

# Writing in PROSE proteomic-based selection for second line treatment in non-small-cell lung cancer

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Since the identification of epidermal growth factor receptor mutations ( $EGFR^{mut+}$ ), biomarker selection of patients for specific targeted agents became the standard of care in advanced non-small-cell lung cancer (NSCLC). Small molecules against the tyrosine kinase domain of EGFR (EGFR TKIs) such as gefitinib, erlotinib or afatinib, induced impressive and durable responses in patients with  $EGFR^{mut+}$ , demonstrating superiority over platinum-based chemotherapy when used in front-line setting (1-3). In addition, previous and recent evidences indicated that in presence of  $EGFR^{mut+}$  an EGFR TKI is preferable to chemotherapy irrespective of line of treatment (4-7). More recently, crizotinib a potent ALK-MET-ROS1 inhibitor demonstrated superiority *vs.* chemotherapy in chemo-naïve and in pretreated NSCLC patients with *ALK* translocations (8,9). Unfortunately, the vast majority of NSCLCs does not harbor any *EGFR* or *ALK* druggable alterations and availability of novel targeted agents is confined to clinical trials. For these patients, platinum-based chemotherapy remains the best front-line treatment, with regimens including pemetrexed and/or bevacizumab for individuals with non-squamous histology (10). Conversely, in second line setting the optimal treatment is not as defined. In 2000, docetaxel was firstly approved for second-line therapy in NSCLC (11) and few years later two other agents, pemetrexed and erlotinib, reached the approval based on the results of phase III trials showing non inferiority in terms of survival *vs.* docetaxel (12) or superiority *vs.* placebo (13). Additional trials, conducted in a general and unselected population of pretreated NSCLC, showed only minimal differences in terms of efficacy among the three drugs, leading to the conclusion that pemetrexed, docetaxel

and erlotinib could be reasonably considered an acceptable option (14-16) in both *EGFR* wild type ( $EGFR^{wt}$ ) or *EGFR* unknown NSCLC. For such reasons, medical oncologists generally based their choice on several factors, including personal experience or familiarity with the drug, toxicity, patient characteristics and preferences, and last but not least, drug costs.

VeriStrat (Biodesix, Boulder, CO, USA) is a serum proteomic test using mass spectrometry, originally developed to identify those NSCLC patients having the best survival outcome with EGFR-TKIs (17). According to the intensity of eight regions in the mass spectra, patients were classified into two categories, good and poor. Previous data have suggested that VeriStrat had a strong prognostic power (18-20) in pretreated NSCLC patients, whereas its predictive role remained controversial (18).

Recently, Greorc *et al.* published the results of the PROSE trial, a phase III study specifically conducted to assess the predictive value of VeriStrat test in the comparison of erlotinib and chemotherapy (pemetrexed or docetaxel) as second line treatment in NSCLC patients. The primary end point was overall survival (OS) (21). Proteomic tests were obtained before starting therapy and results were blinded for patients and for investigators who administered treatment, whereas treatment was blinded for investigators who generated the proteomic stratification. Among 285 subjects included onto the study, the proportion of patients with good and poor proteomic profile was similar in both chemotherapy and erlotinib arm (68% and 72% and 32% and 28%, respectively). In the whole population, there was no significant difference in OS (9.0 *vs.* 7.7 months, HR 1.14, for chemotherapy and erlotinib, respectively), whereas

by splitting results according to the proteomic stratification patients with good profile had a significant longer survival than those classified as poor (11.0 *vs.* 3.7 months,  $P < 0.001$ , HR 2.5). Moreover, treatment with erlotinib associated with shorter survival in patients with poor proteomic classification (3.0 *vs.* 6.4 months,  $P = 0.022$ ; HR 1.72); on the contrary, patients with good proteomic test equally benefited from both chemotherapy and erlotinib treatment (OS 10.9 *vs.* 11.0 months,  $P = 0.714$ , HR 1.06).

Two main conceptual findings emerged from this trial. Authors should be congratulated for their efforts of identifying an alternative and innovative “biomarker” in NSCLC. Actually, PROSE was the first prospective trial in which patient selection was based on proteomic profile of patients rather than on tumor molecular portrait. Second, according to the VeriStrat stratification, they identified a subgroup of patients for which second-line treatment with EGFR TKIs might have a detrimental effect on survival. From this perspective, the PROSE trial seems solving two important medical needs in thoracic oncology. First, VeriStrat stratification could overcome the need of tumor tissue; obviously, in our daily clinical practice, it would be much easier to obtain one microliter of serum suitable for a proteomic test than ten micrometers of tumor slice for molecular analyses. Second, proteomic results could facilitate the choice of a second line treatment in *EGFR*<sup>WT</sup> NSCLC, at least in those patients displaying poor proteomic features.

Unfortunately, the reasons why this trial appears so exciting are the same of its weakness. Proteomic stratification suffers of some limitation. Despite the absolute difference in OS observed in PROSE and in other trials (18-21) is numerically and clinically meaningful, it could be mainly driven by the prognostic effect of proteomic signature. VeriStrat poor classification has been identified in a wide range of cancers including NSCLC, colorectal cancer and head and neck squamous cell carcinoma and in almost all studies poor classification correlated with worse survival (18-22). More interestingly, the proportion of patients having a poor test is invariably of around 30%, irrespective of tumor type, disease stage and line of therapy (22), thus suggesting that for a consistent proportion of patients long-term perspectives are inevitably poor. Although the nature and biologic significance of the proteins detected on proteomic peaks remain largely unknown, it is possible that VeriStrat could measure some “tumor-host” response to the presence of cancer, as demonstrated by the fact that poor classification has not

been observed in other-than-cancer diseases or in healthy individuals (18). Furthermore, the study failed to clarify the other clinical question that is whether chemotherapy should be preferred as second line treatment in good classified patients. Indeed, VeriStrat has been originally designed to refine selection of NSCLC patients more likely to benefit from EGFR TKIs approximately ten years ago, when the predictive role of EGFR mutations was not yet elucidated (17,18); consequently it performs well for EGFR TKIs but it does not for conventional chemotherapy (17). Results of the ongoing phase II trial exploring the significance of VeriStrat test in predicting benefits from first-line platinum based chemotherapy, are urgently awaited in order to define if the proteomic signature could explained different sensitivity to chemotherapeutic drugs (NCT 02055144, [www.clinicaltrials.gov](http://www.clinicaltrials.gov)).

Finally, although the aim of the trial was to evaluate the predictive value of a serum test and not to directly compare chemotherapy *vs.* erlotinib in a general population, Greco *et al.* results seemed not add substantial changes in current state of the art. In unselected NSCLC second line treatments have only modest efficacy (10). In addition, as suggested by two phase III trials (14,16) and in two meta-analyses (7,23), when *EGFR* status was assessed by using a high sensitive method reducing the risk of including false negative patients, chemotherapy resulted superior to EGFR TKIs in terms of PFS confirming that, once again, *EGFR* status cannot be ignored.

Fortunately, the scenario of NSCLC therapy is rapidly evolving and new therapies potentially more effective are emerging. On January 11<sup>th</sup> a press-release by Bristol-Myers Squibb announced that the CheckMate-017, a randomized phase 3 study evaluating the checkpoint inhibitor nivolumab *vs.* docetaxel in previously treated squamous lung cancer met its endpoint, demonstrating increased OS in immunotherapy arm ([news.BMS.com](http://news.BMS.com)). It is possible that in coming years, the role of proteomic test should be re-interpreted on the light of the emergent treatment strategies. We hope that in the next future new therapeutic options and more effective predictive tests will be available for our patients offering the concrete possibility to extend their survival.

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