Effects of anti-cancer targeted therapies on lipid and glucose metabolism

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Abstract

During the past years, targeted therapies for cancer have been developed using drugs that have significant metabolic consequences. Among them, the mammalian target of rapamycin (mTOR) inhibitors and, to a much lesser extent, the tyrosine kinase inhibitors (TKIs) are involved. mTOR plays a key role in the regulation of cell growth as well as lipid and glucose metabolism. Treatment with mTOR inhibitors is associated with a significant increase in plasma triglycerides and LDL cholesterol. mTOR inhibitors seem to increase plasma triglycerides by reducing the activity of the lipoprotein lipase which is in charge of the catabolism of triglyceride-rich lipoproteins. The increase in LDL cholesterol observed with mTOR inhibitors seems to be due to a decrease in LDL catabolism secondary to a reduction of LDL receptor expression. In addition, treatment with mTOR inhibitors is associated with a high incidence of hyperglycemia, ranging from 13 to 50% in the clinical trials. The mechanisms responsible for hyperglycemia with new onset diabetes are not clear, but are likely due to the combination of impaired insulin secretion and insulin resistance. TKIs do not induce hyperlipidemia but alter glucose homeostasis. Treatment with TKIs may be associated either with hyperglycemia or hypoglycemia. The molecular mechanism by which TKIs control glucose homeostasis remains unknown. Owing to the metabolic consequences of these agents used as targeted anti-cancer therapies, a specific and personalized follow-up of blood glucose and lipids is recommended when using mTOR inhibitors and of blood glucose when using TKIs.

Introduction

Targeted therapies refer to the use of medication that blocks the growth of cancer cells by interfering with specific targeted molecules needed for carcinogenesis and tumor growth. During recent years, targeted therapies have been more frequently used to treat cancers. Some of the drugs used for targeted therapies have significant...
metabolic consequences, more particularly the mammalian target of rapamycin (mTOR) inhibitors and to a much lesser extent, the tyrosine kinase inhibitors (TKIs). Indeed, mTOR inhibitors, by interfering with the protein kinase mTOR pathway, which is also involved in glucose and lipid metabolism, are responsible for frequent dyslipidemia and hyperglycemia. Most of the TKIs have no significant effects on lipids and glucose metabolism. However, increasing evidence suggests that some TKIs, such as imatinib, sunitinib, pazopanib, or nilotinib, induce adverse reactions of hypoglycemia or hyperglycemia. We will consider in this review drugs currently approved by the Food and Drug Administration (FDA) in oncology with available phase III trial results but also data on mTOR inhibitors when used at the lowest doses as anti-rejection therapy for transplantation. Moreover, mTOR inhibitors also modify phospho-calcium metabolism. However, we will only consider their effects on lipid and glucose metabolism.

**Main indications for mTOR inhibitors and TKIs**

mTOR inhibitors were first used as anti-rejection agents for transplantation before being developed as targeted anti-cancer therapies (1). Both sirolimus (i.e. rapamycin) and everolimus are used as anti-rejection treatment while everolimus and temsirolimus are used as targeted anti-cancer therapies. Everolimus is used for both indications but with lower doses for anti-rejection (1.5–3.0 mg/day) than for anti-cancer (5–10 mg/day) treatment.

mTOR inhibitors are agents that inhibit signaling in the phosphoinositide 3-kinase (PI3K)–Akt–mTOR pathway, playing a key role in the regulation of cell growth. Although several compounds affecting this pathway are in various stages of clinical development, only two mTOR inhibitors (temsirolimus and everolimus) have been approved in oncology for marketing in the USA by the FDA. Everolimus is approved by the FDA for advanced progressive pancreatic neuroendocrine tumors (P-NETs) (2), advanced renal cell carcinoma (RCC) after failure of treatment with sunitinib or sorafenib (3), advanced hormone receptor-positive breast cancer in combination with exemestane (4), and for subependymal giant cell astrocytoma associated with tuberous sclerosis (5). Temsirolimus is approved by FDA in RCC (6) and mantle cell lymphoma (7).

TKIs are small molecules inhibiting tyrosine kinases that are enzymes playing an important role in the modulation of growth factor signaling. Imatinib is approved for treatment of chronic myeloid leukemia (CML), Philadelphia-positive (Ph+) acute lymphoblastic leukemia, chronic eosinophilic leukemia, and gastrointestinal stromal tumor (GIST) (8). Sunitinib is approved in second line for GIST (9), RCC (10), and recently for advanced P-NETs (11). Pazopanib is approved for advanced soft tissue sarcoma and RCC (12, 13). Nilotinib is approved for Ph+ CML (14).

**mTOR inhibitors**

**Influence of mTOR on lipid and carbohydrate metabolism**

mTOR is a serine/threonine kinase, belonging to the PI3K-related kinase family, playing a key role in the regulation of cell growth as well as lipid and glucose metabolism. mTOR nucleates at least two distinct multi-protein complexes, mTOR complex 1 (mTORC1) and mTORC2 (15, 16). mTORC1 is composed of five subunits and regulates processes involved in the promotion of cell growth and metabolism. mTORC1 responds to amino acids (arginine and branched chain amino acids: leucine, isoleucine, and valine), stress, energy status, oxygen, and growth factors (including insulin). mTORC2, which is composed of six subunits, is activated by growth factors and regulates cell survival, cell metabolism, and cytoskeletal organization (Fig. 1).

Once activated, mTORC1 regulates many cellular processes. mTORC1 promotes protein synthesis by directly phosphorylating the translational regulators eukaryotic translation initiation factor 4E (eIF4E)-binding protein (4E-BP1) and S6 kinase 1 (S6K1) (17). In addition, mTORC1 controls growth by inhibiting autophagy (18), the central degradative process in cells. Finally, mTOR positively regulates cellular metabolism and ATP production by inducing the expression of many glycolytic genes, through the activation of the transcription factor hypoxia inducible factor 1α (19).

Compared with mTORC1, the functions of the mTORC2 pathway are poorly known. mTORC2 controls several protein kinases, including Akt/PKB, serum- and glucocorticoid-induced protein kinase 1 (SGK1), and protein kinase C (PKC)-α (20, 21). By controlling Akt and SGK1, mTORC2 impacts cell growth/survival and metabolism, while it controls actin cytoskeleton dynamics by activating PKC-α (22).

The paradigm states that mTORC1, but not mTORC2, is sensitive to inhibition by rapamycin, although its exact mechanism of action remains unclear. The picture is less clear, however, because long-term treatment with rapamycin can also reduce mTORC2 signaling by suppressing mTORC2 assembly (23).
mTOR and lipid metabolism

mTORC1 and mTORC2 complexes. mTORC1 is composed of six protein components and responds to various stimuli, such as amino acids, stress, growth factors, energy, and oxygen. mTORC1 is acutely inhibited by rapamycin and promotes cell growth by inducing anabolic processes and cell-cycle progression and inhibiting autophagy. mTORC2 is composed of seven protein components and is involved in cytoskeletal organization and cell survival. Both mTORC1 and mTORC2 are involved in metabolic homeostasis.

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mTOR and lipid metabolism

mTORC1 controls lipid metabolism by different ways. Several data indicate that mTORC1 increases the activity of the transcription factor sterol regulatory element-binding protein-1c (SREBP-1c), which upregulates the expression of enzymes involved in lipogenesis (acetyl-CoA carboxylase, fatty acid synthase, and stearoyl-CoA desaturase). It has been shown that impairment of mTORC1 activity with rapamycin blocks the SREBP-1c induced expression of lipogenic enzymes (24). This is in line with data showing that rapamycin reduces the expression of SREBP-1c target genes such as acetyl-CoA carboxylase (25), fatty acid synthase (26), and stearoyl-CoA desaturase (27). Increased de novo lipid synthesis is an important feature of cancer cells and it is suggested that induction of lipid metabolism by SREBP-1c promotes cancer progression by providing lipids required for membranes. mTORC1 has been shown to increase PPARγ expression and then to promote adipogenesis (28). It has also been suggested that mTORC1 increases the activity of lipin 1, which is a phosphatidic acid phosphatase promoting triglyceride synthesis and enhancing the PPARγ adipogenic activity (29). Lipin 1 has also been suggested to upregulate SREBP-1c expression and it has been shown in mice that inhibition of mTORC1 in the liver significantly impairs SREBP function in a lipin 1-dependent fashion (30).

So far, a possible implication of mTORC2 in the control of lipogenesis remains imprecise. Some data suggest that mTORC2 may control sphingolipid de novo synthesis by regulating the activity of the ceramide synthase (31), but further studies are needed to clarify the potential role of mTORC2 in the control of lipid metabolism.

mTOR and carbohydrate metabolism

The effects of mTOR on glucose homeostasis are complex with some opposite results depending on the level of mTORC1 activity (16). mTORC1 promotes insulin resistance in adipose tissue through the S6K1-mediated inhibition of insulin signaling, linked to the serine-phosphorylation of insulin receptor substrate 1 (IRS1) that disrupts the recruitment and activation of PI3K (32). The activity of mTORC1 is increased in insulin-target tissues (adipose tissue, skeletal muscle, and liver) of insulin-resistant rodents, reinforcing the hypothesis of a deleterious effect of sustained mTORs signaling on insulin sensitivity (32, 33, 34). In the liver, the serine phosphorylation of IRS1 by mTORC1/S6K1 also leads to the impairment of the PI3K–Akt metabolic
pathway, thereby promoting gluconeogenesis. Conversely, mTORC1 also plays a key role in the oxidative metabolism of skeletal muscle, as muscle-specific mTORC1 knockout mice display a muscle dystrophy with reduced muscle mass and oxidative function (35). In the liver, mTORC1 activity, which is reduced during fasting, inhibits ketogenesis by blunting the activity of PPARα, the master transcriptional modulator of ketogenic genes (36).

mTORC1 also plays a central role in the regulation of the β-cell mass and function, which are some critical components in the pathogenesis of type 2 diabetes mellitus (T2DM). mTORC1 is a positive regulator of β-cell function, as constitutive activation of mTORC1 in β-cells improves insulin secretion and decreases blood glucose levels, and is linked to an increase in both β-cell size and number (37). Conversely, S6K1-deficient mice exhibit diminished β-cell size, hypoinsulinemia, and glucose intolerance (38). However, a sustained activation of mTORC1 in β-cells ultimately leads to an exhaustion of their insulin secreting capacity and to a deterioration of glucose homeostasis, by promoting apoptosis and reducing cell survival (39).

**Effects of mTOR inhibitors on lipids**

The use of mTOR inhibitors is known to induce dyslipidemia. Indeed, a review of 17 randomized controlled trials shows an increase in total cholesterol and/or triglycerides in all but three trials (40). Among those three trials reporting no difference in lipids between groups, the percentage of mTOR inhibitor-treated patients on lipid lowering therapy were increased in two of them, indicating that mTOR inhibitors might have induced dyslipidemia (41, 42). In a study performed on 474 kidney transplant recipients, it was noted that dyslipidemia is frequent 1 year after transplantation and, in multivariate analysis, that treatment by mTOR inhibitors (sirolimus or everolimus) is an independent factor associated with dyslipidemia (43). As discussed later, dyslipidemia is also observed when mTOR inhibitors are used as targeted anti-cancer therapy.

**Sirolimus (rapamycin)**

A significant increase in total cholesterol ranging from 25 to 55% has been observed with sirolimus (40, 41, 43, 44, 45, 46, 47, 48). In most clinical studies, sirolimus treatment is associated with a significant increase in LDL cholesterol (40, 43, 47). In the study performed by Spinelli et al. (43), sirolimus treatment was an independent factor associated with plasma LDL cholesterol levels above 121 mg/dl. It has also been reported that treatment with sirolimus is associated with a dramatic increase in plasma triglycerides ranging from 65 to 100% (41, 42, 43, 44, 45, 47). Patients receiving sirolimus 2 or 5 mg/day have been shown to have plasma triglycerides respectively 59 or 103 mg/dl greater than that of controls (41). In a multivariate analysis, sirolimus treatment was an independent factor associated with hypertriglyceridemia (43).

Plasma HDL cholesterol has been shown to be unchanged or slightly increased in clinical trials (43, 44, 45, 46, 47). Thus, the detrimental effect of sirolimus on lipids is obvious and also confirmed by the fact that sirolimus-treated patients more frequently require adjunctive treatment with lipid lowering drugs (49). The hyperlipidemic effect of sirolimus is observed in the absence (50) and in the presence of ciclosporin (48, 49). In addition, data from the clinical studies indicate that the hyperlipidemic effect of sirolimus is dose-dependent (41, 49).

**Everolimus**

Increase in total cholesterol with everolimus has been reported in several studies (3, 43, 44, 50). In clinical trials which evaluated the efficacy of everolimus as anti-cancer therapy, hypercholesterolemia has been reported to be significantly more frequent with everolimus vs placebo (3, 50) (Table 1). When measured, LDL cholesterol has been found to be increased with everolimus treatment (43).

Treatment with everolimus is associated with a significant increase in plasma triglycerides in most studies (40, 43, 44, 51, 52). In the phase III trial evaluating everolimus in metastatic RCC, hypertriglyceridemia was significantly more frequent with everolimus (71%) than with placebo (30%) (3). In post-transplantation studies, an increase in plasma HDL cholesterol level from 15 to 21% has been reported (43, 44, 45).

Several studies have pointed out some differences between everolimus and sirolimus as far as their lipid effects are concerned. A lower increase in plasma triglycerides with everolimus compared with sirolimus has been reported in two studies (44, 45), but not in the study performed by Spinelli et al. (43). Total cholesterol has been shown to be increased with sirolimus but not with everolimus (45). However, this different effect on cholesterol levels between the two drugs has not been found in other studies (43, 44). We have to say that the data comparing the lipid effects of the two drugs are limited and sometimes discrepant and do not allow drawing firm conclusions.

**Temsirolimus**

Temsirolimus has also been reported to induce hyperlipidemia. In a phase III trial comparing
temsirolimus and interferon for treatment of advanced RCC, hypercholesterolemia was significantly more frequent with temsirolimus (24%) than that with interferon (4%) (6). Treatment of advanced breast cancer with temsirolimus in association with letrozole is responsible for twice more hypercholesterolemia (12%) than treatment with letrozole alone (6%) (53) (Table 1).

Pathophysiology of mTOR inhibitor-induced dyslipidemia. The pathophysiology of mTOR inhibitor-induced dyslipidemia is not yet totally clear. In vitro studies have shown that mTOR inhibition by rapamycin increases fatty acids β-oxidation in the hepatocytes (25, 54) and skeletal muscles (55). In addition, several studies have shown that rapamycin reduces the gene expression of lipogenic enzymes such as acetyl-CoA carboxylase (25), fatty acid synthase (26), and stearoyl-CoA desaturase (27). In addition, rapamycin has been shown to promote fatty acid β-oxidation while decreasing fatty acid flux into anabolic storage pathways in primary cultures of rat hepatocytes, suggesting that the hyperlipidemia associated with rapamycin is not likely due to increased lipid hepatic synthesis but rather due to delayed peripheral clearance (25). Indeed, in kidney transplant recipients with sirolimus-induced hypertriglyceridemia, it has been shown with an in vivo kinetic study that hypertriglyceridemia was mostly due to decreased very-LDL (VLDL) catabolism (56). In that study, treatment with sirolimus was associated with a 70% decrease in VLDL-apoB100 catabolism and a 28% decrease in IDL-apoB100 catabolism (56). In line with these kinetic results, a significant decrease in post-heparin lipoprotein lipase (LPL) activity (the major enzyme responsible for triglyceride catabolism) was observed in patients showing sirolimus-induced hypertriglyceridemia (56) (Fig. 2). Interestingly, it has been shown that insulin upregulates LPL activity in a PI3K-dependent manner and that this regulation is decreased by 60% in the presence of rapamycin (57). In addition, treatment with rapamycin (sirolimus) is associated with a 92% increase in apolipoprotein CIII, which is a potent inhibitor of LPL (47). Thus, all these data indicate that mTOR inhibitors are likely to decrease LPL activity and

Table 1  Incidence of hyperlipidemia and hyperglycemia induced by mTOR inhibitors in phase III studies.

<table>
<thead>
<tr>
<th>Drug</th>
<th>Types of cancer</th>
<th>Hyperglycemia (%)</th>
<th>Hypercholesterolemia (%)</th>
<th>Hypertriglyceridemia (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>All grades</td>
<td>Grades 3–4</td>
</tr>
<tr>
<td>Everolimus</td>
<td>Renal cell carcinoma</td>
<td>269</td>
<td>50</td>
<td>12</td>
</tr>
<tr>
<td>Placebo (3)</td>
<td></td>
<td>135</td>
<td>23</td>
<td>1</td>
</tr>
<tr>
<td>Everolimus</td>
<td>Pancreatic NETs</td>
<td>204</td>
<td>13</td>
<td>5</td>
</tr>
<tr>
<td>Placebo (2)</td>
<td></td>
<td>203</td>
<td>4</td>
<td>2</td>
</tr>
<tr>
<td>Everolimus + Oct</td>
<td>Gastrointestinal NETs</td>
<td>215</td>
<td>12</td>
<td>5</td>
</tr>
<tr>
<td>Placebo + Oct (64)</td>
<td></td>
<td>211</td>
<td>2</td>
<td>0.5</td>
</tr>
<tr>
<td>Everolimus</td>
<td>Breast</td>
<td>482</td>
<td>13</td>
<td>4</td>
</tr>
<tr>
<td>Placebo (4)</td>
<td></td>
<td>238</td>
<td>2</td>
<td>&lt;1</td>
</tr>
<tr>
<td>Everolimus</td>
<td>Subependymal giant cell astrocytoma with TBS</td>
<td>78</td>
<td>ND</td>
<td>ND</td>
</tr>
<tr>
<td>Placebo (5)</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Temsirolimus</td>
<td>Renal cell carcinoma</td>
<td>208</td>
<td>26</td>
<td>11</td>
</tr>
<tr>
<td>Interferon</td>
<td></td>
<td>200</td>
<td>11</td>
<td>2</td>
</tr>
<tr>
<td>Tensi + interferon (6)</td>
<td></td>
<td>208</td>
<td>17</td>
<td>6</td>
</tr>
<tr>
<td>Temsirolimus + letrozole</td>
<td></td>
<td>550</td>
<td>13</td>
<td>4</td>
</tr>
<tr>
<td>Placebo + letrozole (53)</td>
<td></td>
<td>553</td>
<td>5</td>
<td>1</td>
</tr>
<tr>
<td>Temsirolimus</td>
<td>Mantle cell lymphoma</td>
<td>108</td>
<td>ND</td>
<td>ND</td>
</tr>
<tr>
<td>Investigator’s choice (7)</td>
<td></td>
<td>53</td>
<td>ND</td>
<td>ND</td>
</tr>
</tbody>
</table>

NETs, neuroendocrine tumors; ND, not described (incidence are <10%); Oct, octreotide; TBS, tuberous sclerosis complex. Common terminology criteria for adverse events (CTCAE) grades for hyperglycemia: Grade 1, fasting blood glucose between 126 and 160 mg/dl (7.0–8.9 mmol/l); Grade 2, fasting blood glucose between 161 and 250 mg/dl (8.9–13.9 mmol/l); Grade 3, blood glucose between 251 and 500 mg/dl (13.9–27.8 mmol/l); Grade 4, blood glucose > 500 mg/dl (27.8 mmol/l). CTAE grades for hypercholesterolemia: Grade 1, total cholesterol between the upper limit of the normal range and 300 mg/dl (7.75 mmol/l); Grade 2, total cholesterol between 301 and 400 mg/dl (7.76–10.34 mmol/l); Grade 3, total cholesterol between 401 and 500 mg/dl (10.35–12.92 mmol/l); Grade 4, total cholesterol > 500 mg/dl (12.92 mmol/l). CTAE grades for hypertriglyceridemia: Grade 1, triglycerides between 150 and 300 mg/dl (1.71–3.42 mmol/l); Grade 2, triglycerides between 301 and 500 mg/dl (3.43–5.70 mmol/l); Grade 3, triglycerides between 501 and 1000 mg/dl (5.71–11.4 mmol/l); Grade 4, triglycerides > 1000 mg/dl (11.4 mmol/l).
thus to reduce the catabolism of triglyceride-rich lipoproteins leading to hypertriglyceridemia. It has also been shown that in renal transplant patients, treatment with sirolimus induced a 42% increase in free fatty acid pool, suggesting an augmented release of free fatty acids from the adipose tissue that may be due to a reduction of the inhibition of the hormone-sensitive lipase by insulin (47). It has been postulated that this increase in free fatty acids may increase hepatic VLDL production, but such increase in VLDL production has not been clearly shown in human kinetic studies (56).

As far as LDL cholesterol is concerned, a kinetic study performed in renal transplant patients has shown that treatment with sirolimus was associated with a 68% decrease in LDL-apoB100 catabolism (56). It has been shown in vitro with HepG2 cells that sirolimus reduced LDL receptor (LDLr) expression leading to diminished LDL uptake (58) (Fig. 2). This inhibitory effect is mediated by the downregulation of SREBP cleavage activating protein (SCAP) and SREBP-2 mRNA and by reduced translocation of SCAP–SREBP-2 complex from endoplasmic reticulum to Golgi for activation, thereby reducing LDLr gene transcription (58). Proprotein convertase subtilisin kexin type 9 (PCSK9) is an endogenous inhibitor of the LDLr (for review see (59)). Interestingly, a recent report has demonstrated that rapamycin increased expression of PCSK9, decreased levels of hepatic LDLr protein, and finally increased levels of VLDL/LDL cholesterol in WT but not in PCSK9 knockout mice (60). These results suggest that PCSK9 could be a molecular link for the mTOR inhibitor-induced dyslipidemia.

So far, the mechanisms responsible for increased plasma HDL cholesterol observed with everolimus and sometimes with sirolimus are not known. In one study, an inconsistent increase in plasma apoA-I, the major apolipoprotein of HDL, was observed but it did not totally parallel the increase in HDL cholesterol (47). To date, no clear hypotheses can be made to explain the increase of HDL cholesterol with mTOR inhibitors and studies are needed to clarify this point.

**Figure 2**
Pathophysiology of the dyslipidemia induced by mTOR inhibitors. VLDL, very-LDL; IDL, intermediate density lipoprotein; LPL, lipoprotein lipase; B/E rec., receptor B/E (LDL receptor); ABCA1, ATP-binding cassette transporter A1; ABCG1, ATP-binding cassette transporter G1; TG, triglycerides.
Effects of mTOR inhibitors on glycemia ► The perturbation in glucose homeostasis with mTOR inhibitors is the second main metabolic complication for this class of drugs.

Sirolimus (rapamycin) ► The first data came from the use of sirolimus as immunosuppressive agent in patients with kidney transplant, demonstrating that sirolimus can induce new-onset diabetes in this population, especially when the drug is used in association with calcineurin inhibitors (61). For instance, the 3-year cumulative incidence of new onset-diabetes was 21.9% in patients treated by the combination of sirolimus and ciclosporin A (n=800) and 21.5% in those with sirolimus and tacrolimus (n=1179) (62). In this study, treatment with sirolimus was independently associated with a risk of new-onset diabetes (62). A more direct proof of the direct deleterious effect of sirolimus on glucose homeostasis comes from a small pilot study assessing the metabolic consequences of the withdrawal of calcineurin inhibitors (ciclosporin A or tacrolimus) and the switch to sirolimus in 41 adult kidney transplant recipients. The switch to sirolimus led to a 30% increase in the incidence of impaired glucose tolerance and to new-onset diabetes mellitus in four patients (63).

Everolimus and temsirolimus ► A same tendency for a deterioration of glucose homeostasis is observed when mTOR inhibitors are used as anti-cancer agents. The frequency of hyperglycemia was assessed in randomized trials as an adverse event listed in the Common Terminology Criteria for Adverse Events (see Table 1). Hyperglycemia was reported in 12–50% of patients in phase III studies for everolimus (2, 3, 4, 5, 64) and temsirolimus, but grades 3–4 (G3–4) hyperglycemic events (>13.9 mmol/l) were less frequent (4–22%) (Table 1).

However, all these studies did not report the percentage of patients treated with glucose-lowering agents and their dose. Moreover, these studies did not reflect the real life of patients treated by mTOR inhibitors, as patients with uncontrolled diabetes were excluded in these trials. One can suspect that that the incidence of hyperglycemia might be different according to the treated cancer with, for instance, a higher proportion of diabetes in P-NETs, which in theory are more likely to induce diabetes. However, this is not really the case as, in clinical trials, the highest frequency of diabetes has been found in patients with advanced RCC (3, 6). For instance, in the RADIANT-3 study of treatment with everolimus in patients with advanced P-NETs, 13% of treated patients exhibited hyperglycemia (with 5% G3–4), compared with 4 and 2% in the placebo group respectively (2). In comparison, hyperglycemia appears to be more frequent in patients with advanced RCC, as 50% of everolimus-treated patients exhibited hyperglycemic events in the RECORD-1 trial compared with 23% in the placebo group (3). This indicates that hyperglycemia induced by mTOR inhibitors is observed independently of the origin of treated cancer.

Pathophysiology of hyperglycemia induced by mTOR inhibitors ► The role of mTOR pathway and its manipulation with mTOR inhibitors is undoubtedly complex. Notably, there are some still unexplained discrepancies between both in vitro and in vivo preclinical studies on rodents and clinical data from phase II/III study with everolimus on humans.

It seems that mTOR inhibitors exert a ‘Janus effect’ on glucose metabolism. Indeed, based on the observation that chronic activation of mTORC1 promotes insulin resistance, it could be hypothesized that the chronic inhibition of the mTOR pathway could improve glucose homeostasis. As suggested by Laplante & Sabatini (16), the relation between mTORC1 activity and metabolic homeostasis probably follows a U-shaped curve, where too little or too much mTORC1 activity has a deleterious effect on systemic metabolism (Fig. 3). In addition, it has been demonstrated very recently that the duration of mTOR inhibition with rapamycin is of critical importance for regulating metabolic homeostasis (65). While short-term treatment with rapamycin exerts detrimental effects on glucose homeostasis in mice, prolonged treatment (i.e. 20 weeks) is associated with enhanced insulin sensitivity, increased energy expenditure, and better lipid profile (65). Notably, the antidiabetic drug metformin might improve glucose homeostasis by partially inhibiting mTORC1 signaling (16, 65). Thus, the metabolic effects of treatment with rapamycin have been assessed in several rodent models. Unexpectedly, rapamycin leads to the aggravation of glucose homeostasis with the development of T2DM, linked to a combination of an impaired insulin secretion and severe insulin resistance (67, 68). Moreover, it has been shown recently that chronic treatment with rapamycin disrupts mTORC2 complex and blocks its inhibiting activity on hepatic gluconeogenesis (69). Similar results were observed with a selective mTOR kinase inhibitor AZD8055, with an increase in blood glucose levels in treated mice (70). In humans, long-term treatment with rapamycin impaired insulin signaling assessed by the phosphorylation of Akt in circulating mononuclear cells (PBMC) of kidney-
transplanted patients (71). In accordance with an impairment of insulin signaling, insulin-stimulated glucose uptake was reduced in human adipocytes treated for a short-term (15 min) or long-term treatment (20 h) with rapamycin at therapeutic concentrations (72). At the whole body level, the switch of calcineurin inhibitors to sirolimus induces a worsening of insulin resistance associated with a deterioration of compensatory insulin secretion, leading to new-onset diabetes (63). Interestingly, Kulke et al. (72) first reported in 2009 that the use of everolimus in four patients with insulinoma and refractory hypoglycemia was associated with improved glycemic control. This ‘beneficial’ hyperglycemic effect of mTOR inhibitors in patients with severe insulinoma is another proof of the hyperglycemic effect of this therapeutic class.

Tyrosine kinase inhibitors

TKIs and lipids ★ TKIs seem to have limited effect on lipids. The phase III trials with TKI did not show any hyperlipidemic effects. Interestingly, correction of hyperlipidemia has been reported in hyperlipidemic patients treated with imatinib for either CML or hypereosinophilic syndrome (74). In this study, among nine patients with hypercholesterolemia, not treated with hypolipidemic agents, eight of them normalized their plasma cholesterol 30 days after starting treatment with imatinib, with a significant decrease in mean plasma total cholesterol from 254 to 176 mg/dl (74). In addition, four of these nine patients also had hypertriglyceridemia and three of them normalized their plasma triglycerides with a significant decrease in mean plasma triglycerides from 264 to 130 mg/dl (74). The reasons for this corrective effect of imatinib on hyperlipidemia are not clear. It has been suggested that imatinib, by inhibiting the platelet-derived growth factor, may then reduce the cytoplasmic phosphorylation of LDLr-related protein which is involved in lipid metabolism.

TKIs and glucose homeostasis ★ TKIs influence glucose metabolism, with both elevated and decreased blood glucose levels that have been attributed to TKI treatment. The molecular mechanism by which TKI controls glucose homeostasis remains unknown. More disturbing is the observation that the same TKI (for instance, imatinib or sunitinib) can be associated either with hyperglycemic or hypoglycemic events (75). Hyperglycemia has been reported with sunitinib in 15% of patients with metastatic RCC (10), while it was not reported in phase III trials of patients with advanced P-NETs or GIST (Table 2). In contrast, some patients treated by imatinib or sunitinib also presented hypoglycemia and/or regression of long-standing diabetes (12, 76, 77, 79, 80). Among TKIs, the main drug causing hyperglycemia is nilotinib (Table 2). Hyperglycemia is reported as a common adverse event in its prescribing information form, occurring in 38% of patients (G3–4 Z 5%) with CML treated by nilotinib in the phase III trials (14). Owing to their unpredictable action on blood glucose levels, a tight monitoring of glucose homeostasis (with at least HbA1c dosage and self-monitoring of blood glucose if necessary) is required in patients under TKI. It should also be recommended that patients are educated to recognize any symptomatic sign of hypoglycemia or hyperglycemia (e.g. thirst, polyuria). It should also be mentioned that nilotinib was found to be associated with severe progressive peripheral arterial occlusive disease and, to a lesser extent, with other vascular occlusive events (myocardial infarction and stroke) (81). The underlying molecular mechanism

Figure 3

Relationship between mTORC1 activity and metabolic homeostasis. The U-shaped curve indicates that too little or too much mTORC1 activity is associated with insulin resistance and altered metabolic profile. MS, metabolic syndrome; T2DM, type 2 diabetes mellitus. Adapted from Laplante M & Sabatini DM. mTOR signaling in growth control and disease. Cell 2012 149 274–293.
remains unclear, but these arterial events seem to occur in patients with cardiovascular (CV) risk factors (82).

Metabolic consequences of anti-cancer targeted therapies in clinical practice

The main modifications of plasma lipids and glycemia with mTOR inhibitors and TKI are listed in Table 3. Owing to the significant effect of mTOR inhibitors on lipid and glucose metabolism, a specific management of mTOR-induced lipid disorders or hyperglycemia has been recently proposed by a Task Force of the US National Cancer Institute Investigational Drug Steering Committee (54) and a French expert committee (83).

Before initiation of a treatment with an mTOR inhibitor, it is recommended to measure fasting lipids (total cholesterol, triglycerides, and HDL cholesterol), fasting plasma glucose (FPG), and HbA1c. LDL cholesterol level will be calculated using the Friedewald formula (LDL cholesterol = total cholesterol − HDL cholesterol − triglycerides/5 (mg/dl)). In case of plasma triglycerides above 400 mg/day (4.56 mmol/l), the Friedewald formula may not be used and a direct measurement of LDL cholesterol or plasma apoB will be required.

When an abnormal lipid profile (elevated LDL cholesterol and elevated triglycerides) is present at baseline, it is recommended to initiate an appropriate hypolipidemic treatment before starting the mTOR inhibitor therapy. If the baseline lipid profile is normal, fasting lipid levels will have to be checked every 2 weeks during the first mTOR inhibitor treatment cycle and then with every mTOR inhibitor treatment cycle. During treatment with mTOR inhibitor, a treatment with statin will have to be initiated if plasma LDL cholesterol is above 190 mg/dl (4.90 mmol/l) when one or no CV risk factors are present (age < 50 for male, age < 60 for female, family history of coronary artery disease, hypertension, diabetes, smoking, HDL cholesterol < 40 mg/dl or 1.0 mmol/l) or above 130 mg/dl (3.35 mmol/l) when two or more CV risk factors are present, or above 100 mg/dl (2.58 mmol/l) in case of history of previous CV disease. The recommended LDL cholesterol goals are < 130 mg/dl (3.35 mmol/l) when

<table>
<thead>
<tr>
<th>Drug</th>
<th>Types of cancer</th>
<th>n</th>
<th>Hyperglycemia (%)</th>
<th>Hypoglycemia (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>All grades Grades 3–4 All grades Grades 3–4</td>
<td></td>
</tr>
<tr>
<td>Nilotinib</td>
<td>Chronic myeloid leukemia</td>
<td>556</td>
<td>38 5</td>
<td>ND ND</td>
</tr>
<tr>
<td>Imatinib (14)</td>
<td></td>
<td>280</td>
<td>20 0</td>
<td>ND ND</td>
</tr>
<tr>
<td>Nilotinib</td>
<td>Gastrointestinal stromal tumor</td>
<td>165</td>
<td>ND ND</td>
<td>ND ND</td>
</tr>
<tr>
<td>Control (84)</td>
<td></td>
<td>83</td>
<td>ND ND</td>
<td>ND ND</td>
</tr>
<tr>
<td>Pazopanib</td>
<td>Renal cell carcinoma</td>
<td>290</td>
<td>41 2</td>
<td>17 1</td>
</tr>
<tr>
<td>Placebo (12)</td>
<td></td>
<td>145</td>
<td>33 1</td>
<td>3 0</td>
</tr>
<tr>
<td>Pazopanib</td>
<td>Soft tissue sarcoma</td>
<td>239</td>
<td>ND ND</td>
<td>ND ND</td>
</tr>
<tr>
<td>Placebo (13)</td>
<td></td>
<td>123</td>
<td>ND ND</td>
<td>ND ND</td>
</tr>
<tr>
<td>Sunitinib</td>
<td>Renal cell carcinoma</td>
<td>375</td>
<td>15</td>
<td>ND ND</td>
</tr>
<tr>
<td>Interferon α (10)</td>
<td></td>
<td>360</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sunitinib</td>
<td>Pancreatic NETs</td>
<td>83</td>
<td>ND ND</td>
<td>ND ND</td>
</tr>
<tr>
<td>Placebo (11)</td>
<td></td>
<td>82</td>
<td>ND ND</td>
<td>ND ND</td>
</tr>
<tr>
<td>Sunitinib</td>
<td>Gastrointestinal stromal tumor</td>
<td>202</td>
<td>ND ND</td>
<td>ND ND</td>
</tr>
<tr>
<td>Placebo (9)</td>
<td></td>
<td>102</td>
<td>ND ND</td>
<td>ND ND</td>
</tr>
</tbody>
</table>

Table 2 Incidence of hyperglycemia and hypoglycemia induced by tyrosine kinase inhibitor (TKI) in phase III studies.

Table 3 Modification of plasma lipids and glycemia with mTOR inhibitors and tyrosine kinase inhibitors (TKIs).

<table>
<thead>
<tr>
<th></th>
<th>mTOR inhibitors</th>
<th>TKIs</th>
</tr>
</thead>
<tbody>
<tr>
<td>Total cholesterol</td>
<td>↑</td>
<td>↓ (↓ in one study with imatinib)</td>
</tr>
<tr>
<td>Triglycerides</td>
<td>↑</td>
<td>(↓ in one study with imatinib)</td>
</tr>
<tr>
<td>LDL cholesterol</td>
<td>↑ or ↑</td>
<td>↑</td>
</tr>
<tr>
<td>HDL cholesterol</td>
<td>↑</td>
<td>↑</td>
</tr>
<tr>
<td>Glycemia</td>
<td>↑</td>
<td>↑ or ↓</td>
</tr>
</tbody>
</table>

NETs, neuroendocrine tumors; ND, not described (incidence are < 10%). Common terminology criteria for adverse events (CTCAE) grades for hyperglycemia: Grade 1, fasting blood glucose between 126 and 160 mg/dl (7.0–8.9 mmol/l); Grade 2, fasting blood glucose between 161 and 250 mg/dl (8.9–13.9 mmol/l); Grade 3, blood glucose between 251 and 500 mg/dl (13.9–27.8 mmol/l); Grade 4, blood glucose > 500 mg/dl (27.8 mmol/l). CTCAE grades for hypoglycemia: Grade 1, blood glucose between 55 mg/dl (3 mmol/l) and the lower limit of the normal range; Grade 2, blood glucose between 40 and 54 mg/dl (2.2–2.9 mmol/l); Grade 3, blood glucose between 30 and 39 mg/dl (1.7–2.1 mmol/l); Grade 4, blood glucose < 30 mg/dl (1.7 mmol/l).
one or no CV risk factors are present or <100 mg/dl (2.58 mmol/l) when two or more CV risk factors or history of previous CV disease are present (84). It is recommended to use pravastatin, fluvastatin, or rosuvastatin in patients on mTOR inhibitor therapy. The use of simvastatin or atorvastatin is not recommended because they may interact with mTOR inhibitors (via competitive inhibition of cytochrome P 3A4).

Patients on mTOR inhibitor therapy showing fasting plasma triglycerides above 400 mg/day (4.56 mmol/l) should be instructed in therapeutic lifestyle changes including limitation of alcohol and high-sugar food. LDL cholesterol-lowering therapy with a statin will have to be considered if plasma LDL cholesterol is above the recommended level (see above). If plasma triglycerides remain elevated (above 400 mg/dl or 4.56 mmol/l), a treatment with fenofibrate and/or omega 3 fatty acids should be considered, because of the risk for acute pancreatitis.

As far as glucose metabolism is concerned, baseline measurement of FPG and HbA1c will allow to diagnose patients with prediabetes (FPG, 110–126 mg/dl or 6.0–7.0 mmol/l; HbA1c, 6.0–6.4%) or diabetes (FPG, ≥126 mg/dl or 7.0 mmol/l; HbA1c, ≥6.5%). In normoglycemic patients, FPG should be checked every 2 weeks during the first month of treatment and then monthly. In addition, HbA1c will be measured every 3 months in all patients. In patients with prediabetes and/or metabolic syndrome, once-daily self-monitoring blood glucose (SMBG) assessment should be recommended. For the patients with known diabetes, SMBG should be intensified. It is recommended to initiate a treatment with metformin, the first-line drug for management of hyperglycemia, when FPG is above 126 mg/dl (7.0 mmol/l), when plasma glucose is above 200 mg/dl at any time, or when HbA1c is above 6.5%. The recommended HbA1c target for these patients is between 7.0 and 8.0%. Ketonuria should be checked when blood glucose >250 mg/dl to early diagnose diabetic ketoacidosis. When plasma glucose is above 250 mg/dl (13.8 mmol/l), patients should rapidly be referred to an endocrinologist or diabetologist for the management of hyperglycemia (84).

Because treatment with TKI may be associated with hypo- or hyperglycemia, it is recommended to measure FPG and HbA1c before initiating the TKI treatment. FPG will be checked every 2 weeks during the first month and then once every month. HbA1c will be measured every 3 months. In a patient with known diabetes, SMBG will be reinforced.

Conclusion

Targeted therapies are widely used in oncology for the treatment of a variety of malignancies. Considering the results from the phase III trials, it is obvious that some of these targeted therapies, especially mTOR inhibitors, are associated with an increased risk of metabolic complications. This risk should be anticipated by the identification of high-risk patients and managed by a close monitoring of metabolic parameters, in tight collaboration with physicians specialized in diabetes and endocrinology.

Declaration of interest
The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the review reported.

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