

Accepted Manuscript

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PII: S1556-0864(18)30004-2

DOI: [10.1016/j.jtho.2018.01.003](https://doi.org/10.1016/j.jtho.2018.01.003)

Reference: JTHO 836

To appear in: *Journal of Thoracic Oncology*

Received Date: 12 September 2017

Revised Date: 1 November 2017

Accepted Date: 5 January 2018

Please cite this article as: Akula S, Kamasani S, Sivan S, Manga V, Vudem DR, Kancha RK, Computational analysis of epidermal growth factor receptor mutations predicts differential drug sensitivity profiles towards kinase inhibitors, *Journal of Thoracic Oncology* (2018), doi: 10.1016/j.jtho.2018.01.003.

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Brief Report

Computational analysis of epidermal growth factor receptor mutations predicts differential drug sensitivity profiles towards kinase inhibitors

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Conflicts of interest

SA-None, SK-None, SKS-None, VM-None, DRV-None and RKK-None

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