More Controversy than Ever – Challenges and Promises Towards Personalized Treatment of Gastric Cancer

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Clinical decisions for the treatment of localized gastric cancer have become much more sophisticated and complicated than ever. Two recent large-scale trials published in *NEJM* for East Asian and Western patients strongly support the routine use of adjuvant chemotherapy. However, differences in design, extent of surgery, kind of chemotherapy timing of administration, and survival results in the two trials as well as potential differences in genetic background of Asian and Western gastric cancer patients raise critical questions and grow confusion and uncertainty. Which is the optimum update treatment for Western patients?

Is the Japanese model with standardized D2 surgery followed by one year S-1 chemotherapy applicable in the West and can it produce similar excellent results or should treatment decisions be based on Western patient’s data from the UK MAGIC trial and the USA INT-0116 trial? This editorial approaches this critical question towards a live-saving decision. Emphasis is given to current advances in network biology, cancer genome and functional studies as well as a current comprehensive bench-to-bedside genomic-based protocol for biomarkers-based personalized treatment of gastric cancer.

The landmark ACTS-GC Japanese study demonstrated excellent survival results with primary standardized D2 surgery followed by S-1 chemotherapy for advanced stages II and III. Most patients (89%) had node-positive disease; these advanced tumor stages are associated with poor prognosis in the West. Despite this advanced disease, the overall 3-year survival rate was 70% after D2 surgery alone and 80% in the S-1 chemotherapy group. The hazard ratio for death in the S-1 group, as compared with the surgery-only group, was 0.68 (95% CI, 0.52–0.87; *p* = 0.003). Because of this significant survival difference the trial was stopped.

Local control is now increasingly recognized to have a crucial role in the treatment strategy not only of gastric cancer but also for many other solid tumor including breast cancer despite the use of adjuvant treatment. Appropriate surgery alone or surgery plus radiotherapy for local control not only reduce local recurrence but also improve overall survival. This principle could be achieved by a perfectly conducted standardized D2 surgery. The 109 centers located throughout Japan were selected from among hospitals performing at least 100 operations annually for gastric cancer. The high-quality D2 surgery had resulted in a local relapse rate of only 2.8% and a nodal relapse rate of 8.7%. Adjuvant S-1 chemotherapy significantly reduced all types of recurrence (peritoneal, local, nodal) but questions arise form the finding that systemic treatment did not improve hematogenous relapse (*p* = 0.35).

The ACTS-GC study provides some important messages but also raises several questions. First, can the Japanese excellent results be reproduced in Western patients? There is long-term discussion on whether there are differences between Japanese and
Western gastric cancer and patients. Sakuramoto and colleagues believe that these results can be achieved in the West if a standardized, Japanese-style D2 surgery is performed, because there is no genetic difference between gastric cancers in Japan and Europe. Although this may be correct, for definitive conclusions much more in-depth research work is needed considering gastric cancer genome and interactions between environmental and genetic factors in Japan and Europe or the USA. If all these genetic alterations and environmental variables in carcinogenesis and metastasis are similar, then indeed the Japanese results can be reproduced in the West. It should be noted, however, that survival results for gastric cancer even with adequate D2 surgery in highly specialized Western institutions, are much lower than those reported from Japan.

Second, the Japanese study has demonstrated that a substantial fraction of stage II/III patients had a localized nature of disease. Indeed, the estimated actuarial 5-year relapse-free survival was approximately 50% in the surgery-only group. How can these patients be identified to spare chemotherapy toxicity? Third, the addition of chemotherapy after perfect local and nodal control could increase the survival rate by 10%. How can this small survival benefit be further improved? These two questions are discussed below as they can be approached considering current and future research directions towards the development of both novel biomarkers and targeted agents.

Preoperative and postoperative (perioperative) chemotherapy with epirubicin, cisplatin, and fluorouracil was associated with a 5-year overall survival benefit of 13% (from 23% to 36%) for stage II or III operable gastric cancer patients in the MAGIC trial conducted mainly in the UK. In the USA INT-0116 trial, postoperative chemoradiotherapy improved the 3-year overall survival from 41% to 50%. Both trials have been criticized for inadequate surgery (D0 or D1). Thus, the approximately 30% survival difference between the Japanese trial and Western trials could likely be explained by the fact that chemotherapy or chemoradiotherapy simply compensated mainly surgical failure for local and nodal control. Another reason, beyond the potential differences in body mass index that encourages an “easy” D2 surgery for Japanese patients, cancer genome and environment, is also the inclusion of patients with gastroesophageal junction tumors, which are associated with poorer survival, in the Western trials while such patients were excluded from the Japanese study.

### PERSONALIZED TREATMENT

Worldwide, despite improvements, estimated cure rates for patients with stages IIIA and IIIB remain poor. Given the short mean follow-up of 3 years in the ACTS-GC study and the fact that a subset of patients, approximately 20%, recur after the three first postoperative years, a cure rate of approximately 35% with surgery only and 45% by adding S-1 chemotherapy to D2 surgery could be estimated. These rates are currently the best achieved worldwide; those reported from the West are lower despite D2 surgery and adjuvant treatment. These clinical data strongly indicate the urgent need to develop new therapeutic strategies.

How can long-term survival results be improved? Despite intensive efforts over recent decades, single-gene-based traditional research has produced little clinical success. No robust prognostic or predictive marker has been validated for clinical use despite multiple reports and hundreds of proposals. An isolated success of traditional research towards personalized medicine for various cancer types, including gastric cancer is the prevention and treatment of hereditary diffuse gastric cancer (HDGC). Individuals who are carriers of germ-line mutations in the CDH1 (E-cadherin) gene face a very high lifetime risk of gastric cancer (60–80%). This very small subpopulation can be identified from the very large general population on the basis of family history criteria and genetic testing. Tailoring a risk-reducing prophylactic gastrectomy for these very high-risk individuals can save their lives. Women with CDH1 mutations also have a high risk, approximately 40%, of breast cancer. For these carriers, as well as for women with germ-line BRCA1/2 mutations, prophylactic bilateral mastectomy or close surveillance could be considered.

But the management of sporadic cancer is much more complicated because multiple genes and oncogenic pathways are involved in gastric carcinogenesis. Obviously, single-gene-based traditional research cannot reveal the global signaling pathways network. Gastric cancer is a biologically heterogeneous disease with various molecular tumor subsets that likely respond distinctly differently and require new therapeutics for complete response. In the postgenomic era of network cancer biology, and with the advent of the new high-throughput technologies language, methodologies and research directions should be altered for a faster clinical success.

Currently, a comprehensive bench-to-bedside genomic-based model for personalized treatment of
gastric cancer has been developed and proposed by Roukos. This protocol has been focused on how to overcome current treatment challenges and biases associated with high-throughput analyses. As reported above, the vast majority of patients received an inadequate adjuvant treatment either because they do not need adjuvant systemic treatment, they have a localized nature of disease, or they do not respond to currently used cytotoxic chemotherapy. Focusing on these clinical needs, cancer genome and functional strategies are integrated to reveal the precise role of key genes and interacting pathways. The proposed model can lead to the development of class II prognostic and class III predictive biomarkers. Profiling involved gastric cancer key genes, driver mutations, and interacting pathways will improve our understanding on signaling pathways network. Insights into the presence or absence of small cells subpopulations like cells with high mutation rates or cancer stem-like cells in individual tumors will clarify resistance and sensitivity to cytotoxic and targeted agents. Taken together, the development of both novel biomarkers and targeted agents can be achieved. Prognostic gene signatures might classify patients into low and high metastatic risk. Low-risk patients can successfully be treated by R0 surgery alone, sparing unnecessary chemotherapy toxicity. Among patients at high risk of metastatic relapse, predictive gene signatures can be developed comparing tissue-bank samples from responding and nonresponding patients. The fact that the protocol has considered, from the very initiating stage of design, the current robust clinicopathological factors including the Lauren classification into intestinal-type and diffuse-type gastric cancer, enables the clinical prospective validation of new biomarkers.

**CURRENT CLINICAL PRACTICE**

Although cancer genome and functional studies provide great promise for the future, patients and
Clinicians are currently faced with hard decisions: preoperative or postoperative chemotherapy combined with D2 surgery or D1 plus radiotherapy? Considering the preclinical and clinical evidence, a step-by-step treatment-guided algorithm for Western clinical practice is provided (Fig. 1). Laparoscopy is clinically useful to confirm by biopsy or exclude distant metastasis (peritoneum, liver, distant abdominal enlarged lymph nodes) and serosa status (serosa negative or positive) for more accurate staging than clinical staging with endoscopic ultrasonography, computer tomography or positron emission tomography (PET). Analysis of data from randomized clinical trials (RCTs) and descriptive studies show that, first, survival rates for stage II patients are relatively good not only in Japan but also in the West with appropriate surgery followed by adjuvant treatment. Because preoperative chemotherapy harms some of these patients, primary D2 surgery without delay, converting laparoscopy into an open D2 gastrectomy, now appears to be the preferred approach. Laparoscopic-assisted D2 gastrectomy provides encouraging data but feasibility, safety, and efficacy can be ensured at the present time only in a few highly specialized institutions. Second, because postoperative S-1 chemotherapy for stages IIIA and IIIB did not significantly improve survival in the subgroup analysis of the ACTS-GC study, preoperative cisplatin-based chemotherapy for downstaging and potential early elimination of circulated cancer cells could be considered, although no subgroup analysis from the MAGIC trial is available.

CONCLUSIONS

Gastric cancer still remains a major health problem. Despite improvements in local control and empirical chemotherapy, prognosis particularly for stage III patients remains poor worldwide. New therapeutic strategies are needed.

In the postgenomic era of network cancer biology, clinical practice could be altered when the catalogue of gastric cancer genes has been completed and new high-throughput analyses reveal the whole genome function, including the understanding of noncoding DNA regions. In the meantime, microarray-based and proteomics data profiling genes, mutations, and interacting oncogenic signaling pathways may lead to the development and validation of both biologically targeted agents and prognostic class II and predictive class III biomarkers. Tailoring the best preoperative or postoperative combination of empirical cytotoxic and targeted agents distinctly different in various subsets of individual tumors on the basis of which genes are involved and which pathways are deregulated in a given patient, then indeed personal cancer genome and personalized oncology will revolutionize clinical outcomes not only of gastric cancer but also most solid cancers.

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