Abstracts

deficiency. Whether this protein can only be used as a "surrogate" marker, or is functionally involved in the progression of tumors through this pathway remains to be elucidated.

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FP10

ZEB2 CpG promoter methylation in colorectal cancer is associated with lymph node metastases and invasion in KRAS wild-type patients

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Background. ZEB2 is thought to be involved in epithelial-mesenchymal transition (EMT) through transcriptional repression of E-cadherin and up-regulation of genes including anoikis-resistance factor neurotrophic tyrosine kinase receptor, TrkB. The aim of this study is to determine the clinical relevance of ZEB2 protein expression and ZEB2 promoter methylation in patients with colorectal cancer and their relationship to E-cadherin, TrkB and molecular features KRAS/BRAF mutation and CpG Island Methylator Phenotype

Methods. ZEB2, E-cadherin and TrkB immunohistochemistry was evaluated in 160 colorectal cancer patients using a multi-punch (4-5 tumour punches/patient) tissue microarray. ZEB2 promoter methylation, CIMP, KRAS (codon 12/13) and BRAF (V600E) mutation were determined in 95 patients using pyrosequencing.

Results. High ZEB2 expression was associated with advanced pT (p=0.0308), high-grade tumour budding (i.e., detached tumor cell clusters representing an EMT-like process; p=0.0376) and KRAS mutation (p=0.0141), whereas hypermethylation was frequent in KRAS wild-type cases (p=0.0233). In KRAS wild-type cancers, ZEB2 hypermethylation showed marked relationships with lymph node metastasis (p=0.0004), venous invasion (p=0.0247), lymphatic vessel invasion (p=0.0001), unfavourable survival (p=0.0417; 5-year survival: 27 vs. 74 months) and a positive correlation with TrkB (r=0.40). In KRAS mutated cancers, there were no associations of ZEB2 methylation with clinicopathological features, although a strong inverse correlation between methylation and expression (r=-0.45) was observed. Only weak associations with E-cadherin were noted.

Conclusions. ZEB2 over-expression and CpG methylation occur frequently in colorectal cancers, but may cause their effects in a KRASdependent manner. Among KRAS wild-type patients, methylation of ZEB2 may represent a novel route to tumour cell dissemination.

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FP11

Influence of KRAS mutations on outcome in patients with curatively resected stage III colon cancer treated with adjuvant chemotherapy

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Background. Postoperative chemotherapy improves survival in curatively resected stage III colon cancer (CC) patients and oxaliplatin in combination with fluoropyrimidine-based regimens has been established as a standard treatment. There is no clear evidence for a predictive marker regarding the effect of adjuvant chemotherapy. KRAS mutations could predict oxaliplatin sensitivity in in vitro experiments, as recently demonstrated. Here, we profiled KRAS and correlated it with outcome in stage III CC patients who underwent adjuvant chemotherapy.

Methods. Eligible patients were those with resected stage III CC who underwent 6-months adjuvant chemotherapy, either with single-agent fluoropyrimidine (FP: modulated 5FU or capecitabine) or with oxaliplatin-based regimens (O-FP: FOLFOX or XELOX). DNA extraction was performed on formalin-fixed paraffin-embedded sections, and KRAS mutations were analyzed by direct sequencing. Disease-free survival (DFS) and overall survival (OS) analyses were computed using the Kaplan-Meier method and the Log-rank test. Results. The study population included 261 patients: 115 treated with

FP, 146 O-FP. We identified KRAS mutations in 71/261 (27.4%) cases, of which 33 (46.5%) received FP, and 38 (53.5%) O-FP. In wild-type (wt) KRAS cases, DFS and OS group did not significantly differ between the two treatment modalities [in months, FP vs O-FP-median OS: 62.0 vs 49.6, HR: 1.12 (95% CI: 0.65; 1.92); median DFS: 57.1 vs 44.4, HR: 0.93 (95% CI: 0.55; 1.56)]. In patients treated with FP, a worse DFS [in months, mutant vs wt-median DFS: 46.0 vs 57.1, p=0.04; HR: 1.86 (95% CI: 1.01; 3.41)] and a trend toward a worse OS in months, mutant vs wt—median OS: 56.1 vs 62.0, p=0.08; HR: 1.71 (95% CI: 0.94; 3.10)] were observed in KRAS mutated patients. On the contrary, DFS and OS were not statistically different for mutated and wt KRAS patients treated with O-FP [in months, mutant vs wt—median DFS: 43.7 vs 44.4, p=0.62; HR: 1.17 (95% CI: 0.63; 2.18); median OS: 48.9 vs 49.6, p=0.77; HR: 1.11 (95% CI: 0.55; 2.23).

Conclusions. Our results suggest that curatively resected stage III CC patients exhibiting wt KRAS status might benefit from FP alone thus avoiding oxaliplatin-adverse effects. On the contrary, an oxaliplatin-containing regimen should be recommended in KRAS mutated patients.

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