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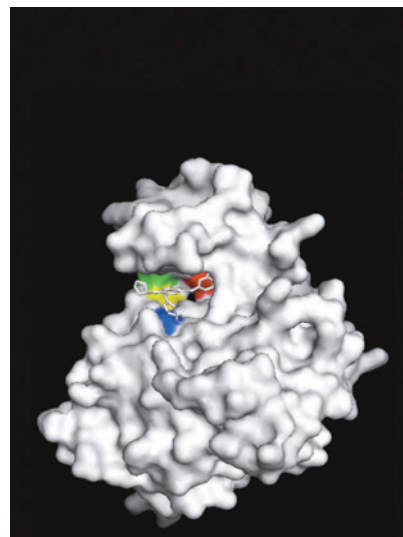
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ABOUT THE COVER

The cover shows a crystal structure of EGFR protein with afatinib, 2nd generation EGFR-tyrosine kinase inhibitor. Amino acid residues of the novel L792F and C797S, in addition to the conventional T790M, mutations as the mechanisms of acquired resistance against afatinib are colored. For details, see the article by Kobayashi and colleagues on page 357 of this issue.



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