

INTERVIEW

Panos Fidias, MD

Dr Fidias is Assistant Professor of Medicine at Harvard Medical School and Medical Director of the Inpatient Oncology Unit at Massachusetts General Hospital's Thoracic Oncology Service in Boston, Massachusetts.

Tracks 1-15

Track 1	Implementing the SNaPshot [®] multiplexed genotyping of NSCLC into routine clinical practice	
Track 2	Response of advanced ROS1- rearranged NSCLC to crizotinib	
Track 3	Vemurafenib in the treatment of advanced BRAF-mutant NSCLC	
Track 4	Early clinical trials of maintenance therapy in advanced NSCLC	
Track 5	PARAMOUNT study of maintenance pemetrexed after cisplatin/pemetrexed for advanced nonsquamous NSCLC	
Track 6	AVAPERL study of maintenance pemetrexed/bevacizumab after first-line therapy for advanced nonsquamous NSCLC	
Track 7	Practical benefits of maintenance therapy compared to second-line chemotherapy	
Track 8	Case discussion: An otherwise healthy 80-year-old woman and former smoker with bilateral lung adenocarcinoma and mediastinal adenopathy with wild-type biomarkers receives carboplatin, pemetrexed and bevacizumab followed by maintenance pemetrexed/	

- Track 9 Reducing dose intensity of therapy in very elderly patients with advanced NSCLC
- Track 10 Selection and duration of maintenance therapy for advanced EGFR wild-type NSCLC
- Track 11 Nanoparticle albumin-bound (*nab*) paclitaxel in the treatment of advanced NSCLC
- Track 12 EGFR expression and the use of cetuximab in advanced squamous cell NSCLC
- Track 13 Case discussion: A 45-year-old man and never smoker undergoes craniotomy and whole-brain radiation therapy for an exon 19 EGFR mutation from a lung adenocarcinoma and receives afatinib on the LUX-Lung 2 clinical trial
- Track 14 Clinical implications of recent Phase III studies — LUX-Lung 3, IPASS, OPTIMAL and EURTAC — of EGFR TKI therapy versus chemotherapy as firstline treatment for EGFR-mutant NSCLC
- Track 15 Viewpoint on the SELECT study results with adjuvant erlotinib in resected EGFR-mutant NSCLC

Select Excerpts from the Interview

bevacizumab

📊 Track 1

DR LOVE: Would you comment on the issue of multiplex genomic testing and the approach used in your oncology group at Mass General?

DR FIDIAS: The platform we use is called SNaPshot. It's a multiplex DNA sequencing platform that began as a research tool in an effort to identify oncogenic drivers, with EGFR being the most frequent. Several years ago we thought it should not be simply a research tool. We believed it should be integrated into clinical practice because

being applicable strictly as a research tool requires a protocol and can therefore only be utilized in 15% to 20% of the population.

We believe that every patient with lung cancer who comes into the clinic should undergo genotyping. The SNaPshot platform has now been supplemented by FISH analysis, primarily for the ALK-translocated gene and, more recently, ROS1 rearrangement. We also test everyone who comes to the clinic for MET and other less common rearrangements and mutations.

This model proved to be successful in clinical practice (Sequist 2011; [3.1]). We obtain a lot of data, many of which are unexpected — you would not have predicted these patients to have tumors with these types of mutations. The model has now spread to other disease centers, where breast cancer and colorectal cancer are screened.

3.1 Multiplex Genotyping of Non-Small Cell Lung Cancer (NSCLC) in Clinical Practice

"While widely agreed that it is important to identify patients with EGFR and ALK given the availability of effective therapeutics, it is also noteworthy that in a short time frame at a single institution, we identified over 30 patients with less common mutations like BRAF, PIK3CA and HER2, which also have relevant candidate targeted therapies.

Among the patients with advanced or recurrent NSCLC seen within these 15 months, 22% began a genotype-specific therapy in response to SNaPshot results. We anticipate that this proportion should increase further in the future, as the scope of genotype-specific clinical trial efforts is rapidly broadening.... Overall, we have demonstrated that broad clinical genotyping with SNaPshot can be tightly integrated into clinical practice and we believe it can make a real difference for patients."

Sequist LV et al. Ann Oncol 2011;22(12):2616-24.

Track 11

DR LOVE: Your group reported on a variation of the ECOG-E4599 regimen with carboplatin/*nab* paclitaxel and bevacizumab. What did you see, and do you have any sense of whether the regimens differ?

DR FIDIAS: In general, no overall survival benefit is evident with *nab* paclitaxel compared to paclitaxel, but we have recorded the highest response rate to date with any regimen (Heist 2011; [3.2]), although this one is more intense and causes more myelo-

3.2 Response to Carbopl and Bevacizumab Nonsquame	atin, Nanoparticle Albumin-Bound (<i>Nab</i>) Paclitaxel in Patients with Previously Untreated Advanced ous Non-Small Cell Lung Cancer (n = 25)
Disease control rate	17/23 (74%)
Partial response	8/23 (35%)
Stable disease	9/23 (39%)
One patient is in cycle 1, and 1 dropp study prior to first restaging: 1 for pa	bed out without receiving any study drug. Four patients came off inful bony disease requiring radiation therapy and 3 for toxicity

(perforated diverticulitis, liver function abnormalities and nausea and vomiting).

Heist RS et al. Proc ASCO 2012; Abstract e18016.

suppression and fatigue. A motivated person can get through it, but it's too early to tell whether *nab* paclitaxel should replace regular paclitaxel.

📊 Track 15

DR LOVE: Would you discuss the SELECT study, which you worked on with Dr Lecia Sequist? What did you report, and how do you interpret those results?

DR FIDIAS: This is an adjuvant trial for patients with EGFR mutation-positive disease, up to Stage IIIA, who can receive erlotinib either immediately if they don't receive chemotherapy or at the end of chemotherapy if the standard treatment for their tumor stage is to receive adjuvant chemotherapy after resection. It's a single-arm Phase II study, so it will not be conclusive, but the disease-free survival rate appears to be remarkably good at 94% after 2 years (Neal 2012; [3.3]).

It's still early for this population, and we have to see how the results fall along historical lines, but I would not be surprised if this regimen became a new standard. My sense is that EGFR TKI therapy will be moved earlier in the treatment algorithm. It's effective, and I can envision using TKIs in both the adjuvant and the metastatic settings. Eventually we'll need a Phase III study to find out whether we have a new standard. Until such time, I would not use this approach outside of a trial setting.

	Jeli Lung Callel
94% (95% Cl: 79.5%-98.5%)*	
Any (%)	Grade ≥3 (%)
89	17
78	3
61	6
oxicities leading to dose redu	ctions
	94% (95% CI: 7 Any (%) 89 78 61 öxicities leading to dose redu

SELECT PUBLICATIONS

Heist RS et al. Phase II trial of carboplatin, Abraxane, and bevacizumab in NSCLC. *Proc ASCO* 2011;Abstract e18016.

Neal JW et al. The SELECT study: A multicenter phase II trial of adjuvant erlotinib in resected epidermal growth factor receptor (EGFR) mutation-positive non-small cell lung cancer (NSCLC). *Proc ASCO* 2012; Abstract 7010.

Oxnard GR et al. Maintained sensitivity to EGFR tyrosine kinase inhibitors in EGFR-mutant lung cancer recurring after adjuvant erlotinib or gefitinib. *Clin Cancer Res* 2011;17(19):6322-8.

Sequist LV et al. Implementing multiplexed genotyping of non-small-cell lung cancers into routine clinical practice. Ann Oncol 2011;22(12):2616-24.

Shao H et al. Improved response to *nab*-paclitaxel compared with Cremophor-solubilized paclitaxel is independent of secreted protein acidic and rich in cysteine expression in non-small cell lung cancer. *J Thorac Oncol* 2011;6(6):998-1005.

Socinski MA et al. Weekly *nab*-paclitaxel in combination with carboplatin versus solvent-based paclitaxel plus carboplatin as first-line therapy in patients with advanced non-small-cell lung cancer: Final results of a phase III trial. *J Clin Oncol* 2012;30(17):2055-62.

Su Z et al. A platform for rapid detection of multiple oncogenic mutations with relevance to targeted therapy in non-small-cell lung cancer. J Mol Diagn 2011;13(1):74-84.