Pharmacology in the Era of Targeted Therapies: The Case of PI3K Inhibitors

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The PI3K pathway is often aberrantly activated in estrogen receptor positive (ER+), breast cancer and therapies combining PI3K inhibitors and antiestrogens are under clinical development. Given that many PI3K inhibitors have substantial toxicities with continuous dosing and that alternate dosing schedules are equally active, further clinical exploration is warranted. Clin Cancer Res; 22(9): 1–3. ©2016 AACR.

In this issue of *Clinical Cancer Research*, Yang and colleagues explore alternative dosing schedules of PI3K inhibition that elicit high antitumor activity (1). Their observations provide a mechanistic rationale for intermittent dosing of PI3K inhibitors in the management of ER+/PIK3CA–mutant breast cancer that should be considered in clinical practice.

At the height of the development of powerful anticancer cytotoxic therapies, the field of cancer pharmacology was characterized by the study of meticulously detailed dose schedules that maximized the benefit/toxicity ratio, particularly for multiple agent combination regimens (2). Currently, in the age of targeted therapies, this attention to classical pharmacology has been somewhat lost for a number of reasons. Targeted agents were anticipated to have a favorable safety profile because they preferentially target oncogenic drivers and even in some cases, like with mAbs, their prolonged half-life did not allow for flexible scheduling. And yet, the promise of high target specificity is not usually the case as many of these targeted agents have substantial toxicities, highlighting the importance of dose and scheduling. Several studies have indeed compared tumor regression achieved by transient potent inhibition versus continuous inhibition of oncogenic drivers (3–5). For example, transient potent BCR-ABL inhibition by the oral ABL kinase inhibitor imatinib, which is frontline therapy for chronic myeloid leukemia, is sufficient to induce tumor regression in cell culture models and in patients (3). But overall, with the majority of these agents, a continued schedule had been favored over alternative dosing schedules because of a general assumption that clinical success might require prolonged target inhibition.

A particular example of a class of targeted therapies that has shown higher than anticipated toxicities with continuous administration is the case of PI3K inhibitors. This increased toxicity is particularly true for pan-PI3K inhibitors such as GDC-0941 and BKM120. In the case of BKM120 (buparlisib), we recently reported the results of the phase III study of BKM120 plus fulvestrant versus placebo plus fulvestrant (6). The study met its primary endpoint of improved progression-free survival but the benefit was modest. The patients who received fulvestrant alone had a progression-free survival of 5 months; those who received buparlisib plus fulvestrant had a progression-free survival of 6.9 months (HR, 0.78; P < 0.001). Patients who had mutant PIK3CA detected in their circulating tumor DNA had much better outcomes if they received buparlisib plus fulvestrant when compared with those who received fulvestrant alone: progression-free survival of 7 months in the buparlisib plus fulvestrant only group versus 3.2 months in the fulvestrant group (HR 0.56; P < 0.001). However, up to 25% of the patients who received buparlisib experienced serious adverse effects, including hyperglycemia and an increase in markers of liver damage leading to treatment discontinuation or dose reductions. Overall, the median exposure to BKM120 was limited to 2 months of therapy due to the observed toxicities. Similar toxicities have also been reported from the FERGI study on GDC-0941 (7). In retrospect, this high toxicity observed is not surprising as the PI3K pathway is also critical for processes including cell growth, proliferation, metabolism, and homeostasis (8).

To deliver the full potential of PI3K inhibitors, two possible nonmutually exclusive solutions to the high toxicity observed with pan-PI3K inhibitors come to mind (Fig. 1). The first is to explore new scheduling alternatives that may be less toxic. A second ongoing approach is to develop isotype-specific PI3K inhibitors that may have a better safety profile. Yang and colleagues (1) have explored in their article the second ongoing approach: studying new scheduling alternatives with pictilisib (GDC-0941), a class I pan-PI3K with a recommended dose of 330 mg once daily (9). They explored different doses and schedules: (i) weekly treatment with a high dose of GDC-0941 (800 mg/kg), (ii) daily treatment with a low dose of GDC-0941 (100 mg/kg), or (iii) bi-daily treatment with a low dose of GDC-0941 for 3 consecutive days per week (100 mg/kg). Transient, complete PI3K inhibition stopped cell growth in vitro more efficiently than...
longer term, continuous PI3K inhibition. Pharmacokinetic analyses revealed that high dose of GDC-0941 increased the drug’s half-life, exposure, and mean residence time. Pharmacodynamics studies demonstrated that high dose GDC-0941 inhibited the PI3K signaling for 72 hours in comparison with 6 to 9 hours for the low dose. Although these different treatment schedules elicited different patterns of pathway activation/inhibition, proliferation, or apoptosis, they all evoked robust anticancer effects in combination with fulvestrant. Hence, both metronomic (daily) and intermittent (weekly) inhibition of PI3K in combination with fulvestrant induces tumor regression. In summary, the work by Yang and colleagues (1) supports the testing of intermittent pathway inhibition in the treatment of ER\(^+\) and PIK3CA-mutant breast cancer. Their critical finding is that potent target inhibition by the PI3K inhibitor remains effective even if it may be achieved transiently. From the clinical point of view, a different range of schedules and doses should be considered and optimized in early-stage clinical trials. This is especially important for those compounds where toxicity has precluded them from continuous therapy. Moreover, further preclinical studies such as those performed by Yang and colleagues can address the temporal response and pharmacodynamic effects of other combinatorial treatments. This would provide rationale for treatment scheduling that might increase the therapeutic window of new agents and combinations. In conclusion, classical pharmacologic principles of dose and schedule continue to be critically important in the era of precision medicine and carefully preclinical exploration of different schedules may enable improved clinical trials design and delivery of efficacious combination regimens.

**Disclosure of Potential Conflicts of Interest**

No potential conflicts of interest were disclosed.
Authors’ Contributions

Conception and design: J. Baselga
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