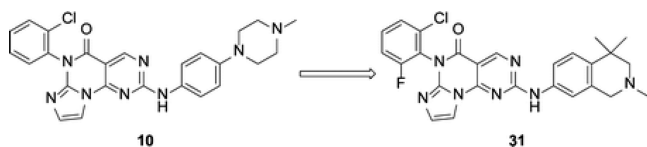


ORAL INHIBITORS OF WEE1 KINASE

Cells utilize checkpoints to ensure genomic integrity. Wee1 kinase is a key player in the process as it enables repair of damaged DNA in G2 phase prior to mitosis. As such, Wee1 inhibitors have been employed in preventing the activation of the G2 cell cycle checkpoint to enhance the antitumor activity of DNA damaging agents.

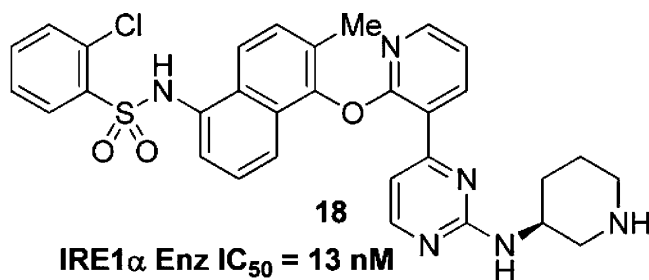
The research featured on the cover of this Special Issue discloses a novel class of potent and efficacious Wee1 inhibitors. Tong et al. (DOI: 10.1021/ml5002745) present the SAR studies and in vivo biomarker and efficacy data of a novel class of inhibitors. The SAR studies could give further insight into the active site of the Wee1 kinase.



RETHINKING THE ROLE OF IRE1 α IN CANCER THERAPY

Target validation efforts in drug discovery are often hampered by the potency and selectivity of the tool compound used, with compounds interacting with other cellular targets resulting in misperceived readouts.

In this Special Issue, Harrington et al. (DOI: 10.1021/ml500315b) address the previously reported potential of the enzyme IRE1 α as an anticancer target. The group identified potent and selective molecules that inhibit the endonuclease function of IRE1 α in cells and demonstrate surprising results showing that the enzyme inhibition has minimal impact on the tumor cell lines. These results may suggest a revision of the reported value of IRE1 α inhibition in cancer therapy.



IRE1 α Enz IC₅₀ = 13 nM

XBP1-Luc IC₅₀ = 99 nM

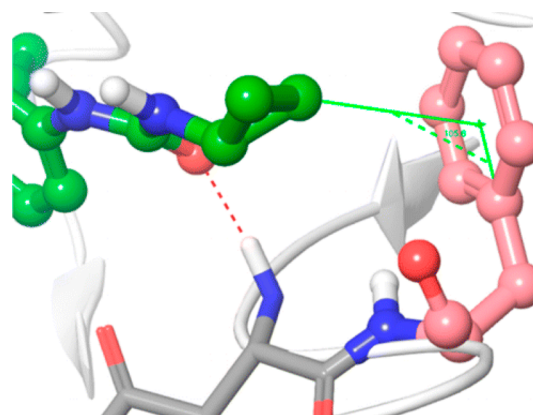
No Impairment of Tumor Cell Viability

BINDING MODE OF POTENTIAL ANTICANCER DRUG LENVATINIB

Lenvatinib is an oral, multitargeted tyrosine kinase inhibitor that is currently under clinical investigation in solid tumors. Lenvatinib selectively inhibits vascular endothelial growth factor receptor 2 (VEGFR2) and is being developed as an anti-cancer drug. Lenvatinib recently achieved the primary end point

in a phase 3 clinical trial in patients with differentiated thyroid cancer.

Here, Okamoto et al. (DOI: 10.1021/ml500394m) discuss the analysis of the interaction kinetics between lenvatinib and VEGFR2, which revealed the rapid association rate constant and a relatively slow dissociation rate constant of lenvatinib in complex with VEGFR2.



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