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Figure S1 Gene-gene network analysis showing E2F1 correlated genes. (**A**) Gene-gene interaction network associated with DNA repair genes. Each colored line represents functional and physical interactions in the literature, respectively. These networks were generated by GeneMANIA (http://genemania.org) (**B**) GeneMANIA function analysis and significant enrichment of selected genes in bladder cancer patients.

DR-GFP vector

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Figure S2 Schematic diagram of HR assay by DR-GFP. (A) DR-GFP vector contains an I-SceI endonuclease site within the coding region, which abolishes GFP expression. Expression of I-SceI induces single DSBs.

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Figure S3 Schematic diagram of bioinformatics. (**A**) RLEL, RAD54L low and E2F1 low; RLEH, RAD54L low and E2F1 high; RHEL, RAD54L high and E2F1 low; RHEH, RAD54L high and E2F1 high.



Figure S4 Hierarchical clustering analysis of gene expression data from GSE13507 for patients with NMIBC. (**A**) Heat map of RAD54L correlated 1,027 genes in 102 NMIBC patients. Two subtypes of patients were revealed from unsupervised clustering analysis using RAD54L correlated 1,027 genes. (**B**) Depending on the expression of E2F1, two subtypes are redefined into four 4 new subtypes.





Figure S5 E2F1 and RAD54L expression levels are not correlated with recurrence in BLC patients. (A) IHC staining with anti-E2F1, RAD54L antibody was performed for non-recurrent BLC tumor tissues and recurrent BLC tumor tissues. Photographs were taken at a magnification of ×. (B) Recurrence-free survival in NMIBC patients (GSE13507, n = 102, p = 0.05 by log-rank test, left panel). Recurrence-free survival in NMIBC patients (GSE13507, n = 102, p = 0.18 by log-rank test, right panel). RLEL, RAD54L low and E2F1 low; RLEH, RAD54L low and E2F1 high; RHEL, RAD54L high and E2F1 low; RHEH, RAD54L high and E2F1 high.