Old Habits Die Hard: Addiction of BRAF-Mutant Cancer Cells to MAP Kinase Signaling

Catherine B. Meador and William Pao

Summary: Dual and triple combination therapies with RAF inhibitors plus other targeted agents have demonstrated promising clinical utility in BRAFV600E-mutant solid tumors. However, despite vertical inhibition at multiple nodes on the MAPK signaling pathway, resistant tumors emerge. Aharonian and colleagues show that in BRAFV600E-mutant colorectal cancer, resistance involves reactivation of RAS/RAF/MEK/ERK signaling and may be overcome by newly emerging ERK inhibitors. Cancer Discov; 5(4): 348–50. © 2015 AACR.

See related article by Aharonian et al., p. 358 (4).

Mutations in the serine–threonine kinase BRAF at amino acid position V600 occur in 8% to 15% of colorectal cancers. BRAFV600E mutations also occur in other tumor types, such as melanoma, non–small cell lung cancer (NSCLC), and hairy cell leukemia (HCL), suggesting that mutant BRAF could be broadly targeted regardless of histology. Unfortunately, response rates to mutant-specific BRAF inhibitors, such as vemurafenib and dabrafenib, vary by tumor type. Although BRAF inhibitors have achieved a mere 5% response rate in BRAFV600E-mutant colorectal cancer, response rates in melanoma, NSCLC, and HCL are approximately 52%, 32%, and 96%, respectively (1–3). Thus, at least in solid tumors, the presence of a BRAF mutation is not a “slam dunk” for single-agent BRAF inhibition.

Mechanisms of resistance to single-agent BRAF inhibition also differ by histologic subtype. In BRAFV600E-mutant colorectal cancer cells treated with vemurafenib, MAPK signaling is reactivated by the feedback activation of EGFR (Fig. 1A; ref. 3). Amplification of mutant BRAF has also been implicated in preclinical resistance to BRAF inhibitors in this context (4). Consistent with these findings, combined BRAF/MEK and BRAF/EGFR inhibition more effectively inhibits growth of BRAFV600E-mutant colorectal cancer cells than BRAF inhibition alone (3). Other studies have shown that PTK736 mutations and PTEN loss may also contribute to vemurafenib resistance in BRAFV600E-mutant colorectal cancer cell lines (5). Additional molecular alterations across different tumor types (mostly melanoma) that modulate response to BRAF inhibition in BRAFV600E-mutant tumors include RAS mutations, loss of the RAS negative regulator NF1, BRAF splice variants, increased expression of CRAF and COT, MEK1 mutations, RB1 inactivation, and activation of MET, IGF1R, and HER3 receptor tyrosine kinases, among others (3). Clearly, reactivation of MAPK signaling is a recurring theme (3, 5).

Based on the identification that EGFR bypass signaling can mediate resistance to vemurafenib in BRAFV600E-mutant colorectal cancer, clinical trials testing combinations of BRAF/MEK and BRAF/EGFR inhibitors in these patients are under way. Such combinations have demonstrated somewhat improved outcomes relative to BRAF inhibition alone, but acquired resistance still emerges (6, 7). In this issue of Cancer Discovery, Aharonian and colleagues (4) provide the first description of mechanisms of acquired resistance to combined BRAF/MEK and combined BRAF/EGFR inhibition in BRAFV600E-mutant colorectal cancer. By utilizing both in vitro cell line models and clinical samples, they describe multiple distinct mechanisms of resistance to these BRAF inhibitor combinations, all of which reactivate the MAPK signaling pathway.

In vitro modeling of acquired resistance to combined BRAF/EGFR inhibition (vemurafenib/cetuximab) and combined BRAF/MEK inhibition (vemurafenib/selumetinib) resulted in acquired KRASG12D and KRASG13D substitution mutations, respectively (Fig. 1B). The authors further showed that these activating KRAS mutations increased phosphorylation of CRAF, MEK, ERK, and RSK, suggesting that mutant KRAS-mediated constitutive activation of the MAPK signaling pathway is responsible for resistance in this setting. Importantly, the cell line models of KRAS-mediated acquired resistance to combined BRAF/EGFR inhibition displayed cross-resistance to combined BRAF/MEK inhibition, and vice versa.

Aharonian and colleagues (4) also performed whole-exome and/or RNA sequencing of paired pretreatment and postresistance biopsies from 3 separate patients with BRAFV600E-mutant colorectal cancer who progressed on either BRAF/EGFR or BRAF/MEK combination therapy. In the two described cases of acquired resistance to BRAF/EGFR combination therapy, the authors found amplification of BRAF (dabrafenib/panitumumab resistance) and KRAS (encorafenib/cetuximab resistance) to be unique to the resistant tumor (Fig. 1B). Of note, the KRAS amplification was identified in a lesion that had progressed through BRAF/MEK (dabrafenib/trametinib) combination therapy and continued to progress after the patient...
Figure 1. Schematic representation of mechanisms of resistance to BRAF/EGFR or BRAF/MEK inhibitor combinations in BRAF<sup>V600</sup>-mutant colorectal cancer (CRC). A, comparison of response of BRAF<sup>V600</sup>-mutant melanoma versus colorectal cancer to single-agent BRAF inhibition. Left, BRAF inhibitors (BRAFi) effectively inhibit MAPK signaling and induce tumor regression in over half of BRAF<sup>V600</sup>-mutant melanomas, when used as monotherapy. Mechanisms of resistance in nonresponsive tumors are not depicted. Right, BRAF inhibitors result in reactivation of EGFR-mediated MAPK signaling in BRAF<sup>V600</sup>-mutant colorectal cancer, contributing to the low overall response rate to single-agent BRAF inhibition. B, mechanisms of resistance to various combinations of BRAF/EGFR or BRAF/MEK inhibitors were identified by Aharonian and colleagues via in vitro modeling of acquired resistance or analysis of tumor samples after progression on combination therapy. KRAS<sup>G12D</sup>, KRAS<sup>G13D</sup>, or amplification of wild-type KRAS or mutant BRAF were found to confer resistance to dual inhibition of BRAF/EGFR and BRAF/MEK. A MEK<sup>F53L</sup> substitution mutation was found to confer resistance to dual BRAF/MEK inhibition. All models of resistance to combination BRAF/EGFR and BRAF/MEK inhibition retained sensitivity to ERK inhibition.

was switched to the BRAF/EGFR (encorafenib/cetuximab) regimen. This finding further validates the authors’ in vitro result that KRAS-mediated activation of MAPK signaling can confer cross-resistance to both BRAF/MEK and BRAF/EGFR inhibition.

In addition, the authors identified co-occurring acquired missense mutations in MEKI1 and ARAF in a BRAF<sup>V600</sup>-mutant colorectal cancer sample with acquired resistance to BRAF/MEK inhibition (dabrafenib/trametinib). Follow-up in vitro analysis revealed that the MEKI<sup>F53L</sup> mutation, but not the ARAF<sup>Q489L</sup>
In the case of BRAFV600E-mutant colorectal cancer, for example, the addition of an ERK inhibitor to the BRAF/MEK or BRAF/EGFR combination could significantly extend progression-free survival. However, if acquired resistance after the addition of an ERK inhibitor occurs within a similar time frame as seen with BRAF/MEK and BRAF/EGFR inhibitor combinations, a patient could potentially benefit from receiving the dual inhibitor combination first, followed by an ERK inhibitor once the tumor has progressed. The question of which strategy will provide the best outcome for patients can only be answered clinically. In the meantime, studies such as those performed by Ahronian and colleagues (4) are critical to pave the road ahead.

Disclosure of Potential Conflicts of Interest

W. Pao is Global Head, Oncology Discovery and Translational Area, at Roche; reports having received commercial research grants from AstraZeneca, Bristol-Myers Squibb, and Symphogen; and has been a consultant/advisory board member for AstraZeneca, Bristol-Myers Squibb, and ExelisRx. No potential conflicts of interest were disclosed by the other author.

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