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Special Reporting on:

Integrated Genomic Analyses of Cancer

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Tumor Patient

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PLUS Meeting Abstract Summaries

Integrated Genomic Analyses of Cancer

In the Keynote Lecture at the 2010 American Society of Clinical Oncology Gastrointestinal Cancers (ASCO GI) Symposium, Dr. Victor E. Velculescu discussed integrated genomic analyses of cancers.¹ He focused on the recent developments resulting from these technologies, including the challenges with their use and the potential therapeutic implications.

Dr. Velculescu and other researchers have focused their work on understanding the “cancer genome” with the goal of translating these findings to patient care. Dr. Velculescu explained that the development and progression of cancer is a genetic process involving the activation of oncogenes and the inactivation of tumor suppressor genes. However, he called these changes “the tip of the iceberg,” as it is becoming clear that other important changes also occur in cancer cells, including gains and losses of chromosomal arms, amplification of subchromosomal regions, homozygous deletions, and various changes in gene expression. These events suggest that additional driver genes are mediating cancer development.

Using a systematic genome-wide screen for signaling genes associated with human cancers, Davies and colleagues found an association between mutations in the *RAF* gene *BRAF* and various cancers, including malignant melanoma, thyroid cancer, and colorectal cancer.² Subsequently, Samuels and colleagues at The Johns Hopkins Uni-

versity reported that mutations in the *PIK3CA* gene are associated with numerous cancers, including cancer of the colon, breast, liver, brain, stomach, and lung.³ “This currently places *PIK3CA* as one of the most highly mutated oncogenes in human cancer,” explained Dr. Velculescu. He said that the discoveries of *BRAF* and *PIK3CA* highlight the need to study the entire coding region of the human genome, rather than specific genes or groups of genes, in order to identify genes relevant to cancer development.

Additional genome-wide analyses have been undertaken in recent years to try to identify gene mutations present in malignant cells but not in normal tissue from the same individual. These studies were originally undertaken using a database of approximately 13,000 genes. In 2006, the Consensus Coding Sequences of Human Breast and Colorectal Cancers was published identifying 189 genes mutated at a significant frequency in cancer cells.⁴ Most of these genes were not previously known to be altered in tumors.

The most recent studies have evaluated the expression of approximately 23,000 transcripts, representing the majority of protein-coding genes within the genome. Dr. Velculescu said that although a few thousand non-protein-coding genes or small RNA-encoding genes remained to be analyzed, the genes available for analysis represent a good “first start of the important

machinery inside of the genome.” These studies have analyzed gene expression in 11 patients with colorectal cancer, 11 patients with breast cancer, 24 patients with pancreatic cancer, and 22 patients with glioblastoma.

The first step in these studies has been to perform a discovery screen to identify tumor-specific mutations—alterations that are present in malignant cells but not in healthy tissue. Identified genes of interest are then analyzed in a larger panel of 96 tumors. Genes with mutations occurring at a frequency above background levels are then studied at a more detailed level.

In order to understand the significance of genetic alterations in cancer cells, other analyses are also performed. For example, copy number analyses, which identify gene amplifications or losses, can reveal oncogenes or tumor suppressor genes. Newer technologies for assessing copy number include high-density oligonuclear-type microarrays and serial analysis of gene expression coupled with next-generation sequencing. These approaches allow a detailed quantitative measurement of gene expression by analyzing several million tags for each sample.

Dr. Velculescu described some of the findings that have resulted from these gene expression studies. An analysis of single nucleotide substitutions in the tumor samples revealed variations in the type of base changes that occur in different tumor types. For example,

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C:G to T:A transitions represented over half of evaluated colorectal cancer mutations but only 35% of evaluated breast cancer mutations. Conversely, C:G to G:C transversions represented less than 10% of colorectal cancer mutations but 29% of breast cancer mutations.⁵ Dr. Velculescu said that these findings, which are not well understood, suggest the involvement of organ-specific carcinogens or mechanisms of repair.

To analyze genomic alterations on the global level, Dr. Velculescu and colleagues have used a technology called digital karyotyping, which provides a high-resolution analysis of copy number alterations (amplifications and deletions) on a genome-wide scale.

By integrating these various genomic analysis tools, researchers are attempting to identify the genes responsible for driving tumorigenesis in different cancer types. For colorectal cancer, the resulting list of the top 20 candidate genes included some genes already associated with tumor development, such as *RAS*, *p53*, *CDC-4*, and *SMAD-4*, but also identified genes not previously known to be important in any cancer, including kinases, metalloproteinases, and other enzymes.

Dr. Velculescu added that there are challenges with these analyses: many mutations occur at low frequencies, and there is interpatient heterogeneity, with no 2 patients expressing identical alterations. He suggested that it may be advantageous to think about targeting molecular processes or signaling pathways rather than specific genes. For example, in pancreatic cancer, 12 pathways have been identified that are affected in at least two-thirds of all patients. "This is," he said, "the beginning of understanding these pathways and the underlying mechanisms in cancer."

A greater understanding of these pathways could identify therapeutic targets and could be used for diagnostic purposes and for monitoring of disease. For example, Dr. Luis Diaz and colleagues at Johns Hopkins developed a technique for monitoring for recurrence

ABSTRACT SUMMARY Panitumumab With FOLFIRI Versus FOLFIRI Alone as Second-line Treatment in Metastatic Colorectal Cancer

In patients with *KRAS* wild-type metastatic colorectal cancer, the addition of panitumumab to FOLFIRI in the second-line setting provides a significant PFS benefit and a significant improvement in patient-reported outcomes, according to results of an open-label, randomized, global phase III trial presented by Dr. Marc Peeters (Abstract 282). Among the 597 evaluable patients with wild-type *KRAS* tumors, panitumumab plus FOLFIRI was significantly more effective than FOLFIRI alone in regard to median PFS (5.9 vs 3.9 months; HR, 0.73; 95% CI, 0.59–0.90; $P=.004$) and objective response rate (35% vs 10%; $P<.001$), and there was a nonsignificant trend toward improved median OS (14.5 vs 12.5 months; HR, 0.85; 95% CI, 0.70–1.04; $P=.12$). The addition of panitumumab to FOLFIRI provided no benefit among the 486 evaluable patients with mutant *KRAS* tumors. Median PFS in these patients was 5.0 months with panitumumab plus FOLFIRI versus 4.9 months with FOLFIRI alone ($P=.14$), objective response rates were 13% and 14%, respectively, and median OS was 11.8 months and 11.1 months, respectively. The investigators also evaluated the impact of panitumumab on patient-reported outcomes. In *KRAS* wild-type patients, panitumumab was associated with a significant benefit as assessed by the EQ-5D Overall Health Rating (OHR) but not the multidimensional Health State Index (HSI). Panitumumab provided no benefit in patient-reported outcomes in *KRAS*-mutant patients. Dr. Peeters and colleagues reported that the combination of panitumumab and FOLFIRI was well tolerated and revealed no unexpected toxicities. The most common grade 3/4 adverse events in *KRAS* wild-type patients receiving panitumumab plus FOLFIRI were skin toxicities (37%), neutropenia (20%), and diarrhea (14%).

in which a mutation is identified from a resected tumor, and the presence of that mutation can be detected in the plasma at very low levels (1 in 10,000 or lower) in patients developing recurrence after surgery.

Dr. Velculescu said that moving forward, researchers will be expanding their analyses into other gastrointestinal cancers, including gastric cancer, hepatocellular carcinoma, and esophageal cancer, to identify relevant genes and pathways. He concluded that, with the tools available today, the greatest challenge lies not in identifying relevant genes, but in determining the functional and pathway implications of those genes. It is hoped that this

research will lead to novel approaches to the diagnosis, evaluation, and treatment of patients with cancer.

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Pursuit of Novel Treatment Strategies in the *KRAS* Wild-type Tumor Patient

Dr. Cornelius J.A. Punt discussed novel treatment strategies for patients with *KRAS* wild-type colorectal cancer.¹ Today, chemotherapy plus bevacizumab is often considered to be the standard first-line treatment for patients with metastatic colorectal cancer, based on data from randomized trials. However, multiple recent trials have also demonstrated the efficacy of anti-epidermal growth factor receptor (anti-EGFR) therapy combined with chemotherapy in patients with *KRAS* wild-type tumors. Therefore, Dr. Punt posed the question of which is preferable—chemotherapy plus bevacizumab, or chemotherapy plus an EGFR agent? Moreover, which chemotherapy regimen should be used? To address these questions, Dr. Punt first reviewed the key studies evaluating anti-EGFR therapy in metastatic colorectal cancer.

The randomized, phase III Cetuximab Combined with Irinotecan in First-Line Therapy for Metastatic Colorectal Cancer (CRYSTAL) study showed that the addition of cetuximab to FOLFIRI (leucovorin, fluorouracil, irinotecan) in the first-line setting is associated with a modest but significant increase in median progression-free survival (PFS) over FOLFIRI alone (9.9 vs 8.4 months; hazard ratio [HR], 0.696; $P=.0012$) and a more significant increase in median overall survival (OS) (23.5 vs 20.0 months; HR, 0.798; $P=.0093$).² (The complete CRYSTAL results are discussed elsewhere in this report.) In patients with *KRAS* mutant tumors, cetuximab had no detrimental effect, but it also was not beneficial.

In the randomized phase II Oxaliplatin and Cetuximab in First-line Treatment of Metastatic Colorectal

Cancer (OPUS) trial, the addition of cetuximab to FOLFOX (leucovorin, fluorouracil, oxaliplatin) was also associated with a significant improvement in median PFS over FOLFOX alone (8.3 vs 7.2 months) and a nonsignificant trend toward improved OS.³ In this trial, the addition of cetuximab to chemotherapy was detrimental in patients with *KRAS* mutated tumors, as it was associated with a significant decrease in median PFS and a shorter OS compared with chemotherapy alone.

In contrast to these findings, results from A Three-arm Randomised Controlled Trial Comparing Either Continuous Chemotherapy Plus Cetuximab or Intermittent Chemotherapy With Standard Con-

tinuous Palliative Combination Chemotherapy With Oxaliplatin and a Fluoropyrimidine in First Line Treatment of Metastatic Colorectal Cancer (COIN) presented at the 2009 European Cancer Organisation/European Society for Medical Oncology (ECCO/ESMO) meeting failed to show an efficacy benefit with the addition of cetuximab to FOLFOX in the first-line treatment of metastatic colorectal cancer.⁴ There was no difference in median PFS or OS with FOLFOX plus cetuximab compared with FOLFOX alone in patients with *KRAS* wild-type or *KRAS* mutated tumors.

The Randomized Phase III Study of Panitumumab With FOLFOX4 Compared to FOLFOX4 Alone as

ABSTRACT SUMMARY Safety of Panitumumab in Combination With Chemotherapy in Metastatic Colorectal Cancer Patients With Wild-type *KRAS* Tumors

The safety of panitumumab and chemotherapy in patients with metastatic colorectal cancer is consistent and as expected for an EGFR monoclonal antibody in combination with chemotherapy, according to a meta-analysis of 5 multicenter clinical trials presented by Dr. Jean-Yves Douillard (Abstract 409). The most common grade 3/4 adverse events among the 763 patients with *KRAS* wild-type tumors randomized to FOLFIRI plus panitumumab (473 patients) or FOLFOX plus panitumumab (322 patients) were skin-related toxicity (20–37%), neutropenia (15–42%), diarrhea (13–24%), pulmonary embolism (0–8%), nausea (3–5%), dehydration (2–10%), and hypomagnesemia (3–8%). Two patients died due to adverse events: 1 due to diarrhea and 1 due to pulmonary embolism. The evaluated studies included 3 randomized trials (1 phase II and 2 phase III) and 2 single-arm phase II trials. The phase II studies evaluated FOLFIRI plus panitumumab and irinotecan plus panitumumab, whereas the phase III studies evaluated FOLFOX or FOLFIRI with or without panitumumab. Common eligibility criteria included a diagnosis of metastatic colorectal cancer, age of 18 years or older, and adequate organ function.

1st-Line Treatment for Metastatic Colorectal Cancer (PRIME), presented at the same meeting, evaluated the addition of the humanized anti-EGFR antibody panitumumab to FOLFOX in patients with previously untreated metastatic colorectal cancer.⁵ Compared with FOLFOX alone, FOLFOX plus panitumumab was associated with a significant 1.6-month improvement in PFS in patients with *KRAS* wild-type tumors and a significant decrease in PFS and OS in patients with *KRAS* mutated tumors. (Updated data from the PRIME study, presented the same day as Dr. Punt's talk, confirmed these findings.)

In the Randomized Phase III Study of Capecitabine, Oxaliplatin, and Bevacizumab With or Without Cetuximab in Advanced Colorectal Cancer (CAIRO2), Dr. Punt and his colleagues in the Dutch Colorectal Cancer Group evaluated capecitabine, oxaliplatin, and bevacizumab with or without cetuximab in 755 patients.⁶ In the overall population, the addition of cetuximab was associated with a significant reduction in median PFS (9.4 vs 10.7 months). A subgroup analysis by *KRAS* status found no survival differences between treatments in patients with *KRAS* wild-type tumors but a significant detriment in patients with *KRAS* mutated tumors in regards to median PFS (8.1 vs 12.5 months) and median OS (17.2 vs 24.9 months). Dr. Punt noted that toxicity differences did not account for the detrimental effect of cetuximab. However, the incidence of hypertension, an adverse event associated with bevacizumab, was lower in the cetuximab-containing arm, suggesting a negative interaction between the 2 agents.

The Panitumumab Advanced Colorectal Cancer Evaluation (PACCE) study used a similar design, evaluating chemotherapy and bevacizumab with or without panitumumab for the first-line treatment of metastatic colorectal cancer.⁷ Panitumumab was

ABSTRACT SUMMARY Skin Toxicity in Metastatic Colorectal Cancer Patients Taking Panitumumab and FOLFIRI

In a study evaluating the safety of panitumumab and chemotherapy, Dr. Meinolf Karthaus reported results from a single-arm phase II study showing that panitumumab plus FOLFIRI is well tolerated in patients with previously untreated metastatic colorectal cancer (Abstract 429). Skin toxicities reported with the combination of panitumumab and FOLFIRI were similar to those previously observed with panitumumab monotherapy. Among the 145 patients enrolled, 98% developed any-grade skin nail toxicity or ocular toxicity, with 36% developing grade 3/4 dermatologic toxicities. The most common grade 3/4 dermatologic toxicities included rash (10%), acne (10%), paronychia (6%), and dermatitis acneiform (5%). Of the 145 patients evaluated, 59% had *KRAS* wild-type tumors. The median duration of therapy was 6.9 months for these patients and 5.8 months for patients with *KRAS* mutant tumors. The overall incidence and severity of skin toxicity were similar regardless of *KRAS* status. However, grade 3/4 dermatologic toxicities were more common among patients who responded to therapy than in those who did not respond in both *KRAS* strata. In the overall population, 51% of responders developed grade 3 skin toxicity, compared with 19% of nonresponders. The authors noted, however, that no definitive conclusions regarding the relationship of *KRAS* status, skin toxicity, and response could be drawn from this small sample size. Medications used for the treatment and management of skin and nail toxicities included antibiotics/antifungals (62% of *KRAS* wild-type and 69% of *KRAS* mutant patients), steroids (19% and 39%, respectively), antihistamines (14% in both groups), and other agents. While patients started topical treatments upon the development of any skin toxicity, oral antibiotics were incorporated into the treatment plan for patients developing severe acneiform dermatitis and/or paronychia of at least grade 2 severity. The authors also noted that some of the last patients who enrolled in the trial may have received prophylactic skin toxicity treatments.

discontinued after a significant decrease in median PFS was seen in the panitumumab arm versus the control arm (10.0 vs 11.4 months; HR, 1.27; 95% CI, 1.06–1.52). Analyses by *KRAS* status and chemotherapy (oxaliplatin-based vs irinotecan-based) showed that in *KRAS* wild-type patients, the addition of panitumumab to oxaliplatin-based chemotherapy was associated with a decrease in median PFS and OS. Panitumumab was also associated with a reduction in PFS in irinotecan-treated patients, although

the patient numbers in these groups were small.

Panitumumab was associated with worse toxicity in the PACCE study. Among oxaliplatin-treated patients, panitumumab was associated with a higher incidence of grade 3/4 skin toxicity (36% vs 1%), diarrhea (24% vs 13%), infections (19% vs 10%), and pulmonary embolism (6% vs 4%).

Dr. Punt also discussed the 181 study of FOLFIRI with or without panitumumab in the second-line treatment of metastatic colorectal cancer;

ABSTRACT SUMMARY Correlation of Number of Nodes Examined With Colon Cancer Recurrence

The number of nodes examined and the 12-gene colon cancer recurrence score both independently predict recurrence in patients with stage II colon cancer, according to an analysis of the Quick and Simple and Reliable (QUASAR) study presented by Dr. Richard Gray (Abstract 331). QUASAR randomized 3,239 patients with resected colorectal cancer (66% stage II colon cancer) to adjuvant chemotherapy with fluorouracil and leucovorin or observation. The current analysis focused on 657 patients with stage II colon cancer, assessing the prognostic value of the number of nodes examined and the recurrence score, which has previously been validated in patients with stage II colon cancer from this study. Overall, patients had a median of 10 nodes examined (interquartile ratio, 7–14); 37% had at least 12 nodes examined. The number of nodes examined increased over time; the proportion of patients with at least 12 nodes examined increased from 21% in 1994–1995 to 54% in 2002–2003. Several parameters were significantly associated with the increased number of nodes examined, including later year of randomization ($P<.001$), younger age ($P<.001$), deficient mismatch-repair ($P<.010$), and higher tumor grade ($P=.001$). After controlling for these factors plus lymphovascular invasion and recurrence score, the number of nodes examined was significantly associated with recurrence risk ($P=.004$). In a multivariate analysis, there was a significant association between recurrence score and risk of recurrence and between nodes examined and risk of recurrence. Recurrence rates were 25% in patients with ≥ 12 nodes examined, 28% in patients with 8–11 nodes examined, and 37% in patients with < 8 nodes examined.

results were presented at the 2009 ECCO-ESMO meeting. Among patients with *KRAS* wild-type tumors, the addition of panitumumab to FOLFIRI was associated with a significant increase in median PFS (5.9 vs 3.9 months) and a nonsignificant trend toward improved OS.⁸ In patients with *KRAS* mutated tumors, no significant differences in the 2 treatment groups were noted; thus, this study found no detrimental effects of adding an EGFR inhibitor in these patients.

Overall, these trials indicate that the benefit of anti-EGFR antibodies is limited to patients with *KRAS* wild-type tumors. In patients with *KRAS*

mutant tumors, the addition of an anti-EGFR antibody to an oxaliplatin-based regimen is associated with a detrimental effect. Dr. Punt said that “based on CAIRO2 and PACCE, the combination of bevacizumab with cetuximab or panitumumab should not be used.” He added that the currently available data suggest “no outright superiority for anti-EGFR agents over bevacizumab.” The Cancer and Leukemia Group B (CALGB) trial X0405, which is ongoing in patients with *KRAS* wild-type tumors, is randomizing patients treated with FOLFOX or FOLFIRI to bevacizumab or cetuximab or a combination

of the 2. This study should provide important insight into the optimal first-line treatment strategy.

Regarding the role of EGFR-targeted therapy, Dr. Punt concluded that the absolute benefits of anti-EGFR antibodies appear to increase in later-line treatments, with these agents showing the greatest benefit when used as monotherapy in chemotherapy-refractory patients. However, bevacizumab has not been evaluated in this setting, and bevacizumab does appear to be better tolerated than anti-EGFR agents in most patients. “When you want to expose your patients to all of the available drugs,” he explained, “there is still a preference for bevacizumab in first-line, and a role for cetuximab or panitumumab in salvage treatments.”

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Cetuximab Plus FOLFIRI in the Treatment of Metastatic Colorectal Cancer

The addition of the anti-EGFR monoclonal antibody cetuximab to FOLFIRI in the first-line treatment of metastatic colorectal cancer is associated with a significant survival improvement in patients with *KRAS* wild-type tumors, according to results of the open-label, randomized, multicenter phase III CRYSTAL trial (Abstract 281).¹ These findings show, for the first time in a randomized study, that the addition of cetuximab to FOLFIRI is associated with a significant survival benefit compared with FOLFIRI alone, explained Dr. Eric Van Cutsem in his presentation. In this updated analysis, the CRYSTAL investigators also confirmed the predictive value of *KRAS* mutation status across all efficacy endpoints, and identified *BRAF* mutation status as a poor prognostic factor in previously untreated metastatic colorectal cancer.

The CRYSTAL study evaluated the efficacy and safety of cetuximab plus FOLFIRI versus FOLFIRI alone in 1,198 patients with previously untreated EGFR-expressing metastatic colorectal cancer. In a previous publication of the CRYSTAL data, Van Cutsem and colleagues reported that in patients with *KRAS* wild-type tumors, cetuximab plus FOLFIRI was associated with a significant 32% reduction in risk of disease progression ($P=.02$) and a nearly 2-fold increase in the likelihood of tumor response compared with FOLFIRI alone.² The benefit of cetuximab was limited to this subset of patients with *KRAS* wild-type tumors.

At the time of the 2009 publication, *KRAS* mutation status had only been determined in 45% of patients. To further clarify the benefit of cetuximab in this population, the CRYSTAL

investigators analyzed *KRAS* status in additional patients and evaluated outcomes after a longer median follow-up of approximately 46 months. Of 1,063 evaluated patients, 63% (666 patients) had *KRAS* wild-type tumors. Updated efficacy analyses for these patients presented at the 2009 ASCO GI Symposium included data from 316 patients randomized to cetuximab plus FOLFIRI and 350 patients randomized to FOLFIRI alone.

These updated analyses confirmed the benefit of cetuximab in patients

with *KRAS* wild-type tumors. Compared with FOLFIRI alone, cetuximab plus FOLFIRI was associated with a significant improvement in median OS (23.5 vs 20.0 months; HR, 0.798; 95% confidence interval [CI], 0.670–0.946; $P=.0093$), median PFS (9.9 vs 8.4 months; HR, 0.696; 95% CI, 0.558–0.867; $P=.0012$), and objective response rate (57.3% vs 39.7%; odds ratio, 2.07; 95% CI, 1.52–2.83; $P<.0001$).

An analysis of tumor regression showed a qualitative and quantitative

ABSTRACT SUMMARY Cetuximab With Chemotherapy as First-line Treatment for Metastatic Colorectal Cancer

Cetuximab plus chemotherapy is superior to chemotherapy alone for the first-line treatment of patients with wild-type *KRAS* metastatic colorectal cancer, according to a meta-analysis of the CRYSTAL and OPUS trials (Abstract 406). In a pooled analysis of 845 *KRAS* wild-type patients randomized to FOLFOX4 (OPUS) or FOLFIRI (CRYSTAL) with or without cetuximab, the addition of cetuximab was associated with significant improvements over chemotherapy alone in regard to objective response rate (57.3% vs 38.5%; odds ratio, 2.16; 95% CI, 1.64–2.86; $P<.0001$), median PFS (9.6 vs 7.6 months; HR, 0.66; 95% CI, 0.55–0.80; $P<.0001$), and median OS (23.5 vs 19.5 months; HR, 0.81; 95% CI, 0.69–0.94; $P=.0062$). These findings are based on an expanded number of samples evaluable for *KRAS* mutation status in both trials, which included 89% of samples from the CRYSTAL study (1,063 of 1,198) and 93% of samples from the OPUS study (315 of 337). Previous analyses were based on 45% of patients from CRYSTAL and 69% of patients from OPUS. In their analysis, Dr. Claus-Henning Köhne and colleagues also evaluated the effect of *BRAF* mutations on responses to cetuximab. However, as in the CRYSTAL analysis by Dr. Eric Van Cutsem and colleagues, the current analysis found no difference in the cetuximab treatment effect according to *BRAF* mutation status. *BRAF* mutation status appeared to have negative prognostic value, with shorter survival and lower response rates in both treatment arms in the 8% of patients with *BRAF* mutant tumors. However, even in this small subset of patients, there was a trend toward better outcomes with the addition of cetuximab to chemotherapy.

ABSTRACT SUMMARY Neoadjuvant or Adjuvant Chemoradiotherapy and Bevacizumab in Rectal Cancer

Bevacizumab can be added to standard neoadjuvant or adjuvant chemoradiotherapy in most patients with localized rectal cancer and may provide an efficacy benefit, although toxicities are a concern, according to a nonrandomized phase II trial presented by Dr. David Spigel (Abstract 459). Among 66 patients with stage II/III rectal cancer treated with bevacizumab in the neoadjuvant setting (35 patients) or the adjuvant setting (31 patients), 62 patients (94%) remained alive and free of recurrence after a median follow-up of 14 months. Post-combined modality treatment for all patients included FOLFOX6 plus bevacizumab for up to 4 cycles. Those patients with no evidence of disease could continue single-agent bevacizumab for up to 1 year of total treatment. Of the 35 patients who received bevacizumab in the neoadjuvant setting, 4 patients did not undergo surgery due to disease progression, coagulopathy, bowel perforation, and patient request (1 patient each). Of the 31 patients undergoing surgery, 10 patients (29%) had a pathologic complete response and 21 patients had residual disease (17 with gross residual disease and 4 with microscopic residual disease). Grade 3/4 toxicities reported in the neoadjuvant cohort were stomatitis/mucositis (23%), neutropenia (15%), diarrhea (14%), leukopenia (11%), postoperative infection (9%), dehydration (6%), and fatigue (6%). Grade 3/4 toxicities reported in the adjuvant cohort included diarrhea (29%), neutropenia (22%), fatigue (10%), hypertension (9%), dehydration (6%), and stomatitis/mucositis (6%). Eight patients discontinued neoadjuvant therapy due to treatment-related toxicities that included small-bowel perforations, prolonged surgical healing, postoperative wound complications, perirectal abscess, nausea/vomiting/diarrhea, methicillin-resistant *Staphylococcus aureus* infection, and coagulopathy. Five patients receiving adjuvant bevacizumab discontinued therapy due to rectovaginal fistula, perianal infection, hypoxia, hematochezia, and dehydration. The investigators concluded that although “the addition of bevacizumab may add to treatment efficacy . . . bowel perforations, infection, and wound-healing complications are potential serious toxicities that warrant cautious use of these regimens.”

tive improvement in response rate with cetuximab plus FOLFIRI versus FOLFIRI alone, with a 13.9% difference in the best percentage change in lesion size based on the World Health Organization (WHO) criteria.

The analysis also confirmed the predictive value of *KRAS* status, showing significant interactions between treatment outcomes and *KRAS* mutation status for all efficacy measures,

including tumor response ($P=.0005$), PFS ($P=.003$), and OS ($P=.046$).

The investigators also explored the association of another potential biomarker, *BRAF*, for predicting responses to cetuximab and FOLFIRI. *BRAF* is a serine-threonine kinase that is a downstream effector of *KRAS*. Previous data have suggested that *BRAF* mutations, which are present in approximately 8% of colorectal tumors, are predictive of

responses to cetuximab in previously treated patients. Van Cutsem and colleagues determined *BRAF* mutation status in 83% of patients in the CRYSTAL trial (1,000 of 1,198). *BRAF* mutations were detected in 6% of evaluable samples, including in 1 patient with a *KRAS*-mutant tumor. Of the 625 *KRAS* wild-type tumors, 555 (88%) were *BRAF* wild-type and 59 (9%) were *BRAF* mutant.

BRAF mutation status had a clear prognostic value in this study, explained Dr. Van Cutsem in his presentation, noting a striking difference in survival and response rates in both treatment arms. In *KRAS* mutation-positive patients receiving cetuximab and FOLFIRI, median OS was 25.1 months in *BRAF* wild-type patients and 14.1 months in *BRAF*-mutant patients. Median PFS was 10.9 months and 8.0 months, respectively, and the overall response rates were 61.0% and 19.2%, respectively. *BRAF* status was also a negative prognostic factor in patients receiving chemotherapy alone.

Although the small sample size of *BRAF* mutant tumors precludes statistical comparisons of the treatment arms, the poor outcomes in these patients are evident. Dr. Van Cutsem concluded that *BRAF* mutation status “does not appear to be a strong predictive biomarker for the addition of cetuximab to FOLFIRI in the first-line treatment of metastatic colorectal cancer,” considering also the low frequency of *BRAF* mutations in this population.

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Panitumumab With FOLFOX4 Compared to FOLFOX4 Alone as First-line Treatment in Metastatic Colorectal Cancer

In the first-line treatment of metastatic colorectal cancer, the addition of panitumumab to FOLFOX4 appears beneficial in patients with *KRAS* wild-type tumors but detrimental in patients with *KRAS* mutant tumors, according to a prospective analysis of the open-label, randomized, global phase III PRIME trial presented by Dr. Salvatore Siena (Abstract 283).¹

Panitumumab, a fully human monoclonal antibody targeting EGFR, is currently approved for use as a single agent in patients with metastatic colorectal cancer with wild-type *KRAS* tumors. The PRIME trial was designed to evaluate panitumumab in combination with chemotherapy in the first-line setting, and prospectively analyzed outcomes according to *KRAS* status.

The study enrolled 1,183 patients with previously untreated metastatic colorectal cancer who were randomized to FOLFOX4 with panitumumab (593 patients) or FOLFOX4 alone (590 patients). Availability of paraffin-embedded tumor tissue was an eligibility requirement, although EGFR expression and *KRAS* status were not required for study entry. “The role of *KRAS* as a biomarker for clinical outcome was found . . . while this study was enrolling,” explained Dr. Siena in his presentation, “and therefore the decision was made to amend the protocol to focus primarily on the efficacy by *KRAS* status.” Thus, the study was amended prior to efficacy analyses and completion of enrollment to focus on the subset of patients with *KRAS* wild-type tumors.

In the 60% of patients with *KRAS* wild-type tumors, the addition of

cetuximab to FOLFOX4 was associated with a significant 20% reduction in the risk of progression or death over FOLFOX4 alone (median PFS, 9.6 vs 8.0 months; HR, 0.80; 95% CI,

0.66–0.97; $P=.02$). There was also a trend toward an improvement in OS with panitumumab (median OS, 23.9 vs 19.7 months; HR, 0.83; 95% CI, 0.67–1.02; $P=.07$) and a numerical

ABSTRACT SUMMARY Panitumumab Immunogenicity in Metastatic Colorectal Cancer

The development of anti-panitumumab antibodies in patients with metastatic colorectal cancer receiving panitumumab plus FOLFOX or FOLFIRI is rare and independent of *KRAS* status, according to an analysis of 2 phase III trials presented by Dr. Marta Starcevic (Abstract 433). Of 559 patients receiving panitumumab plus FOLFIRI, 22 patients (3.9%) tested positive for anti-panitumumab antibodies via Biacore (16 patients) or enzyme-linked immunosorbent assay (ELISA; 7 patients). However, 19 of these patients (3.8%) had pre-existing antibodies detectable at or before baseline. Thus, only 4 patients (0.8%) had newly developing anti-panitumumab antibodies. The incidence of anti-panitumumab antibodies was similar among *KRAS* wild-type and *KRAS* mutant patients in regards to pre-existing antibodies (4.4% and 3.3%, respectively) and developing antibodies (0% and 1.5%, respectively). Similar trends were observed in FOLFOX-treated patients. Of 558 patients evaluated, 36 (6.5%) had detectable anti-panitumumab antibodies, with 22 patients (4.3%) harboring pre-existing antibodies and 14 patients (3.0%) developing new antibodies. The incidence of anti-panitumumab antibodies was similar among *KRAS* wild-type and *KRAS* mutant patients for pre-existing antibodies (3.9% and 3.7%, respectively) and developing antibodies (3.9% and 1.7%, respectively). “Panitumumab monotherapy is associated with a low rate of immunogenicity . . . the rate appears to be even lower among patients receiving combination chemotherapy,” concluded the researchers in their report. The investigators used a neutralizing antibody bioassay to measure the ability of panitumumab to mediate EGFR phosphorylation in vitro. By this assay, anti-panitumumab antibody activity was detected in 0 of 501 samples from patients receiving panitumumab plus FOLFIRI and in 2 of 470 samples from patients receiving panitumumab plus FOLFOX. The presence of anti-panitumumab antibodies did not appear to alter the safety of the regimens, and the low incidence of these antibodies precluded an evaluation of their impact on efficacy endpoints.

Grade 3/4 Adverse Event	Panitumumab + FOLFOX4		FOLFOX 4	
	<i>KRAS</i> Wild-type (n=322)	<i>KRAS</i> Mutant (n=217)	<i>KRAS</i> Wild-type (n=322)	<i>KRAS</i> Mutant (n=217)
Any event	84	80	69	73
Skin toxicity	36	30	2	1
Neutropenia	42	37	41	47
Diarrhea	18	20	9	10
Neurologic toxicity	16	17	16	17

Table 1. Most Common Grade 3/4 Adverse Events With FOLFOX4 Plus Panitumumab*

*Adverse events that were observed in $\geq 10\%$ of patients in any arm. FOLFOX=leucovorin, fluorouracil, oxaliplatin.

Data from Siena S. Randomized phase III study of panitumumab (pmab) with FOLFOX4 compared to FOLFOX4 alone as first-line treatment (tx) for metastatic colorectal cancer (mCRC): PRIME trial. Paper presented at: 2010 American Society of Clinical Oncology Gastrointestinal Cancers Symposium; January 22-24, 2010; Orlando, FL.

improvement in response rate (55% vs 48%; $P=.07$).

However, in the 40% of patients with *KRAS* mutant tumors, median PFS was significantly shorter with panitumumab plus FOLFOX4 vs FOLFOX4 alone (7.3 vs 8.8 months; HR, 1.29; 95% CI, 1.04–1.62; $P=.02$). There was also a trend toward a shorter median OS (15.5 vs 19.3 months; HR, 1.24; 95% CI, 0.98–1.57; $P=.07$). Dr. Siena said that these results could not be explained by treatment exposure, as the median number of cycles and dose intensity were similar regardless of *KRAS* status.

The adverse event profile of the combination was as expected for an anti-EGFR antibody plus chemotherapy. The most common grade 3/4 adverse events were neutropenia, skin toxicity, diarrhea, and neurologic toxicities. Three patients receiving panitumumab died from treatment-related adverse events, including 2 due to pulmonary embolism and 1 from febrile neutropenia. Two patients developed grade 3 panitumumab-related infusion reactions, for a total incidence of less than 1%.

Subgroup analyses calculated in the subset of patients with *KRAS* wild-type tumors showed a consistent benefit in favor of panitumumab in most planned subgroups. The PFS benefit was not significant in women (HR, 1.00; 95% CI, 0.73–1.39), patients

ages 65 years or older (HR, 1.02; 95% CI, 0.75–1.38), and a small population (38 patients) with Eastern Cooperative Oncology Group (ECOG) performance status 2 (HR, 1.99; 95% CI, 0.96–4.15).

Reference

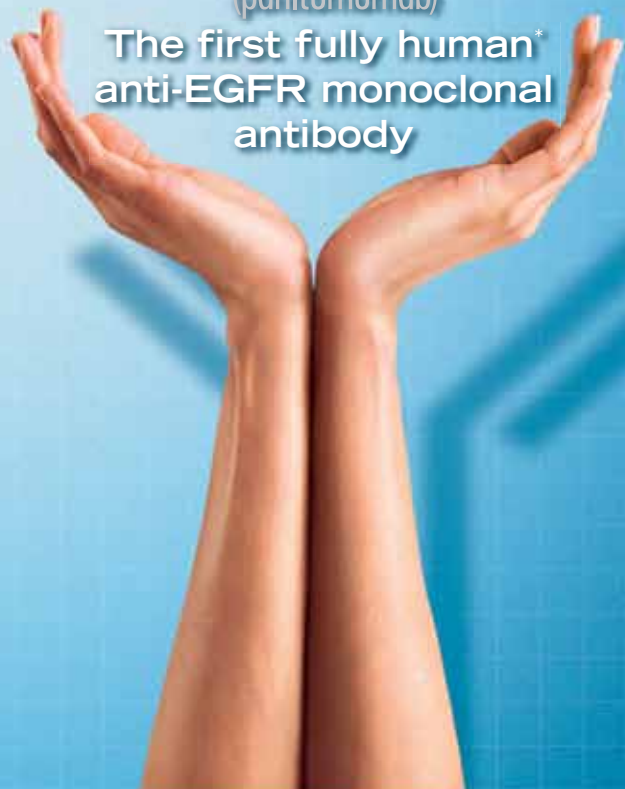
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ABSTRACT SUMMARY Capecitabine Versus 5-Fluorouracil in Colorectal and Gastric Cancers

A meta-analysis of 6,171 patients enrolled in 6 large, multicenter, randomized, noninferiority, phase III clinical trials confirmed the efficacy of capecitabine in patients with colorectal and gastric cancer (Abstract 404). In this analysis, which was undertaken on the advice of European health authorities, Dr. James Cassidy and colleagues compared outcomes in 3,097 patients receiving capecitabine-containing regimens and 3,074 patients receiving 5-fluorouracil/leucovorin (5-FU/LV)-containing regimens. The analysis comprised 3 trials in first-line metastatic colorectal cancer, 1 trial in resected stage III colon cancer, 1 trial in second-line metastatic colorectal cancer, and 1 trial in first-line advanced gastric cancer. In an unadjusted analysis stratified by study, there was no significant difference in median OS with capecitabine (23.1 months) versus 5-FU/LV (22.4 months), with a hazard ratio of 0.94 (95% CI, 0.89–2.00; $P=.0489$). A multivariate Cox regression analysis evaluating the influence of various prognostic factors on OS found that only Eastern Cooperative Oncology Group (ECOG) performance score at baseline was significantly associated with OS. Treatment arm (capecitabine vs 5-FU/LV), age, and sex were not independent prognostic factors. Compared with an ECOG performance score of 0, a score of 1 or higher was associated with a significant increase in the risk of death, with a hazard ratio of 1.56 (95% CI, 1.46–1.66; $P<.0001$). Overall, 66% of 5-FU/LV-treated patients and 67% of capecitabine-treated patients had an ECOG performance score of 0 at baseline; 33% and 32%, respectively, had an ECOG performance score of 1. The investigators concluded that these findings support the “already extensive evidence” regarding the therapeutic equivalence of intravenous 5-FU and oral capecitabine, and suggested that capecitabine can be considered a suitable alternative to 5-FU.

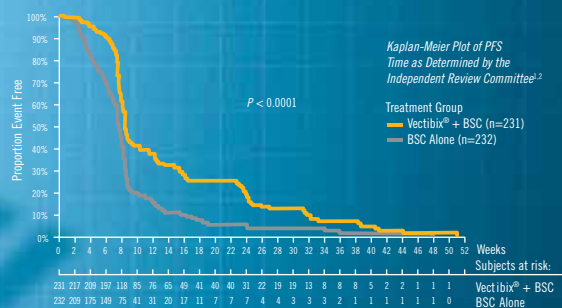


The first fully human* anti-EGFR monoclonal antibody



*Correlation with safety and efficacy is unknown

Based on independent review of disease progression, a statistically significant prolongation in PFS was observed in patients receiving Vectibix® plus BSC vs those patients receiving BSC alone^{1,2}



Prolonged PFS

● Statistically significant prolongation in PFS time vs BSC alone^{1,2}

Q2W dosing

- The recommended dose of Vectibix® is 6 mg/kg administered over 60 minutes (for doses >1000 mg infuse over 90 minutes) as an intravenous infusion every 14 days¹
- Use of premedication was not standardized in clinical trials (the utility of premedication in preventing infusion toxicity is unknown)¹

Infusion reactions

- ~1% incidence of severe infusion reactions reported¹
- Please see Important Safety Information including Boxed WARNINGS for infusion reactions

INDICATION: Vectibix® (panitumumab) is indicated as a single agent for the treatment of epidermal growth factor receptor (EGFR)-expressing, metastatic colorectal carcinoma (mCRC) with disease progression on or following fluoropyrimidine-, oxaliplatin-, and irinotecan-containing chemotherapy regimens. The effectiveness of Vectibix® as a single agent for the treatment of EGFR-expressing mCRC is based on progression-free survival. Currently, no data demonstrate an improvement in disease-related symptoms or increased survival with Vectibix®.

Retrospective subset analyses of metastatic colorectal cancer trials have not shown a treatment benefit for Vectibix® in patients whose tumors had KRAS mutations in codon 12 or 13. Use of Vectibix® is not recommended for the treatment of colorectal cancer with these mutations.

Important Safety Information, including Boxed WARNINGS

WARNING: DERMATOLOGIC TOXICITY and INFUSION REACTIONS

Dermatologic Toxicity: Dermatologic toxicities occurred in 89% of patients and were severe (NCI-CTC grade 3 or higher) in 12% of patients receiving Vectibix® monotherapy. [See Brief Summary: Dosage and Administration, Warnings and Precautions, and Adverse Reactions].

Infusion Reactions: Severe infusion reactions occurred in approximately 1% of patients. [See Brief Summary: Warnings and Precautions and Adverse Reactions]. Although not reported with Vectibix®, fatal infusion reactions have occurred with other monoclonal antibody products. [See Brief Summary: Dosage and Administration].

Vectibix® is not indicated for use in combination with chemotherapy. In an interim analysis of a randomized clinical trial, the addition of Vectibix® to the combination of bevacizumab and chemotherapy resulted in decreased overall survival and increased incidence of NCI-CTC grade 3-5 (87% vs 72%) adverse reactions. NCI-CTC grade 3-4 adverse reactions occurring at a higher rate in Vectibix®-treated patients included rash/dermatitis/acneiform (26% vs 1%), diarrhea (23% vs 12%), dehydration (16% vs 5%), primarily occurring in patients with diarrhea, hypokalemia (10% vs 4%), stomatitis/mucositis (4% vs <1%) and hypomagnesemia (4% vs 0%). NCI-CTC grade 3-5 pulmonary

embolism occurred at a higher rate in Vectibix®-treated patients (7% vs 4%) and included fatal events in 3 (<1%) Vectibix®-treated patients.

In a single-arm study of 19 patients receiving Vectibix® in combination with IFL, the incidence of NCI-CTC grade 3-4 diarrhea was 58%; in addition, grade 5 diarrhea occurred in 1 patient. In a single-arm study of 24 patients receiving Vectibix® plus FOLFIRI, the incidence of NCI-CTC grade 3 diarrhea was 25%.

Pulmonary fibrosis occurred in less than 1% (2/1467) of patients enrolled in clinical studies of Vectibix®. Of the 2 cases, 1 involved a patient with underlying idiopathic pulmonary fibrosis and resulted in death. The second patient had symptoms of pulmonary fibrosis, which was confirmed by CT. Additionally, a third patient died with bilateral pulmonary infiltrates of uncertain etiology with hypoxia.

In the randomized, controlled clinical trial, median magnesium levels decreased by 0.1 mmol/L in the Vectibix® arm; hypomagnesemia (NCI-CTC grade 3 or 4) requiring oral or IV electrolyte repletion occurred in 2% of patients. Patients' electrolytes should be periodically monitored during and for 8 weeks after the completion of Vectibix® therapy.

Exposure to sunlight can exacerbate dermatologic toxicity. It is recommended that patients wear sunscreen and hats, and limit sun exposure while receiving Vectibix®.

The most common adverse events of Vectibix® are skin rash with variable presentations, hypomagnesemia, paronychia, fatigue, abdominal pain, nausea, and diarrhea, including diarrhea resulting in dehydration. The most serious adverse events of Vectibix® are pulmonary fibrosis, severe dermatologic toxicity complicated by infectious sequelae and septic death, infusion reactions, abdominal pain, hypomagnesemia, nausea, vomiting, and constipation.

Please see brief summary of Prescribing Information on next page.

References: 1. Vectibix® (panitumumab) prescribing information, Amgen. 2. Van Cutsem E, Peeters M, Siena S, et al. Open-label phase III trial of panitumumab plus best supportive care compared with best supportive care alone in patients with chemotherapy-refractory metastatic colorectal cancer. *J Clin Oncol.* 2007;25:1658-1664.



Amgen
One Amgen Center Drive
Thousand Oaks, CA 91320-1799
www.amgen.com