Selective Targeting of CDK2 Using Molecular Glue Degraders for the Treatment of **HR-Positive/HER2-Negative Breast Cancer**

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CDK2 phosphorylates RB, freeing E2F transcription factors to modulate gene expression that drives the G1/S progression. CDKs 4 and 6 also impact the G1/S checkpoint and can be targeted with conventional kinase inhibitors in the clinic.





(A) Novel binding mode of MRT-9643, which does not engage a G-loop or the catalytic site. CDK2-MGD-CRBN-DDB1 cryo-EM structure (DDB1 not shown). (B-C) MRT-9643 is a potent CDK2-directed MGD. (B) Western blot of CDK2 degradation in MDA-MB-157 cells, with increasing concentration of MRT-9643. (C) Degradation of CDK2 in multiple HR-positive breast cell lines, assessed by Western blot. (D) MRT-9643 induces potent and highly selective CDK2 degradation. Global protein expression determined with TMT proteomics at 24 hours, with 1µM MRT-9643 treatment in MCF7 cells. Other cyclin-dependent are highlighted in blue.

MRT-9643 Inhibits Proliferation of CDK2-Dependent Cancer Cells Protein fold-change (log₂ G1 phase Proliferation **S** phase - CDK2 levels (WB G2/M phase E2F Targets

(A) Effect of MRT-9643 on cellular proliferation and CDK2 protein level. Proliferation was assessed by CyQuant Direct after 7 days of treatment in MDA-MB-157 cells. Error bars show SD, N=3 CDK2 protein was determined by western blot. (B) CDK2 degradation by MRT-9643 arrests CDK2-dependent cells in G1 phase. Cell cycle phases determined with Click-EdU at 24 hours in MDA-MB-157 cells. Bars show mean, N=3. (C) CDK2 degradation results in reduction of E2F pathway proteins. Global protein expression determined with TMT proteomics at 24 hours, with 1µM MRT-9643 treatment in MDA-MB-157 cells. Transcriptional E2F targets are highlighted blue.



(A) Clinical-stage CDK2 inhibitors show off-target activity in biochemical kinome **profiling.** Inhibition was determined by Carna Biosciences' mobility shift assay, 1µM CDK2 inhibitor or CDK2 MGD, with 323 human kinases assessed. (B-C) CDK2 inhibitors, but not a CDK2 MGD, display CDK2-independent activity. Assessment of growth inhibition of CDK2 MGD MRT-9643 and CDK2 inhibitors in (B) MDA-MB-157 cells with wild type or knocked out RB and (C) MCF7 cells with wild type or knocked out CDK2. Proliferation was assessed by CyQuant Direct after 7 days of treatment. N=3, with error bars showing SD.

MRT-9643 Displays Superior Selectivity Compared to Clinical CDK2 Inhibitors





Growth of MCF7 cells treated with MRT-9643 (1µM), ribociclib (1µM) or combination, and % confluence was measured using Incucyte confluence monitoring. Error bars represent SD, N=2. (B) Combination of MRT-9643 and ribociclib enhances reduction of CDK2 and pCDK2. Western blot of CDK2 and pCDK2 [T160] degradation in T47D cells, with or without 1µM ribociclib treatment. (C-D) Combination of MRT-9643 and ribociclib induces robust tumor regression of T47D xenografts. (C) T47D cells (10x106/mouse) were grown subcutaneously in Balb/c nude mice. Tumors of ~200mm³ were randomized and treated with vehicle, MRT-9643 (30mg/kg PO BID), ribociclib (75mg/kg PO QD), fulvestrant (5mg/mouse SC QD), or combination. (D) Waterfall plot of individual tumor growth, relative to vehicle and starting tumor volume.

- catalytic site or G-loop
- Degradation of CDK2 inhibits CDK2-dependent cancer cell proliferation while displaying superior selectivity compared to clinical CDK2 inhibitors CDK2 degradation delays resistance to CDK4/6 inhibition *in vitro* and exhibits activity in combination with CDK4/6 inhibitors *in vivo* • CDK2 degradation in combination with CDK4/6 inhibition achieves equivalent efficacy compared to anti-CDK4/6 and anti-estrogen standard of care

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Tumors of ~200mm³ were randomized and treated with vehicle, MRT-9643 (30mg/kg PO BID) ribociclib (75mg/kg PO QD), fulvestrant (5mg/mouse SC QD), or combination. (B) Waterfall plot of individual tumor growth, relative to vehicle and starting tumor volume. (C-D) MRT-9643 in combination with HR-positive standard of care treatments induce significant pathway suppression. (C) Relative CDK2 and p-RB [S807/811] protein after indicated treatments in representative xenograft tumors. Samples harvested on Day 28, N=3 tumors per group, error bars represent SEM. (D) RNAseq analysis of representative MCF7 xenograft tumors. Samples harvested on Day 28, N=3 tumors per group. Gene expression is shown as fold change relative to vehicle. E2F genes are colored in purple.

Conclusions

Want to see what else molecular glues can do?



• The CDK2-directed molecular glue degrader MRT-9643 exhibits potency, selectivity, and favorable drug-like properties – without engaging the