Targeted Treatment of Non-Small Cell Lung Cancer

Current Algorithms and New Agents

CME Information

TARGET AUDIENCE

This activity is intended for hematologists, medical oncologists and other healthcare providers involved in the treatment of non-small cell lung cancer (NSCLC).

OVERVIEW OF ACTIVITY

Lung cancer is a devastating disease with broad-reaching impact on public health, as it accounts for 15% of all new cancer cases in the US and the most cancer-related deaths among both men and women. In the year 2015, it is estimated that 221,200 individuals will be diagnosed and 158,040 individuals will die from the disease. Despite the many advances over the past few decades related to surgery, radiation therapy and chemotherapy, death rates attributable to lung cancer have remained relatively unchanged. Today, however, there is renewed optimism that these trends have started to change as recent research advances have led to an explosion in lung cancer genetic and biologic knowledge among scientists and clinicians working in this area of cancer medicine. A major focus of recent lung cancer research has been the development — and subsequent approval — of a number of molecular-targeted agents and the identification of related biomarkers to help guide treatment selection for those individuals who harbor specific oncogenic alterations.

These video proceedings from a CME symposium held in conjunction with the 16th World Conference on Lung Cancer feature discussions with leading researchers regarding actual cases of patients with NSCLC and tumor driver mutations from the practices of general medical oncologists and related clinical research findings to address existing uncertainties and help keep clinicians up to date and informed on the targeted treatment of NSCLC.

LEARNING OBJECTIVES

- Discriminate among molecular determinants that may be used to refine NSCLC prognosis and/or predict therapeutic response to an individual treatment, and apply available clinical guidelines to appropriately select patients for biomarker assessment.
- Employ an understanding of personalized medicine to individualize the use of available EGFR inhibitors in the long-term management of EGFR mutation-positive NSCLC.

- Describe mechanisms of tumor resistance to EGFR tyrosine kinase inhibitors (TKIs), and identify investigational therapeutic opportunities to circumvent this process.
- Communicate the efficacy and safety of crizotinib, ceritinib and other emerging ALK inhibitors to appropriate patients with NSCLC, considering the predictive utility of ALK and ROS1 mutation testing.
- Consider available clinical data and investigator perspectives when caring for patients with EGFR- or ALK-positive NSCLC and brain metastases.
- Assess new oncogenic pathways mediating the growth of unique NSCLC tumor subsets, and recall emerging data with experimental agents exploiting these targets.
- Recognize the abilities and limitations of multiplex and next-generation sequencing platforms, and determine their clinical and/or research application for patients with NSCLC.
- Appreciate the scientific rationale for ongoing investigation of novel agents or therapeutic approaches in NSCLC, and counsel appropriately selected patients about study participation.

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FACULTY — The following faculty (and their spouses/partners) reported real or apparent conflicts of interest, which have been resolved through a conflict of interest resolution process:

D Ross Camidge, MD, PhD

Director, Thoracic Oncology Clinical Program Associate Director for Clinical Research University of Colorado Cancer Center Aurora, Colorado

Consulting Agreements: Array BioPharma Inc, AstraZeneca Pharmaceuticals LP, Biodesix Inc, Genentech BioOncology, ImmunoGen Inc, Lilly, Novartis Pharmaceuticals Corporation, Roche Laboratories Inc.

Pasi A Jänne, MD, PhD

Director, Lowe Center for Thoracic Oncology Dana-Farber Cancer Institute Professor of Medicine Harvard Medical School Scientific Director Belfer Institute for Applied Cancer Science Boston, Massachusetts

Consulting Agreements: AstraZeneca Pharmaceuticals LP, Boehringer Ingelheim Pharmaceuticals Inc, Chugai Pharmaceutical Co Ltd, Clovis Oncology, Genentech BioOncology, Merrimack Pharmaceuticals Inc, Pfizer Inc, Roche Laboratories Inc, Sanofi; **Research Funding:** Astellas Pharma Global Development Inc, AstraZeneca Pharmaceuticals LP; **Stock Ownership:** Gatekeeper Pharmaceuticals Inc; **Other:** LabCorp.

Mark G Kris, MD

William and Joy Ruane Chair in Thoracic Oncology Attending Physician Thoracic Oncology Service Memorial Sloan Kettering Cancer Center New York, New York

Advisory Committee: Daiichi Sankyo Inc; Consulting Agreements: Array BioPharma Inc, AstraZeneca Pharmaceuticals LP, Clovis Oncology; Contracted Research: Pfizer Inc, Puma Biotechnology Inc; Other Remunerated Activities: Roche Laboratories Inc.

Tony SK Mok, MD

Professor, Department of Clinical Oncology The Chinese University of Hong Kong Hong Kong, China Advisory Committee: Amgen Inc, AstraZeneca Pharmaceuticals LP, Boehringer Ingelheim Pharmaceuticals Inc, Genentech BioOncology, GlaxoSmithKline, Lilly, Merck, Novartis Pharmaceuticals Corporation, Pfizer Inc, Roche Laboratories Inc; **Speakers Bureau:** AstraZeneca Pharmaceuticals LP, Boehringer Ingelheim Pharmaceuticals Inc, Lilly, Merck, Pfizer Inc, Roche Laboratories Inc.

CONSULTING ONCOLOGISTS — The following consulting oncologists (and their spouses/partners) reported real or apparent conflicts of interest, which have been resolved through a conflict of interest resolution process:

Paul Fishkin, MD

Illinois CancerCare, PC Peoria, Illinois

William Harwin, MD

Florida Cancer Specialists Fort Myers, Florida

Elizabeth D Simmons, MD Los Angeles, California

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Hardware/Software Requirements:

A high-speed Internet connection A monitor set to 1280 x 1024 pixels or more Internet Explorer 7 or later, Firefox 3.0 or later, Chrome, Safari 3.0 or later Adobe Flash Player 10.2 plug-in or later Adobe Acrobat Reader (Optional) Sound card and speakers for audio **Last review date:** November 2015

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Mark G Kris, MD

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