019). doi:10.3389/fncel.2019.00212.
- **151.** Smith QR, Takasato Y, Sweeney DJ, Rapoport SI: Regional cerebrovascular transport of leucine as measured by the in situ brain perfusion technique, J Cereb Blood Flow Metab. 1985, 300-311.
- **152.** Smith QR, Momma S, Aoyagi M, Rapoport SI. Kinetics of neutral amino acid transport across the blood-brain barrier. J. Neurochem. 1987, 1651– 165.
- **153.** Smith QR (1991). The blood-brain barrier and the regulation of amino acid uptake and availability to brain. Adv. Exp. Med. Biol.414–416, 1991
- **154.** Zaragoza R. Transport of amino acids across the blood-brain barrier. Front Physiol. (2020). doi:10.3389/fphys.2020.00973

Vol 15 Suppl 2 Year 2022 International Journal of Nutrology (Official Journal of Nutrology (Official Journal of Nutrology

- **155.** Martinez-Hernandez A, Bell KP, Norenberg MD. Glutamine synthetase: glial localization in brain. Science. 1977, 1356-1358.
- **156.** Yudkoff M. Brain metabolism of branched-chain amino acids. Glia. 1997, 92–98.
- **157.** Hutson SM, Lieth E, LaNoue KF. Function of leucine in excitatory neurotransmitter metabolism in the central nervous system. J Nutr. 2001, 846-850.
- **158.** Yudkoff M, Daikhin Y, Nissim I et al. Brain amino acid requirements and toxicity: the example of leucine. J Nutr.1531-1538, 2005.
- **159.** Nissim I, States B, Yudkoff M, Segal S. Characterization of amino acid metabolism by cultured rat kidney cells: study with 15N. Am J Physiol. 1987, 1243-1252.
- **160.** Hausmann ON. Post-traumatic inflammation following spinal cord injury. Spinal Cord. 2003, 369–378.
- **161.** Park E, Alexander A. Velumian AA, Fehlings MG: The role of excitotoxicity in secondary mechanisms of spinal cord injury: a review with an emphasis on the implications for white matter degeneration. Journal of Neurotrauma. 754-774, 2004.
- 162. Lewerenz J, Maher P. Chronic glutamate toxicity in neurodegenerative diseases—what is the evidence? Front Cell Neurosci. (2015). Doi 10.3389/fnins.2015.00469
- **163.** Kirdajova DB, Kriska J, TureckovaJ, Anderova Ischemia-triggered glutamate excitotoxicity from the perspective of glial cells. Front Cell Neurosc (2020). doi.org/10.3389/fncel.2020.00051.
- **164.** Shambaugh GE 3rd, Koehler RA. Fetal fuels VI. Metabolism of alphaketoisocaproic acid in fetal rat brain. Metabolism. 421-427, 1983.
- **165.** Yudkoff M. Interactions in the metabolism of glutamate and the branchedchain amino acids and ketoacids in the CNS. Neurochem Res. 10-18, 2017.
- **166.** Yudkoff M, Daikhin Y, Nelson D et al: Neuronal metabolism of branchedchain amino acids: flux through the aminotransferase pathway in synaptosomes. J Neurochem. 2136–2145, 1996.
- **167.** Araque A, Parpura V, Sanzgiri RP, Haydon PG. Tripartite synapses: glia, the unacknowledged partner. Trends Neurosci. 208-215, 1999.
- **168.** Syková E. Extrasynaptic volume transmission and diffusion parameters of the extracellular space. Neurosci 861–876, 2004.
- **169.** Volterra A, Meldolesi J. Astrocytes, from brain glue to communication elements: the revolution continues. Nat Rev Neurosci , 626–640, 2005.

- **170.** Schafer DP, Lehrman EK, Stevens B. The "quadpartite" synapse: microglia-synapse interactions in the developing and mature CNS. Glia. 24-36, 2013.
- Cabral A, Portela R, Tasso T e col: Doenças dos aminoácidos de cadeia ramificada. Acta Med Port. 659-665, 1998.
- 172. Soldin SJ, Brugnara C, Wong EC: Pediatric Reference Intervals. 5^a ed. Washington.AACC Press, 2005.
- 173. Chuang DT, Chuang JL, Wynn RM. Lessons from genetic disorders of branched-chain amino acid metabolism. J Nutr. 243-249, 2006.
- 174. Hoffmann B, Helbling C, Schadewaldt P et al. Impact of longitudinal plasma leucine levels on the intellectual outcome in patients with classic MSUD. Pediatr Res 17–20, 2006.
- 175. Valadares ER. Leucinose: Doença do xarope de bordo. In: Martins AM, organizador. Protocolo brasileiro de dietas: erros inatos do metabolismo. São Paulo: Segmento Farma, 53-58, 2007.
- **176.** Zinnanti WJ, Lazovic J: Interrupting the mechanisms of brain injury in a model of maple syrup urine disease encephalopathy. J Inherit Metab Dis. 71-79, 2012.
- Manoli I, Venditti CP. Disorders of branched chain amino acid metabolism. Transl Sci Rare Dis. 91-110, 2016.
- 178. Camandola S, Mattson MP. Brain metabolism in health, aging, and neurodegeneration. EMBO J. 1474-1492, 2017.
- **179.** Jensen NJ, Wodschow HZ, Nilsson M, Rungby J. Effects of ketone bodies on brain metabolism and function in neurodegenerative diseases. Int J Mol Sci. (2020). doi: 10.3390/ijms21228767.
- 180. Polis B, Samson AO. Role of the metabolism of branched-chain amino acids in the development of Alzheimer's disease and other metabolic disorders. Neural Regen Res. 1460-1470, 2020.
- 181. Socha E, Kośliński P, Koba M et al. Serum amino acid profiles in patients with mild cognitive impairment and in patients with mild dementia or moderate dementia. Amino Acids. 97-109, 2021.
- 182. Shang X, Hill E, Li Y, He M. Energy and macronutrient intakes at breakfast and cognitive declines in community-dwelling older adults: a 9year follow-up cohort study. Am J Clin Nutr. (2021). doi: 10.1093/ajcn/nqaa403.
- 183. Sato H, Takado Y, Toyoda S et al: Neurodegenerative processes accelerated by protein malnutrition and decelerated by essential amino acids in a tauopathy mouse model. (2021). Science Advances.

doi/abs/10.1126/sciadv.abd5046

- **184.** Oddo S.The role of mTOR signaling in Alzheimer disease. Front Biosci. 941-952, 2012.
- **185.** Tramutola A, Triplett JC, Di Domenico F, Alteration of mTOR signaling occurs early in the progression of Alzheimer disease (AD): analysis of brain from subjects with pre-clinical AD, amnestic mild cognitive impairment and late-stage AD.J Neurochem. 739-749, 2015.
- **186.** Crino PB. The mTOR signalling cascade: paving new roads to cure neurological disease. Nat Rev Neurol. 379-392, 2016.
- **187.** Li H, Ye D, Xie W et al: Defect of branched-chain amino acid metabolism promotes the development of Alzheimer's disease by targeting the mTOR signaling. Biosci Rep. (2018). doi: 10.1042/BSR20180127.
- **188.** Norwitz NG, Querfurth H. mTOR mysteries: nuances and questions about the mechanistic target of rapamycin in neurodegeneration. Front Neurosci. (2020). doi: 10.3389/fnins.2020.00775
- **189.** Park KK, Liu K, Hu Y, et al. Promoting axon regeneration in the adult CNS by modulation of the PTEN/mTOR pathway. Science. 963-966, 2008
- **190.** Liu K, Lu Y, Lee JK et al: PTEN deletion enhances the regenerative ability of adult corticospinal neurons. Nat Neurosci. 1075-1081, 2010.
- 191. Suzuki H, Yamashiro D, Ogawa S et al: Intake of seven essential amino acids improves cognitive function and psychological and social function in middle-aged and older adults: a double-blind, randomized, placebo-controlled trial. Front Nutr. (2020). doi: 10.3389/fnut.2020.586166.
- **192.** Tynkkynen J, Chouraki V, van der Lee SJ et al: Association of branched-chain amino acids and other circulating metabolites with risk of incident dementia and Alzheimer's disease: A prospective study in eight cohorts. Alzheimers Dement. (2018). doi: 10.1016/j.jalz.2018.01.003
- **193.** Fernando WMADB, Rainey-Smith SR, Gardener SL et al: Associations of dietary protein and fiber intake with brain and blood amyloid-β. J Alzheimers Dis. (2018) doi: 10.3233/JAD-170742.
- **194.** Hanger DP, Byers HL, Wray S, et al. Novel phosphorylation sites in tau from Alzheimer brain support a role for casein kinase 1 in disease pathogenesis. J Biol Chem. (2007). doi: 10.1074/jbc.M703269200.
- **195.** Croft CL, Kurbatskaya K, Hanger DP, Noble W. Inhibition of glycogen synthase kinase-3 by BTA-EG4 reduces tau abnormalities in an organotypic brain slice culture model of Alzheimer's disease.

Sci Rep. (2017). doi: 10.1038/s41598017-07906-

- 196. Yang L, Wang H, Liu L, Xie A. The role of insulin/IGF-1/PI3K/Akt/GSK3beta signaling in Parkinson's disease dementia. Front Neurosci. (2018). DOI=10.3389/fnins.2018.00073
- **197.** Xu F, NaL, Li Y. et al. Roles of the PI3K/AKT/mTOR signalling pathways in neurodegenerative diseases and tumours. Cell Biosci (2020). doi.org/10.1186/s13578-020-00416-0.
- **198.** Caccamo A, Magrì A, Medina DX et al. mTOR regulates tau phosphorylation and degradation: implications for Alzheimer's disease and other tauopathies. Aging Cell. 370-80, 2013.
- 199. Kitagishi Y, Nakanishi A, Ogura Y, Matsuda S. Dietary regulation of PI3K/AKT/GSK-3beta pathway in Alzheimer's disease. Alzheimers Res Ther (2014) doi: 10.1186/alzrt265
- **200.** Episcopo F, Drouin-Ouellet J, Tirolo C et al: GSK-3β-induced Tau pathology drives hippocampal neuronal cell death in Huntington's disease: involvement of astrocyte-neuron interactions. Cell Death Dis. (2016). doi: 10.1038/cddis.2016.104.
- 201. Duda P, Wiśniewski J, Wójtowicz T e col. Targeting GSK3 signaling as a potential therapy of neurodegenerative diseases and aging. Expert Opin Ther Targets. 833-848, 2018.
- **202.** Hosios AM, Hecht VC, Danai LV et al. Amino acids rather than glucose account for the majority of cell mass in proliferating mammalian cells. Dev Cell.540-549, 2016.
- **203.** Brandhorst S, Longo VD. Fasting and caloric restriction in cancer prevention and treatment. Recent Results Cancer Res. 241-266, 2016.
- **204.** Ananieva EA, Wilkinson AC. Branched-chain amino acid metabolism in cancer. Curr Opin Clin Nutr Metab Care.64-70, 2018.
- 205. Brandhorst S, Longo VD. Protein quantity and source, fasting-mimicking diets, and longevity. Adv Nutr. 340-350, 2019.
- 206. Martin SB, Reiche WS, Fifelski NA et al: Leucine and branched-chain amino acid metabolism contribute to the growth of bone sarcomas by regulating AMPK and mTORC1 signaling. Biochem J. 1579-1599, 2020.
- **207.** Bröer S. Amino acid transport across mammalian intestinal and renal epithelia. Physiol Rev. 249–286, 2008.
- **208.** Taylor PM. Role of amino acid transporters in amino acid sensing. Am J Clin Nutr. 223-230, 2014.
- **209.** Bröer S, Fairweather SJ. Amino Acid Transport Across the Mammalian Intestine. Compr Physiol,

Vol 15 Suppl 2 Year 2022 International Journal of Nutrology (Official Journal of Nutrology (Official Journal of Nutrology (Official Journal of Nutrology)

343-373, 2018.

- **210.** Nicklin P, Bergman P, Zhang B. Bidirectional transport of amino acids regulates mTOR and autophagy. Cell. 2009 521-534, 2009.
- **211.** Palm W, Park Y, Wright K et al. The utilization of extracellular proteins as nutrients is suppressed by mTORC1. Cell. 259-270, 2015.
- **212.** Chen R, Zou Y, Mao D et al. The general amino acid control pathway regulates mTOR and autophagy during serum/glutamine starvation. J Cell Biol.173-182, 2014.
- 213. Bhutia YD, Babu E, Ramachandran S, Ganapathy V. Amino acid transporters in cancer and their relevance to "glutamine addiction": novel targets for the design of a new class of anticancer drugs. Cancer Res. 1782–1788, 2015.
- **214.** Cormerais Y, Vučetić M, Parks SK, Pouyssegur J. Amino acid transporters are a vital focal point in the control of mTORC1 signaling and cancer. Int J Mol Sci. (2020). doi:10.3390/ijms22010023.
- **215.** Errasti-Murugarren E, Palacín M. Heteromeric Amino Acid Transporters in Brain: from Physiology to Pathology. Neurochem Res (2021) doi: 10.1007/s11064-021-03261-w.
- **216.** Wang Q, Holst J. The L-type amino acid transporter family serves as an important route for EAA entry into cells and consists of four members (LAT1–4). L-type amino acid transport and cancer: targeting the mTORC1 pathway to inhibit neoplasia. J Cancer Res. 1281-94, 2015.
- **217.** Bhutia YD, Ganapathy V. Glutamine transporters in mammalian cells and their functions in physiology and cancer. Biochim Biophys Acta. 2531-2539, 2016.
- **218.** Lukey MJ, Katt WP, Cerione RA. Targeting amino acid metabolism for cancer therapy. Drug Discov Today. 796-804, 2017.
- **219.** Zhang J, Xu Y, Li D et al. Review of the correlation of LAT1 with diseases: mechanism and treatment. Front Chem. (2020) doi:10.3389/fchem.2020.564809.
- **220.** Jewell JL, Kim YC, Russell RC et al. Differential regulation of mTORC1 by leucine and glutamine. Science. 194-198, 2015.
- 221. Drummond MJ, Glynn EL, Fry CS et al: An increase in essential amino acid availability upregulates amino acid transporter expression in human skeletal muscle. Am J Physiol Endocrinol Metab. 1011-1018, 2010.
- 222. Hayashi K, Jutabha P, Endou H et al. LAT1 is a critical transporter of essential amino acids for immune reactions in activated human T cells. J Immunol. (2013). doi: 10.4049/jimmunol.

1300923

- 223. Salisbury TB, Arthur S. The regulation and function of the L-type amino acid transporter 1 (LAT1) in cancer. Int J Mol Sci. (2018). doi: 10.3390/ijms19082373
- **224.** Tărlungeanu DC, Deliu E, Dotter CP et al. Impaired amino acid transport at the blood brain barrier is a cause of autism spectrum disorder. Cell. 1481-1494, 2016.
- **225.** Suzuki A, Iwata J. Amino acid metabolism and autophagy in skeletal development and homeostasis. Bone. (2021). doi: 10.1016/j.bone.2021.
- **226.** Ozaki K, Yamada T, Horie T et al. The L-type amino acid transporter LAT1 inhibits osteoclastogenesis and maintains bone homeostasis through the mTORC1 pathway. Sci Signal. (2019). doi:10.1126/scisignal.aaw3921.
- **227.** Wang Q, Tiffen J, Bailey CG et al: Targeting amino acid transport in metastatic castration-resistant prostate cancer: effects on cell cycle, cell growth, and tumor development. Journal of the National Cancer Institute. 9-21, 2013.
- **228.** Marshall AD, van Geldermalsen M, Otte N J et al: LAT1 is a putative therapeutic target in endometrioid endometrial carcinoma. Int. J. Cancer, 2529–2539, 2016.
- **229.** Cormerais Y, Pagnuzzi-Boncompagni M, Schrötter S et al: Inhibition of the amino-acid transporter LAT1 demonstrates anti-neoplastic activity in medulloblastoma. J Cell Mol Med. 2711–2718, 2019.
- **230.** Häfliger P, Charles RP. The L-Type Amino Acid Transporter LAT1-An emerging target in cancer. Int J Mol Sci. (2019). doi:10.3390/ijms20102428.
- **231.** Sato K, Miyamoto M, Takano M et al: Significant relationship between the LAT1 expression pattern and chemoresistance in ovarian clear cell carcinoma. Virchows Arch 701–710, 2019.
- **232.** Kaira K, Kawashima O, Endoh H et al. Expression of amino acid transporter (LAT1 and 4F2hc) in pulmonary pleomorphic carcinoma. Hum. Pathol. 2142–149, 2019.
- **233.** Lu JJ, Li P, Yang Y et al: Prognostic value of LAT-1 status in solid cancer: A systematic review and meta-analysis. PLoS One. (2020). doi: 10.1371/journal.pone.0233629.
- 234. Sinclair LV, Rolf J, Emslie E et al: Control of amino-acid transport by antigen receptors coordinates the metabolic reprogramming essential for T cell differentiation. Nat Immunol. 2013, 500-508.
- 235. Ananieva EA, Patel CH, Drake CH et al. Cytosolic

branched chain aminotransferase (BCATc) regulates mTORC1 signaling and glycolytic metabolism in CD4+ T cells. J Biol Chem. 18793-18804, 2014.

- **236.** Wang Q, Holst J. L-type amino acid transport and cancer: targeting the mTORC1 pathway to inhibit neoplasia. J Cancer Res. 1281-1294, 2015.
- **237.** Ananieva EA, Powell JD, Hutson SM. Leucine metabolism in T cell activation: mTOR signaling and beyond. Adv Nutr. 798-805, 2016.
- **238.** Häfliger P, Charles RP. The L-type amino acid transporter LAT1: An emerging target in câncer. (2019). Int J Mol Sci. doi: 10.3390/ijms20102428.
- 239. Danay Cibrian D, Castillo-Gonzalez R, Fernandez-Gallego N et al: Targeting L-type amino acid transporter 1 in innate and adaptive T cells efficiently controls skin inflammation. J Allergy Clin Immunol (2020) doi.org/10.1016/j.jaci.2019.09.025
- 240. Puris E, Gynther M, Auriola S. et al. L-Type amino acid transporter 1 as a target for drug delivery. Pharm Res (2020). https://doi.org/10.1007/s11095-02002826-8.
- **241.** Hayashi K, Kaminuma O, Nishimura T et al: LAT1specific inhibitor is effective against T cellmediated allergic skin inflammation. Allergy. 463– 467, 2020.
- **242.** Kaminuma O, Nishimura T, Saeki M et al: (2020). L-type amino acid transporter 1 (LAT1) specific inhibitor is effective against T cell-mediated nasal hyperresponsiveness. Allergology International. DOI: 69. 10.1016/j.alit.2019.12.006.
- 243. Ito D, Miura K, Saeki M et al: L-type amino acid transporter 1 inhibitor suppresses murine Th2 cell-mediated bronchial hyperresponsiveness independently of eosinophil accumulation. Asia Pac Allergy. (2021). doi: 10.5415/apallergy.2021.11.e33
- 244. Hayashi K, Anza N: L-type amino acid transporter 1 as a target for inflammatory disease and cancer immunotherapy, Journal of Pharmacological Sciences, 2021,1347-8613.
- **245.** Yoshida S, Pacitto R, Inoki K, Swanson J. Macropinocytosis, mTORC1 and cellular growth control. Cell Mol Life Sci. 1227-1239, 2018.
- **246.** Hoeller O, Bolourani P, Clark J, et al. Two distinct functions for PI3kinases in macropinocytosis. J Cell Sci. 4296-4307, 2013.
- **247.** Kay RR, Williams TD, Paschke P. Amplification of PIP3 signalling by macropinocytic cups. Biochem J. 643-648. 2018.
- **248.** Salloum G, Jakubik CT, Erami Z et al: PI3Kβ is selectively required for growth factor-stimulated

macropinocytosis. Journal of Cell Science (2019). doi: 10.1242/jcs.231639

- 249. Swanson JA, Watts C Macropinocytosis.Trends Cell Biol. 424-428,1995.
- **250.** Swanson JA. Shaping cups into phagosomes and macropinosomes. Nat Rev Mol Cell Biol. 639-649, 2008.
- **251.** Yoshida S, Hoppe AD, Araki N, Swanson JA. Sequential signaling in plasma-membrane domains during macropinosome formation in macrophages. J Cell Sci. 3250-3561, 2009.
- **252.** Commisso C, Davidson SM, Soydaner-Azeloglu RG et al. Macropinocytosis of protein is an amino acid supply route in Ras-transformed cells. (2013) Nature. doi: 10.1038/nature12138.
- **253.** Palm W, Park Y, Wright K et al: The utilization of extracellular proteins as nutrients is suppressed by mTORC1. Cell , 259–270, 2015.
- **254.** Palm W. Metabolic functions of macropinocytosis. Philos Trans R Soc Lond B Biol Sci. (2019). doi: 10.1098/rstb.2018.0285.
- **255.** Shaojuan S, Yanan Z, Tingting D et al: The dual role of macropinocytosis in cancers: promoting growth and inducing methuosis to participate in anticancer therapies as targets (2021). Frontiers in Oncology. doi. 10.3389/fonc.2020.570108.
- **256.** Shibutani S, Okazaki H, Iwata H. Dynamindependent amino acid endocytosis activates mechanistic target of rapamycin complex 1 (mTORC1). J Biol Chem. 18052-18061, 2017.
- **257.** Swanson JA, Yirinec B, Burke E et al. Effect of alterations in the size of the vacuolar compartment on pinocytosis in J774.2 macrophages. J. Cell. Physiol. 195–201, 1986.
- **258.** Swanson JA, Burke E, Silverstein SC. Tubular lysosomes accompany stimulated pinocytosis in macrophages. J. Cell Biol. 1217–1222, 1987.
- **259.** Swanson JA, Yoshida S. Macropinosomes as units of signal transduction. Philos Trans R Soc Lond B Biol Sci. (2019). doi:10.1098/rstb.2018.015.
- 260. Shaw RJ, Cantley LC. Ras, PI3K and mTOR signalling controls tumour cell growth. Nature. 424–430, 2006.
- **261.** Yoshida S, Pacitto R, Yao Y et al: Growth factor signaling to mTORC1 by amino acid-laden macropinosomes. J Cell Biol. 159-172, 2015.
- **262.** Lewis WH. Pinocytosis. Johns Hopkins Hosp Bull, 17–26, 1931.
- **263.** Bridges D, Fisher K, Zolov SN et al: Rab5 proteins regulate activation and localization of target of rapamycin complex 1. J. Biol. Chem. 20913–20921, 2012.
- 264. Nagano M, Toshima JY, Siekhaus DE et al. Rab5-

mediated endosome formation is regulated at the trans-Golgi network. Commun Biol (2019). Doi. 10.1038/s42003-019-0670-5.

- **265.** Tang W, Tam JH, Seah C et al: Arf6 controls betaamyloid production by regulating macropinocytosis of the amyloid precursor protein to lysosomes. (2015). Mol Brain. doi: 10.1186/s13041-015-0129-7.
- 266. Zeineddine R, Yerbury JJ. The role of macropinocytosis in the propagation of protein aggregation associated with neurodegenerative diseases. (2015). Front Physiol. doi: 10.3389/fphys.2015.00277.
- 267. Kabayama H, Nakamura T, Takeuchi M et al: Ca2+ induces macropinocytosis via F-actin depolymerization during growth cone colapse. (2009) Mol. Cell Neurosci. doi: 10.1016/j.mcn.2008.08.009.
- **268.** Fitzner D, Schnaars M, van Rossum D et al. Selective transfer of exosomes from oligodendrocytes to microglia by macropinocytosis. J Cell Sci. 2011:447458, 2011.
- **269.** Lin XP, Mintern JD, Gleeson PA. Macropinocytosis in different cell types: similarities and differences. Membranes (Basel). (2020) doi:10.3390/membranes10080177.
- **270.** Petrova V, Nieuwenhuis B, Fawcett JW, Eva R. Axonal organelles as molecular platforms for axon growth and regeneration after injury. Int J Mol Sci. (2021). doi: 10.3390/ijms22041798.
- **271.** DeBerardinis RJ, Mancuso A, Daikhin E et al. Beyond aerobic glycolysis: transformed cells can engage in glutamine metabolism that exceeds the requirement for protein and nucleotide synthesis. Proc Natl Acad Sci USA. 19345-19350, 2007.
- **272.** Filipp FV, Ratnikov B, De Ingeniis J, et al: Glutamine-fueled mitochondrial metabolism is decoupled from glycolysis in melanoma. Pigment Cell Melanoma Res. 732-739, 2012.
- 273. Sun H, Chen L, Cao S et al: Warburg effects in cancer and normal proliferating cells: two tales of the same name. Genomics, Proteomics & Bioinformatics, 273-286, 2019.
- 274. Birkeland ES, Koch LM, Dechant R. Another consequence of the warburg effect? metabolic regulation of na+/h+ exchangers may link aerobic glycolysis to cell growth. Front Oncol. (2020). doi: 10.3389/fonc.2020.01561.
- **275.** DeBerardinis RJ, Chandel NS. We need to talk about the Warburg effect. Nat Metab. 127–129, 2020.
- **276.** Lieu EL, Nguyen T, Rhyne S, Kim J. Amino acids in cancer. Exp Mol Med. 15-30, 2020.

- 277. Keenan MM, Chi JT. Alternative fuels for cancer cells. Cancer J. 49-55, 2015.
- **278.** Sivanand S, Vander Heiden MG. Emerging roles for branched-chain amino acid metabolism in cancer. Cancer Cell.147-156, 2020.
- **279.** Ananieva EA, Wilkinson AC. Branched-chain amino acid metabolism in cancer. Curr Opin Clin Nutr Metab Care. 64-70, 2018.
- 280. Peng H, Wang Y, Luo W. Multifaceted role of branched-chain amino acid metabolism in cancer. Oncogene. 6747-6756, 2020.
- 281. Wei Z, Liu X, Cheng C, Yu W, Yi P. Metabolism of amino acids in cancer. Front Cell Dev Biol. (2021) doi: 10.3389/fcell.2020.603837.
- **282.** Hattori A, Tsunoda M, Konuma T et al: Cancer progression by reprogrammed BCAA metabolism in myeloid leukaemia. Nature. 500-504, 2017.
- **283.** Charpentier JC, Chen D, Lapinski PE et al: Macropinocytosis drives T cell growth by sustaining the activation of mTORC1. Nat Commun. (2020). doi: 10.1038/s41467-019-13997-3.
- **284.** Zheng YH, Hu WJ, Chen BC et al: BCAT1, a key prognostic predictor of hepatocellular carcinoma, promotes cell proliferation and induces chemoresistance to cisplatin. Liver Int, 1836-1847, 2016.
- 285. Song Y, Zhao B, Xu Y et al: Prognostic significance of branched-chain amino acid transferase 1 and CD133 in triple-negative breast cancer. BMC Cancer. (2020). doi: 10.1186/s12885-020-07070-2
- **286.** Luo L, Sun W, Zhu W et al: BCAT1 decreases the sensitivity of cancer cells to cisplatin by regulating mTOR-mediated autophagy via branched-chain amino acid metabolism. Cell Death Dis. (2021) doi: 10.1038/s41419-021-03456-7.
- **287.** Zhang L, Han J. Branched-chain amino acid transaminase 1 (BCAT1) promotes the growth of breast cancer cells through improving mTOR-mediated mitochondrial biogenesis and function. Biochem Biophys Res Commun. 224-231, 2017.
- **288.** Tabe Y, Lorenzi PL, Konopleva M. Amino acid metabolism in hematologic malignancies and the era of targeted therapy. Blood. 1014-1023, 2019.
- **289.** Manning BD, Cantley LC. Rheb fills a gap between TSC and TOR. Trends Biochem Sci. 573–576, 2003.
- **290.** Wullschleger S, Loewith R, Hall MN.TOR signaling in growth and metabolism. Cell,471-484, 2006.
- **291.** Avruch J, Long X, Ortiz-Vega S et al: Amino acid regulation of TOR complex 1. Am J Physiol Endocrinol Metab. (2009). doi:

10.1152/ajpendo.90645.2008.

- **292.** Dodd KM, Tee AR. Leucine and mTORC1: a complex relationship. Am J Physiol Endocrinol Metab. 1329-1342, 2012.
- **293.** Dibble CC, Manning BD: Signal integration by mTORC1 coordinates nutrient input with biosynthetic output. Nat Cell Biol, 555–564, 2013.
- **294.** Dunlop EA, Tee AR. mTOR and autophagy: a dynamic relationship governed by nutrients and energy. Semin. Cell Dev. Biol. 121–129, 2014.
- **295.** Ben-Sahra I, Manning BD. mTORC1 signaling and the metabolic control of cell growth. Curr Opin Cell Biol. 72-82, 2017.
- **296.** Blenis J. TOR, the gateway to cellular metabolism, cell growth, and disease. Cell. 10-13, 2017.
- **297.** Kim J, Guan KL. mTOR as a central hub of nutrient signalling and cell growth. Nat Cell Biol. 63-71, 2019.
- **298.** Liu GY, Sabatini DM. mTOR at the nexus of nutrition, growth, ageing and disease.Nat Rev Mol Cell Biol. 183-203, 2020.
- **299.** Thoreen CC, Chantranupong L, Keys HR et al: A unifying model for mTORC1-mediated regulation of mRNA translation. Nature.109-113, 2012.
- **300.** Jewell JL, Russell RC, Guan KL. Amino acid signalling upstream of mTOR. Nat Rev Mol Cell Biol, 133–139, 2013.
- **301.** Chantranupong L, Wolfson RL, Sabatini DM. Nutrient-sensing mechanisms across evolution. Cell. 67-83, 2015.
- **302.** Tan VP, Miyamoto, S. Nutrient-sensing mTORC1: integration of metabolic and autophagic signals. J. Mol. Cell Cardiol, 31–41, 2016.
- **303.** Li XZ, Yan XH. Sensors for the mTORC1 pathway regulated by amino acids. J Zhejiang Univ Sci B. 699-712, 2019.
- **304.** Szwed A, Kim E, Jacinto E. Regulation and metabolic functions of mTORC1 and mTORC2. Physiol Rev. 1371-1426, 2021.
- **305.** Sancak Y, Peterson TR, Shaul YD e col. Rag GTPases bind RAPTOR and mediate amino acid signaling to mTORC1. Science. 1496-1501, 2008.
- **306.** Li SC, Kane PM. The yeast lysosome-like vacuole: endpoint and crossroads. Biochim. Biophys. Acta, 650–663, 2009.
- **307.** Sancak Y, Bar-Peled L, Zoncu R et al: Ragulator-Rag complex targets mTORC1 to the lysosomal surface and is necessary for its activation by amino acids. Cell: 290-303, 2010.
- **308.** Efeyan A, Zoncu R, Sabatini DM. Amino acids and mTORC1: from lysosomes to disease. Trends Mol Med. 524-533, 2012.
- 309. Groenewoud MJ, Zwartkruis FJ. Rheb and Rags

come together at the lysosome to activate mTORC1. Biochem Soc Trans. 951-955, 2013.

- **310.** Zhu M, Wang X, Regulation of mTORC1 by small GTPases in response to nutrients, J Nutr, 1004–1011, 2020.
- **311.** Bar-Peled L, Sabatini DM. Regulation of mTORC1 by amino acids. Trends Cell Biol. 400-406, 2014.
- **312.** Dibble CC, Cantley LC. Regulation of mTORC1 by PI3K signaling. Trends Cell Biol. 545-555, 2015.
- **313.** Abraham RT. Making sense of amino acid sensing. Science. 128-129, 2015.
- **314.** Saxton RA, Sabatini DM. mTOR signaling in growth, metabolism, and disease. Cell. 960-976, 2017.
- **315.** Zhuang Y, Wang XX, He J et al: Recent advances in understanding of amino acid signaling to mTORC1 activation. Front Biosci. 971-982, 2019.
- **316.** Meng D, Yang Q, Wang H, et al. Glutamine and asparagine activate mTORC1 independently of Rag GTPases. J Biol Chem. 2890-2899, 2020.
- **317.** Laplante M, Sabatini DM. mTOR signaling in growth control and disease. Cell. 274-293, 2012.
- **318.** Zheng L, Zhang W, Zhou Y et al (2016). Recent advances in understanding amino acid sensing mechanisms that regulate mTORC1. Int J Mol Sci.doi:10.3390/ijms17101636.
- **319.** Takahara T, Amemiya Y, Sugiyama R. et al. Amino acid-dependent control of mTORC1 signaling: a variety of regulatory modes. J Biomed Sci (2020). doi.org/10.1186/s12929-020-00679-2.
- **320.** Nicklin P, Bergman P, Zhang B et al: Bidirectional transport of amino acids regulates mTOR and autophagy. Cell. 521-34, 2009.
- **321.** Jewell JL, Kim YC, Russell RC, Yu FX. Differential regulation of mTORC1 by leucine and glutamine.Science. 194-198, 2015.
- **322.** Li XZ, Yan XH. Sensors for the mTORC1 pathway regulated by amino acids. J Zhejiang Univ Sci B. 699-712, 2019.
- **323.** Han JM, Jeong SJ, Park MC, et a. Leucyl-tRNA synthetase is an intracellular leucine sensor for the mTORC1-signaling pathway. Cell. 410-424, 2012.
- **324.** Durán RV, Hall MN. Leucyl-tRNA synthetase: double duty in amino acid sensing. Cell Res. 1207-1209, 2012.
- **325.** Segev N, Hay N. Hijacking leucyl-tRNA synthetase for amino aciddependent regulation of TORC1. Mol Cell. 4-6, 2012.
- **326.** Yoon MS, Son K, Arauz E et al. Leucyl-tRNA synthetase activates vps34 in amino acid-sensing mtorc1 signaling. Cell Rep. 1510-1517, 2016.
- **327.** Yu YC, Han JM, Kim S. Aminoacyl-tRNA

Vol 15 Suppl 2 Year 2022 International Journal of Nutrology (Official Journal of Nutrology (Official Journal of Nutrology (Official Journal of Nutrology

synthetases and amino acid signaling. Biochim Biophys Acta Mol Cell Res (2021). doi: 10.1016/j.bbamcr.2020.118889.

- **328.** Melick C, Jewell JL. Regulation of mTORC1 by upstream stimuli. Genes (2020). doi:10.3390/genes11090989.
- **329.** Wolfson RL, Chantranupong L, Saxton RA et al. Sestrin2 is a leucine sensor for the mTORC1 pathway. Science. 43-48, 2016.
- **330.** Chantranupong L, Wolfson RL, Orozco JM et al: The sestrins interact with GATOR2 to negatively regulate the amino-acid-sensing pathway upstream of mTORC1. Cell Rep. 1-8, 2014.
- **331.** Kimball SR, Gordon BS, Moyer JE et al. Leucine induced dephosphorylation of Sestrin2 promotes mTORC1 activation. Cell. Signal. 896–906, 2016
- **332.** Wolfson RL, Sabatini DM. The dawn of the age of amino acid sensors for the mTORC1 pathway. Cell Metab. 301-309, 2017.
- **333.** Wang S, Tsun ZY, Wolfson RL et al. Lysosomal amino acid transporter SLC38A9 signals arginine sufficiency to mTORC1. Science. 188-194, 2015.
- **334.** Chantranupong L, Scaria SM, Saxton RA et al. The CASTOR proteins are arginine sensors for the mTORC1 pathway. Cell 165, 153–164, 2016.
- **335.** Ho A, Cho CS, Namkoong S et al. Biochemical basis of sestrin physiological activities. Trends Biochem Sci. 41, 621–632, 2016.
- **336.** Lee JH, Cho US, Karin M. Sestrin regulation of TORC1: is sestrin a leucine sensor? Sci. Signal. (2016). doi:10.1126/scisignal.aaf2885
- **337.** Gu X, Orozco JM, Saxton RA et al: SAMTOR is na S-adenosylmethionine sensor for the mTORC1 pathway. Science. 813–818, 2017.
- **338.** Gulati P, Gaspers LD, Dann SG et al: Amino acids activate mTOR complex 1 via Ca2+/CaM signaling to hVps34.Cell Metab.456-65, 2008.
- **339.** Mercan F, Lee H, Kolli S, Bennett AM. Novel role for SHP-2 in nutrientresponsive control of S6 kinase 1 signaling. Mol Cell Biol. 293-306, 2013.
- **340.** Son SM, Park SJ, Lee H et al. Leucine signals to mTORC1 via its metabolite acetyl-coenzyme A. Cell Metab. 192-201, 2019.
- **341.** Weckhuysen S, Marsan E, Lambrecq V et al; Involvement of GATOR complex genes in familial focal epilepsies and focal cortical dysplasia. Epilepsia. 994-1003, 2006.
- **342.** Baldassari S, Licchetta L, Tinuper P, Bisulli F, Pippucci T. GATOR1 complex: the common genetic actor in focal epilepsies. J Med Genet. 503-510, 2016.
- **343.** Baldassari S, Picard F, Verbeek NE et al: The landscape of epilepsyrelated GATOR1 variants.

Genet Med.: 398-408, 2019.

- **344.** Dawson RE, Nieto Guil AF, Robertson LJ et al. Functional screening of GATOR1 complex variants reveals a role for mTORC1 deregulation in FCD and focal epilepsy. Neurobiol Dis. (2020). doi: 10.1016/j.nbd.2019.104640.
- 345. Iffland PH 2nd, Carson V, Bordey A, Crino PB. GATORopathies: The role of amino acid regulatory gene mutations in epilepsy and cortical malformations. Epilepsia. 2163-2173, 2019.
- **346.** Specchio N, Pepi C, De Palma L et al: Neuroimaging and genetic characteristics of malformation of cortical development due to mTOR pathway dysregulation: clues for the epileptogenic lesions and indications for epilepsy surgery. Expert Rev Neurother. 1333-1345, 2021.
- **347.** Lee WS, Baldassari S, Stephenson SEM et al: Cortical Dysplasia and the mTOR pathway: How the study of human brain tissue has led to insights into epileptogenesis. Int J Mol Sci. (2022) doi: 10.3390/ijms23031344.
- **348.** Griffith JL, Wong M. The mTOR pathway in treatment of epilepsy: a clinical update. Future Neurol. 49-58, 2018.
- **349.** Karalis V, Bateup HS. Current approaches and future directions for the treatment of mTORopathies. Dev Neurosci. 143-158, 2021.
- **350.** Moloney PB, Cavalleri GL, Delanty N. Epilepsy in the mTORopathies: opportunities for precision medicine. Brain Commun. (2021). doi: 10.1093/braincomms/fcab222.
- 351. Nguyen LH, Bordey A. Convergent and divergent mechanisms of epileptogenesis in mTORopathies.
 Front Neuroanat. (2021) doi: 10.3389/fnana.2021.664695.
- **352.** Buerger C, DeVries B, Stambolic V. Localization of Rheb to the endomembrane is critical for its signaling function. Biochem Biophys Res Commun. 869-880, 2006.
- **353.** Dennis MD, Baum JI, Kimball SR, Jefferson LS. Mechanisms involved in the coordinate regulation of mTORC1 by insulin and amino acids. J Biol Chem. 8287-8296, 2011.
- **354.** Zoncu R, Efeyan A, Sabatini DM. mTOR: from growth signal integration to cancer, diabetes and ageing. Nat Rev Mol Cell Biol. 21-35, 2011.
- **355.** Hao F, Kondo K, Itoh T et al; Rheb localized on the Golgi membrane activates lysosome-localized mTORC1 at the Golgi-lysosome contact site. J Cell Sci. (2018). doi: 10.1242/jcs.208017.
- **356.** Angarola B, Ferguson SM. Coordination of Rheb lysosomal membrane interactions with mTORC1 activation. (2020) Faculty Rev.

Vol 15 Suppl 2 Year 2022 International Journal of Nutrology (Official Journal of Nutrology (Official Journal of Nutrology (Official Journal of Nutrology

doi:10.12688/f1000research.22367.1

- **357.** Makhoul C, Gleeson PA. Regulation of mTORC1 activity by the Golgi apparatus. Faculty Rev (2021) DOI: 10.12703/r/10-50. PMID: 34195689; PMCID: PMC8204759.
- **358.** Demetriades C, Doumpas N, Teleman AA. Regulation of TORC1 in response to amino acid starvation via lysosomal recruitment of TSC2. Cell, 786799, 2014.
- **359.** Menon S, Dibble CC, Talbott G et al: Spatial control of the TSC complex integrates insulin and nutrient regulation of mTORC1 at the lysosome. Cell. 771–785, 2014.
- **360.** Demetriades C, Plescher M, Teleman AA. Lysosomal recruitment of TSC2 is a universal response to cellular stress. Nat Commun. (2016) doi:10.1038/ncomms10662.
- **361.** Carroll B, Maetzel D, Maddocks OD et al. Control of TSC2-Rheb signaling axis by arginine regulates mTORC1 activity. Elife (2016). doi:10.7554/eLife.11058
- **362.** Orlova KA, Crino PB. The tuberous sclerosis complex. Ann NY Acad Sci. 87-105, 2010.
- **363.** Wong M. Mammalian target of rapamycin (mTOR) inhibition as a potential antiepileptogenic therapy: From tuberous sclerosis to common acquired epilepsies. Epilepsia. 27-36, 2010.
- **364.** Curatolo P, Moavero R, de Vries PJ. Neurological and neuropsychiatric aspects of tuberous sclerosis complex. The Lancet. Neurology. 2733-745, 2015.
- **365.** Leclezio L, de Vries PJ. Advances in the treatment of tuberous sclerosis complex. Curr Opin Psychiatry. 113-120, 2015.
- **366.** Specchio N, Pietrafusa N, Trivisano M et al. Autism and epilepsy in patients with tuberous sclerosis complex. Frontiers in Neurology. (2020). DOI: 10.3389/fneur.2020.00639.
- **367.** Wong M. Mammalian target of rapamycin (mTOR) pathways in neurological diseases. Biomed J. 40-50, 2013.
- **368.** Liu J, Reeves C, Michalak Z et al. Evidence for mTOR pathway activation in a spectrum of epilepsy-associated pathologies. Acta Neuropathol Commun. (2014). doi: 10.1186/2051-5960-2-71.
- **369.** Zhong S, Zhao Z, Xie W et al: GABAergic interneuron and neurotransmission are mTOR-dependently disturbed in experimental focal cortical dysplasia. Mol Neurobiol. 156-169, 2021.
- **370.** Guertin DA, Sabatini DM. Defining the role of mTOR in cancer, Cancer Cell, 9-22, 2007.
- 371. Laplante M, Sabatini DM. mTOR signaling in

growth control and disease. Cell. 274–293, 2012.

- **372.** Krueger DA, Wilfong AA, Holland-Bouley K et al: Everolimus treatment of refractory epilepsy in tuberous sclerosis complex. Ann Neurol. 679-687, 2013.
- **373.** Benjamin D, Hall MN. mTORC1: turning off is just as important as turning on.Cell. 627-628, 2014.
- 374. Liu J, Reeves C, Michalak Z et al. Evidence for mTOR pathway activation in a spectrum of epilepsy-associated pathologies. Acta Neuropathol Commun (2014). doi: 10.1186/2051-5960-2-71.
- **375.** Crino PB. The mTOR signalling cascade: paving new roads to cure neurological disease. Nat Rev Neurol. 379-92, 2016.
- **376.** Blenis J. TOR, the gateway to cellular metabolism, cell growth, and disease. Cell. 10-13, 2017.
- 377. Wang F, Chen F, Wang G et al. Rapamycin provides anti-epileptogenic effect in a rat model of post-traumatic epilepsy via deactivation of mTOR signaling pathway. Exp Ther Med. 4763-4770, 2018.
- 378. Hillmann P, Fabbro D. PI3K/mTOR pathway inhibition: opportunities in oncology and rare genetic diseases. Int J Mol Sci. (2019) doi:10.3390/ijms20225792.
- **379.** Pópulo H, Lopes JM, Soares P. The mTOR signalling pathway in human cancer. Int J Mol Sci.1886-918, 2012.
- **380.** Hosios AM, Hecht VC, Danai LV et al: Amino acids rather than glucose account for the majority of cell mass in proliferating mammalian cells. Dev Cell. 540-549, 2016.
- **381.** Gao X, Zhang Y, Arrazola P et al. TSC tumour suppressor proteins antagonize amino-acid-TOR signalling. Nat Cell Biol. 699-704, 2002.
- **382.** Guba M, von Breitenbuch P, Steinbauer M et al. Rapamycin inhibits primary and metastatic tumor growth by antiangiogenesis: involvement of vascular endothelial growth factor. Nat Med. 128-35, 2002.
- **383.** Rupertus K, Dahlem C, Menger MD et al. Rapamycin inhibits hepatectomy-induced stimulation of metastatic tumor growth by reduction of angiogenesis, microvascular blood perfusion, and tumor cell proliferation. Ann Surg Oncol. 2629-2637, 2009.
- 384. Álvarez-García O,García-Lopez E, Loredo V et al. Rapamycin induces growth retardation by disrupting angiogenesis in the growth plate. Kidney International, 561 – 568, 2010.
- **385.** Roy D, Sin SH, Lucas A et al. mTOR inhibitors block Kaposi sarcoma growth by inhibiting

essential autocrine growth factors and tumor angiogenesis. Cancer Res. 2235-2246, 2013.

- **386.** Faes S, Demartines N, Dormond O. Mechanistic target of rapamycin inhibitors in renal cell carcinoma: potential, limitations, and perspectives. Front Cell Dev Biol. (2021) doi: 10.3389/fcell.2021.636037
- **387.** Witzig TE, Geyer SM, Ghobrial I et al. Phase II trial of single-agent temsirolimus (CCI-779) for relapsed mantle cell lymphoma. J Clin Oncol. 53475356, 2005.
- **388.** Hess G. Temsirolimus for the treatment of mantle cell lymphoma. Expert Rev Hematol. 631-40, 2009.
- **389.** Lee JS, Vo TT, Fruman DA. Targeting mTOR for the treatment of B cell malignancies. Br J Clin Pharmacol. 1213-1228, 2016.
- **390.** Hess G, Wagner K, Keller U, et al. Final results of a phaseII/IIIi trial of the combination bendamustine and rituximab with temsirolimus (bert) in relapsed mantle cell lymphoma and follicular lymphoma. Hemasphere. (2020) doi:10.1097/HS9.00000000000398.
- **391.** Martelli AM, Evangelisti C, Chiarini F, McCubrey JA. The phosphatidylinositol 3-kinase/Akt/mTOR signaling network as a therapeutic target in acute myelogenous leukemia patients. Oncotarget, 89-103, 2010.
- **392.** Barrett D, Brown VI, Grupp SA, Teachey DT. Targeting the PI3K/AKT/mTOR signaling axis in children with hematologic malignancies. Paediatr Drugs. 299-316, 2012.
- **393.** Renner C, Zinzani P L, Gressin R et al: Swiss SAKK and French GOELAMS group from European Mantle Cell Lymphoma Network (2012). A multicenter phase II trial (SAKK 36/06) of singleagent everolimus (RAD001) in patients with relapsed or refractory mantle cell lymphoma. Haematologica

/doi.org/10.3324/haematol.2011.053173

- **394.** Ghosh J, Kapur R: Role of mTORC1–S6K1 signaling pathway in regulation of hematopoietic stem cell and acute myeloid leucemia. Experimental Hematology 13–21, 2017.
- **395.** Tabe Y, Tafuri A, Sekihara K et al. Inhibition of mTOR kinase as a therapeutic target for acute myeloid leukemia. Expert Opin Ther Targets. 705714, 2017.
- **396.** Feng Y, Chen X, Cassady K et al. The role of mtor inhibitors in hematologic disease: from bench to bedside. Front Oncol. (2021) doi:10.3389/fonc.2020.611690.
- 397. Mekki M, Bridson JM, Sharma A, Halawa A. mTOR

inhibitors in kidney transplantation: A comprehensive review J Kidney, (2017) doi: 10.4172/24721220.1000146.

- **398.** Franz DN, Leonard J, Tudor C et al. Rapamycin causes regression of astrocytomas in tuberous sclerosis complex. Ann Neurol. 490–498, 2006.
- **399.** Li XY, Zhang LQ, Zhang XG et al. Association between AKT/mTOR signalling pathway and malignancy grade of human gliomas. J Neurooncol. 453458, 2011.
- **400.** Duzgun Z, Eroglu Z, Biray C. Role of mTOR in glioblastoma. Gene. 187190, 2016.
- **401.** Ryskalin L, Lazzeri G, Flaibani M et al: mTOR-Dependent cell proliferation in the brain. Biomed Res Int. (2017). doi: 10.1155/2017/7082696.
- **402.** Waetzig R, Matthes M, Leister J et al. Comparing mTOR inhibitor rapamycin with Torin-2 within the RIST molecular-targeted regimen in neuroblastoma cells. Int J Med Sci. 137-149, 2021.
- 403. Lenzi P, Ferese R, Biagioni F et al: Rapamycin ameliorates defects in mitochondrial fission and mitophagy in glioblastoma cells. Int J Mol Sci. (2021) doi: 10.3390/ijms22105379.
- **404.** Mukhopadhyay S, Frias MA, Chatterjee A et al: the enigma of rapamycin dosage. Mol Cancer Ther 347–353, 2016.
- 405. Xie J, Wang X, Proud CG. mTOR inhibitors in cancertherapy. (2016). doi:10.12688/f1000research.9207.1
- **406.** Alzahrani AS. PI3K/Akt/mTOR inhibitors in cancer: At the bench and bedside. Semin Cancer Biol. 125-132, 2019.
- **407.** Hua H, Kong Q, Zhang H et al. Targeting mTOR for cancer therapy. J Hematol Oncol (2019). doi.org/10.1186/s13045-019-0754-1.
- **408.** Harsha C, Banik K, Ang HL et al: Targeting AKT/mTOR in oral cancer: mechanisms and advances in clinical trials. Int J Mol Sci. (2020) doi: 10.3390/ijms21093285
- **409.** Wang H, Liu Y, Ding J et al. Targeting mTOR suppressed colon cancer growth through 4EBP1/eIF4E/PUMA pathway. Cancer Gene The, 448–460, 2020.
- **410.** Bockaert J, Marin P. mTOR in brain physiology and pathologies. Physiol Rev. 1157-1187, 2015.
- **411.** Norwitz NG, Querfurth H. mTOR Mysteries: Nuances and questions about the mechanistic target of rapamycin in neurodegeneration. Front Neurosci. (2020). doi: 10.3389/fnins.2020.00775.
- **412.** Heras-Sandoval D, Pérez-Rojas JM, Pedraza-Chaverri J. Novel compounds for the modulation of mTOR and autophagy to treat

neurodegenerative diseases. Cell Signal. (2020) doi: 10.1016/j.cellsig.2019.109442.

- **413.** Querfurth H, Lee HK. Mammalian/mechanistic target of rapamycin (mTOR) complexes in neurodegeneration. Mol Neurodegener. (2021) doi: 10.1186/s13024-021-00428-5.
- **414.** Ferri CP, Prince M, Brayne C et al: Alzheimer's disease international global prevalence of dementia: a Delphi consensus study. Lancet, 2112-2117, 2005.
- **415.** Pei JJ, Hugon J. mTOR-dependent signalling in Alzheimer's disease. J Cell Mol Med. 2525-2532, 2008.
- **416.** Oddo S. The role of mTOR signaling in Alzheimer disease. Front Biosci , 941-952, 2012.
- **417.** Caccamo A, Magrì A, Medina DX et al: mTOR regulates tau phosphorylation and degradation: implications for Alzheimer's disease and other tauopathies. Aging Cell. 370-380, 2013.
- **418.** Tramutola A, Triplett JC, Di Domenico F, Alteration of mTOR signaling occurs early in the progression of Alzheimer disease (AD): analysis of brain from subjects with pre-clinical AD, amnestic mild cognitive impairment and late-stage AD.J Neurochem. 739-49, 2015.
- **419.** Perluigi M, Di Domenico F, Butterfield DA. mTOR signaling in aging and neurodegeneration: At the crossroad between metabolism dysfunction and impairment of autophagy. Neurobiol Dis. 39-44, 2015.
- **420.** Perluigi M, Di Domenico F, Barone E, Butterfield DA. mTOR in Alzheimer disease and its earlier stages: Links to oxidative damage in the progression of this dementing disorder. Free Radic Biol Med. 169:382-396, 2021.
- **421.** Iyer AM, van Scheppingen J, Milenkovic I et al: mTOR Hyperactivation in Down syndrome hippocampus appears early during development. J Neuropathol Exp Neurol. 671-683, 2014.
- **422.** Perluigi M, Pupo G, Tramutola A et al: Neuropathological role of PI3K/Akt/mTOR axis in Down syndrome brain. Biochim Biophys Acta. 11441153, 2014.
- **423.** Di Domenico F, Tramutola A, Foppoli C et al: mTOR in Down syndrome: Role in Aß and tau neuropathology and transition to Alzheimer disease-like dementia. Free Radic Biol Med. 94-101, 2018.
- **424.** Bordi M, Darji S, Sato Y et al. mTOR hyperactivation in Down syndrome underlies deficits in autophagy induction, autophagosome formation, and mitophagy. Cell Death Dis (2019). doi.org/10.1038/s41419-019-1752-5

- **425.** Martínez-Cué C, Rueda N. Signalling pathways implicated in Alzheimer's disease neurodegeneration in individuals with and without Down syndrome. Int J Mol Sci. (2020). doi: 10.3390/ijms21186906.
- **426.** Troca-Marín JA, Casañas JJ, Benito I, Montesinos ML. The Akt-mTOR pathway in Down's syndrome: the potential use of rapamycin/rapalogs for treating cognitive deficits. CNS Neurol Disord Drug Targets. 34-40, 2014.
- **427.** Urbano-Gámez JD, Casañas JJ, Benito I, Montesinos ML. Prenatal treatment with rapamycin restores enhanced hippocampal mGluR-LTD and mushroom spine size in a Down's syndrome mouse model. Mol Brain. (2021). doi: 10.1186/s13041-021-00795-6.
- **428.** Di Domenico F, Tramutola A, Barone E et al: Restoration of aberrant mTOR signaling by intranasal rapamycin reduces oxidative damage: Focus on HNE-modified proteins in a mouse model of Down syndrome. Redox Biol. (2019) doi: 10.1016/j.redox.2019.101162.
- **429.** Tramutola, A., Lanzillotta, C., Barone, E. et al. Intranasal rapamycin ameliorates Alzheimer-like cognitive decline in a mouse model of Down syndrome. Transl Neurodegener 7, 28 (2018). https://doi.org/10.1186/s40035018-0133-9.
- **430.** Maiese K, Chong ZZ, Shang YC, Wang S. mTOR: On target for novel therapeutic strategies in the nervous system. Trends Mol. Med. 51–60, 2013.
- **431.** Nagahama Y, Shimoda M, Mao G et al: Regnase-1 controls colon epithelial regeneration via regulation of mTOR and purine metabolism. Proc Natl Acad Sci USA. 11036-11041, 2018.
- 432. Zhou JY, Huang DG, Qin YC et al: mTORC1 signaling activation increases intestinal stem cell activity and promotes epithelial cell proliferation. J. Cell. Physiol. 19028–19038, 2019.
- **433.** Wei X, Luo L, Chen J. Roles of mTOR signaling in tissue regeneration. Cells (2019). doi.org/10.3390/cells8091075.
- **434.** Lund-Ricard Y, Cormier P, Morales J, Boutet A. mTOR signaling at the crossroad between metazoan regeneration and human diseases. Int J Mol Sci. (2020). doi: 10.3390/ijms21082718.
- **435.** Qian J, Su S, Liu P. Experimental approaches in delineating mTOR signaling. Genes (2020). doi: 10.3390/genes11070738.
- **436.** Maiese K. Targeting the core of neurodegeneration: FoxO, mTOR and SIRT1. Neural Regen Res. 448-455, 2021.
- **437.** Lu F, Leach LL, Gross JM. mTOR activity is essential for retinal pigment epithelium

regeneration in zebrafish. PLoS Genet. (2022). doi: 10.1371/journal.pgen.1009628.

- **438.** Hilton, BJ, Bradke, F. Can injured adult CNS axons regenerate by recapitulating development? Development, 3417–3429, 2017.
- **439.** Schelski M, Bradke F. Neuronal polarization: From spatiotemporal signaling to cytoskeletal dynamics. Mol. Cell. Neurosci. 11–28, 2017.
- **440.** Wang F, Chen F, Wang G et al. Rapamycin provides anti-epileptogenic effect in a rat model of post-traumatic epilepsy via deactivation of mTOR signaling pathway. Exp Ther Med. 2018;15(6):4763-4770. doi:10.3892/etm.2018.6004

441. Petrova V, Nieuwenhuis B, Fawcett JW, Eva R. Axonal organelles as molecular platforms for axon growth and regeneration after Injury. Int 1 Mol

- growth and regeneration after Injury. Int J Mol Sci. (2021) doi.org/10.3390/ijms22041798. **442.** Leibinger M, Andreadaki A, Fischer D. Role of
- 442. Leibinger M, Andreadaki A, Fischer D. Role of mTOR in neuroprotection and axon regeneration after inflammatory stimulation. Neurobiol Dis. 314–324, 2012.
- **443.** Poulopoulos A, Murphy AJ, Ozkan A. et al: Subcellular transcriptomes and proteomes of developing axon projections in the cerebral cortex. Nature. 356– 360, 2019.
- **444.** Chen CH, Sung CS, Huang SY et al: The role of the PI3K/Akt/mTOR pathway in glial scar formation following spinal cord injury. Exp. Neurol. 27–41, 2016.
- **445.** Yang P, Wen H, Ou S et al: IL-6 promotes regeneration and functional recovery after cortical spinal tract injury by reactivating intrinsic growth program of neurons and enhancing synapse formation. Exp. Neurol. 19–27, 2012.
- **446.** Sivanand S, Heiden MGV: Emerging roles for branched-chain amino acid metabolism in câncer. Cancer Cell, 147-156, 2020.
- 447. Villa-González M, Martín-López G, Pérez-Álvarez MJ. Dysregulation of mTOR signaling after brain ischemia. Int J Mol Sci. (2022). doi: 10.3390/ijms23052814.

