



**Fall 2025**

**Biochemistry and Molecular Biology  
Brown Bag Series**

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*A missense mutation in Rad3<sup>ATR</sup> bypasses Rad9-  
Rad1-Hus1 (9-1-1) phosphorylation to activate  
Cds1<sup>CHK2</sup> under replication stress in  
Schizosaccharomyces pombe (Fission yeast)*

**Tuesday, September 30, 2025**

**11:00 AM**

**Location 135 Oelman Hall**

**Lab: Yong-jie Xu, Ph.D.**



Boonshoft  
**School of Medicine**  
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## Abstract

Under replication stress, DNA replication forks can slow, pause, or even collapse, which activates DNA replication checkpoint (DRC) and DNA damage checkpoint (DDC) pathways for maintenance of genome integrity and cell survival. Mutations of the checkpoint pathways cause genome instability, a hallmark of cancer. In *S. pombe*, Rad3 kinase (the ATR homolog) together with its binding partner Rad26 monitors replication stress and activate the replication checkpoint by phosphorylating several substrates, including Mrc1-Thr645/653, Rad9-Thr412, and Cds1-Thr11. The phosphorylation defective mutations of *mrc1* and *rad9* impair Cds1 phosphorylation, thus sensitizing the cells to hydroxyurea (HU), and methyl methane sulfonate (MMS), by inhibiting DNA replication and causing DNA damage respectively. Although the role of phosphorylated Mrc1 in promoting Cds1 activation is well established, the function of phosphorylated Rad9 in this context has remained less clear. To address this problem, we randomly mutagenized the genome in the *rad9*-Thr412Ala mutant and screened a suppressor mutant that confers HU resistance in *rad9*-T412A, as well as in *rad9*- $\Delta$ C ( $\Delta$ 411-426). Consistent with the drug resistance, phosphorylation of Cds1-Thr11 by Rad3 is restored. Whole genome sequencing identified a single missense mutation within *rad3*, and subsequent experiments demonstrated that overexpression or genomic integration of the mutant *rad3* allele confers HU resistance and restored Cds1-Thr11 phosphorylation in *rad9*- $\Delta$ C cells. In vitro kinase assays further revealed that the mutant Rad3 protein is constitutively active even in the absence of replication stress, and this hyperactive form of Rad3 can also rescue the HU sensitivity seen in *rad26* mutants with N-terminal mutations. These results collectively demonstrate that a *rad3* constitutive mutation can bypass the canonical requirement for 9-1-1 complex phosphorylation, thereby directly activating Cds1 and stabilizing replication forks under stress conditions.