

Expert Opinion

Rapalogs in viral cancers

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At present, 150 clinical trials are registered with the National Cancer Institute, which investigate the efficacy of inhibitors of the PI3K/Akt/mTOR pathway against multiple cancers. Efficacy varies not so much with drug action, but with tumor type, as different cancer types (and different pre-clinical models) exhibit widely differing susceptibilities to mTOR inhibitors, such as rapamycin. Viral cancers appear to be among the most mTOR-addicted and most rapamycin-sensitive cancers. We discuss the different mTOR inhibitors that are currently available and in clinical trials. We also speculate how the molecular makeup of viral cancers could guide the selection and use of known and novel mTOR inhibitors to treat virus-associated malignancies.

Keywords: BEZ235, everolimus, Kaposi sarcoma, mTOR, rapalog, rapamycin, sirolimus, virus

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The term 'Rapalogs' describes a class of allosteric inhibitors that target the mammalian target of rapamycin (mTOR) pathway. More recently, ATP-competitive inhibitors of mTOR as well as so-called 'dual-kinase' inhibitors have been developed, which target other kinases in the mTOR signaling cascade in addition to mTOR itself. Rapamycin (sirolimus) was discovered in the 1970s, and is in widespread use as a second-generation oral immune suppressant in solid organ transplantation. Rapamycin inhibits IL-2 translation and secretion in T cells and thus T cell proliferation (Figure 1). In addition, it also inhibits IL-2-dependent (and other ligand)-dependent signaling in the same cells. In this context, the cell-autonomous G₁ arrest phenotype induced by protein translation arrest is augmented by inhibition of IL-2, which is a paracrine and autocrine growth factor for T cells. The first-generation immune suppressants, cyclosporine and FK506, also inhibit IL-2 expression in T cells and thereby T cell proliferation. However, their inhibition is T cell specific, because the inhibitory mechanism ultimately depends on NFAT (nuclear factor of activated T cells), a T cell lineage-restricted transcriptional transactivator of the IL-2 promoter. By contrast, rapalogs inhibit the ubiquitously required mTOR kinase and thereby inhibit protein translation in all cell types, including cancer cells.

Rapamycin is 'tumorstatic' rather than 'tumortoxic' because mTOR controls protein synthesis and volume growth rather than DNA replication-driven cell proliferation. This mechanism of action limits rapamycin's potency as an anti-cancer agent, except in those cancers where mTOR does not just regulate translation in general, but regulates translation of specific autocrine-acting cytokines required for cancer cell survival. Virus-associated cancers (predominantly herpesvirus-associated B and T cell lymphomas) are examples of this tumor class. Here, rapalogs display nanomolar IC₅₀s in cell culture and in pre-clinical models [1-5]. The efficacy of rapalogs against other subtypes of cancer have been observed in clinical trials, notably in sarcomas, mantle cell lymphoma and renal cell carcinoma, and most dramatically in Kaposi sarcoma (KS), which is associated with human herpesvirus 8 or Kaposi sarcoma-associated herpesvirus (KSHV).

In transplant-associated KS, switching from the immunosuppressant drug cyclosporine A to the immunosuppressant drug rapamycin (sirolimus) resulted in resolution of cutaneous KS [6]. All tumor lesions disappeared but graft function did not decline.

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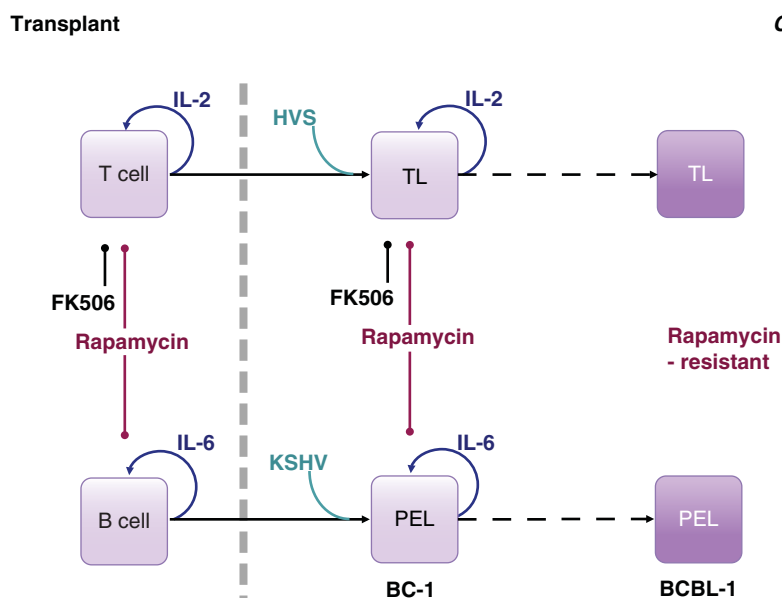


Figure 1. Model of rapamycin modes of action in transplantation (left) and cancer (right). If used as immune suppressants in solid organ transplantation, both rapamycin and FK506 inhibit translation of essential cytokines for activated T cells (IL-2). Rapamycin also inhibits the translation of essential cytokines for activated B cells (IL-6). If used as anti-cancer drugs for viral cancers, both rapamycin and FK506 inhibit IL-2 in herpesvirus saimiri (HVS)-induced T cell lymphoma (TL). Rapamycin also inhibits IL-6 in KSHV-induced primary effusion lymphoma (PEL). Eventually, clones of TL and PEL evolve, which no longer depend on IL-6 or in which IL-6 expression is rapamycin insensitive [3].

This study thus separated rapamycin's immunosuppressive function (on T cells) from its anti-cancer effects on the endothelial lineage tumor KS. Since then, similar results have been reported by others [7,8], although exceptions have been noted as well [9]. Discordant case studies are part of the norm, particularly in a highly pre-treated patient population. This should not detract from the general mechanism. A randomized clinical trial to formally establish the efficacy of any rapalog against KS is still missing.

KS tumor cells are firmly addicted to mTOR signaling. KS lesions are characterized molecularly by high-level phosphorylation of Akt, mTOR and the mTORC1 targets, p70 S6 kinase and ribosomal protein S6 [6,10,11]. In other systems, rapamycin blocked focus formation induced by oncogenic alleles of PI3K or of Akt [12]. These observations place mTOR downstream of, and epistatic to, PI3K and Akt.

Modern mTOR inhibitors promise to improve on the clinical efficacy of rapamycin in several ways. The first class of modern mTOR inhibitors or rapalogs are allosteric inhibitors of mTORC1. They display better bioavailability and pharmacokinetics than sirolimus, but they follow the same molecular mechanism. Everolimus, temsirolimus and ridaforolimus form a mTORC1:FKBP:rapalog complex analogous to rapamycin (sirolimus). Prior binding to FKBP is required and mTORC1 is the direct target of inhibition; a second complex, mTORC2, is not affected. The interactions are a bit more complicated, since the same catalytic subunit of mTOR kinase

participates in both mTORC1 and mTORC2 complexes, with specificity being contributed by different co-factors (raptor in case of mTORC1 and rictor in case of mTORC2). Depending on tumor type and stoichiometry, this can lead to opposing indirect effects: sequestering mTOR kinases into an inactive mTORC1:FKBP:rapalog complex may deplete mTOR and thus diminish mTORC2 kinase activity as well; alternatively mTORC2 activity may increase and in effect compensate for loss of mTORC1 through feedback activation of Akt [13]. Which scenario ultimately prevails is cell type-specific and may help explain the wide range of rapamycin responsiveness among different tumors.

Direct clinical comparisons among the rapalogs have not been reported, but in our animal studies, all allosteric-acting rapalogs exhibit equivalent efficacy across a wide range of tumor xenograft models (unpublished observation); cell lines/tumor types that exhibit a high IC_{50} against rapamycin tend to also be partially resistant to allosteric-acting rapalogs.

The second class of modern mTOR inhibitors target the active site of mTOR kinase (Torin/PP242/Ku-0063794/WYE-354) and in some cases have been shown to inhibit tumor cells more effectively than rapamycin [14]. It appears that active site mTOR inhibitors have weaker effects on the proliferation and function of normal lymphocytes than rapalogs. Additionally, one would expect active site inhibitors to be associated with the rapid development of resistance as well as correspondingly greater

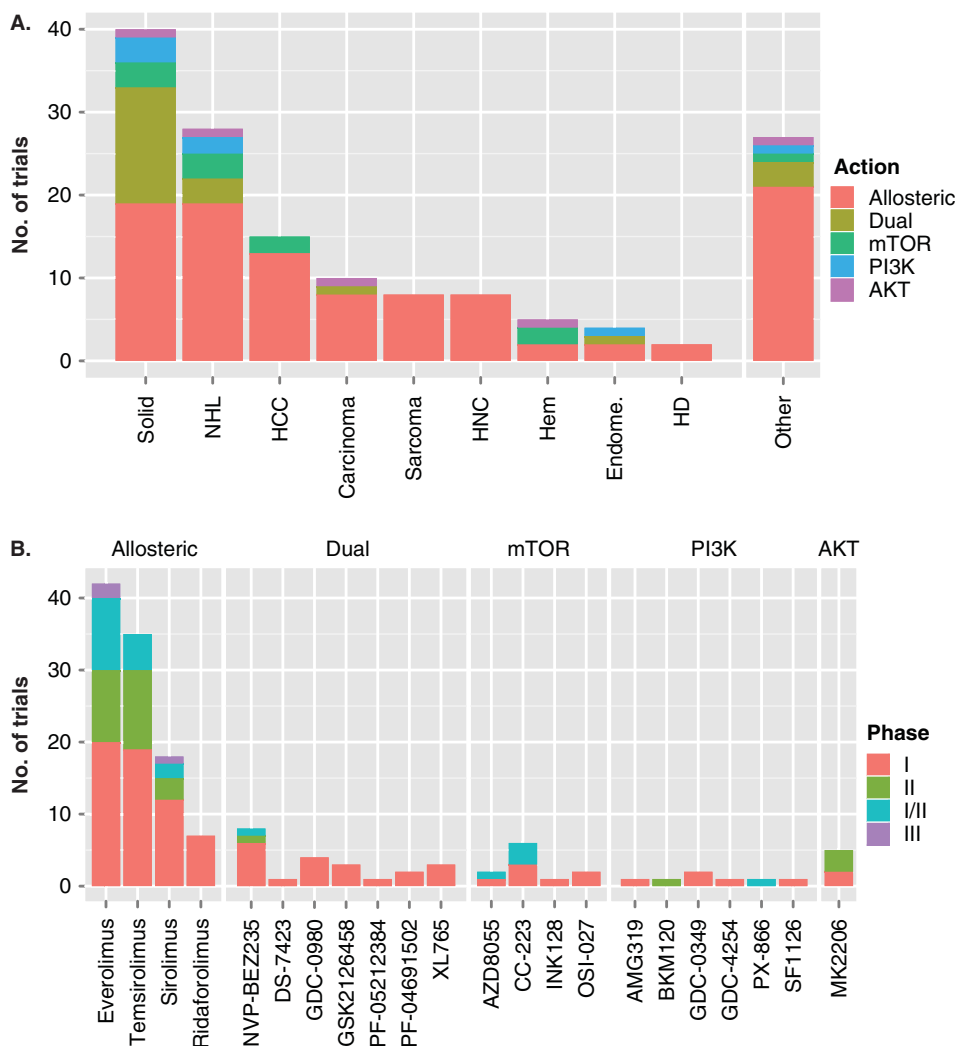


Figure 2. Current NCI registered clinical trials using inhibitors of the PI3K/Akt/mTOR pathway. (A) Stacked bar chart showing the number of trials (on the vertical axis) for different cancer categories (on the horizontal axis). The class of inhibitor is indicated by color. NHL, non-Hodgkin lymphoma; HCC, any liver cancer including hepatocellular carcinoma; HNC, head and neck cancer; Hem, hematopoietic malignancies including multicentric Castlemann disease and leukemia; endome., endometrial cancer; HD, Hodgkin disease. **(B)** Stacked bar chart showing the number of trials (on the vertical axis) for different mTOR inhibitors (on the horizontal axis) grouped by mechanism of action. The phase of the trial is indicated by color.

potential for dose-limiting side effects than rapalogs, which are allosteric inhibitors.

Some active site mTOR inhibitors (e.g., NVP-BEZ335, PI-103) also inhibit PI3K. Dual inhibitors like NVP-BEZ235 are being tested in clinical trials (Figure 2) (reviewed in [15]). These tend to be associated with greater efficacy as well as a wider range of susceptible tumor types based on pre-clinical experiments [1,2,14,16-18]. The exact molecular mechanism for this increased potency probably lies in the expanded range of kinase targets that are inhibited. Inhibiting PI3K affects many more targets (e.g., mTOR-independent targets of Akt). Additionally, PI3K inhibition results in Akt inhibition as well. Consequently, mTORC2-driven Akt activation (due to mTORC1 inhibition-induced feedback) should also be

suppressed. At present, the biological significance of only a few targets of mTORC1 and mTORC2 has been established. A very recent study combined an allosteric rapalog with a dual PI3K/mTOR active site inhibitor [14,19], which resulted in additive efficacy. It remains to be seen if the increased tumor efficacy of dual inhibitors is associated with increased suppression of normal lymphocyte proliferation.

Recently, a new molecular phenotype of mTOR inhibitors has been uncovered. Inhibition of mTOR either by allosteric or active site inhibitors, as well as the dual PI3K/mTOR inhibitors has been shown to induce autophagy [20,21]. In cancer cells, autophagy is one mechanism used by tumor cells to survive anti-neoplastic chemotherapies. Interestingly, although rapamycin by itself could induce autophagy in glioma cells, competitive mTOR inhibitors

like Ku-0063794 induced autophagy to a greater degree than rapamycin. Chloroquine is an inhibitor of autophagy and the combination of chloroquine with dual inhibitors of PI3K and mTOR was found to effectively induce apoptosis in glioma [21]. Thus, the addition of autophagy inhibitors to PI3K/mTOR inhibitors may be warranted in the future.

Currently, extensive safety studies on active site mTOR inhibitors or PI3K/mTOR dual inhibitors have not been reported. Hence, our predictions of clinical utility are highly speculative. Several questions remain to be answered. Should we prioritize individualized therapy and find out when, where and why the established, safe allosteric rapalogs work? Or should we bet on the power of combinatorial therapy, assuming that dual PI3K/mTOR inhibitors will be inherently

more efficacious? What is the safety profile when combination inhibitors are added to the dual inhibitors? In this era of targeted therapies, targeting multiple pathways critical for tumor cell survival is warranted. However, as we target more pathways, the potential for side effects and cytotoxicity is greatly increased and the safety profiles of such combinations need to be considered in equal proportion to their efficacy.

Declaration of interest

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