

# The *POLD1*<sup>R689W</sup> variant increases the sensitivity of colorectal cancer cells to ATR and CHK1 inhibitors

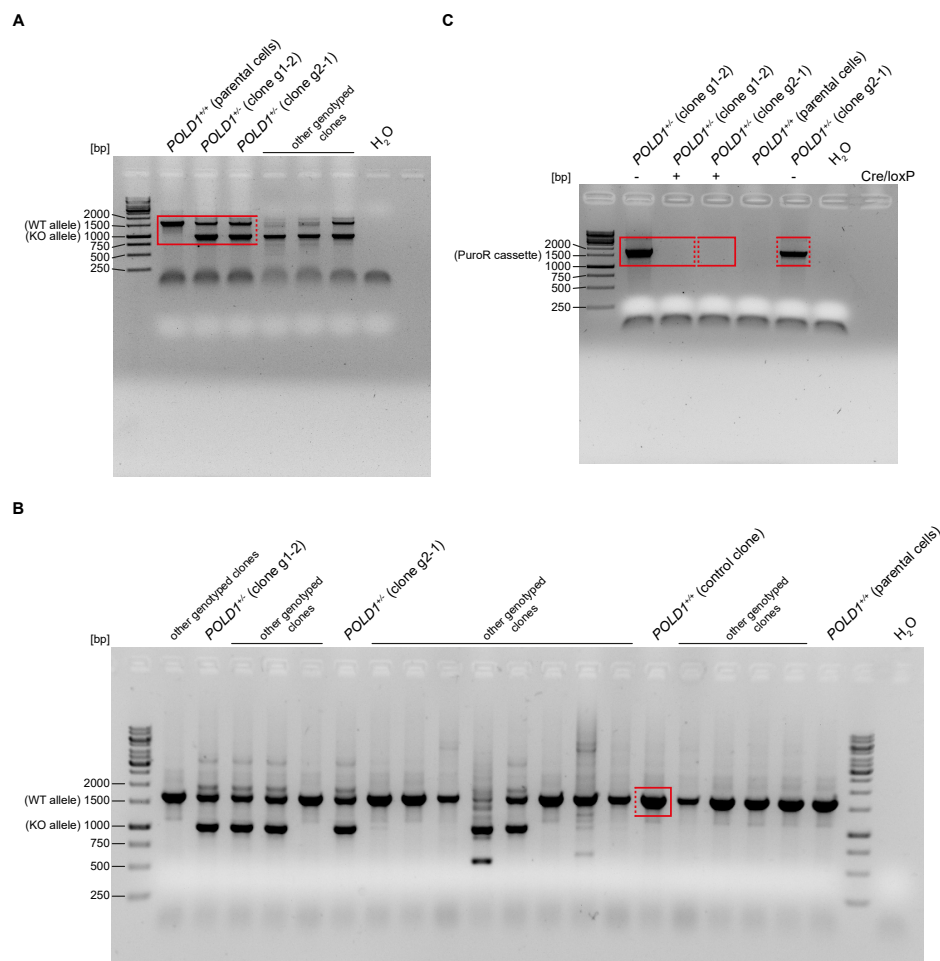
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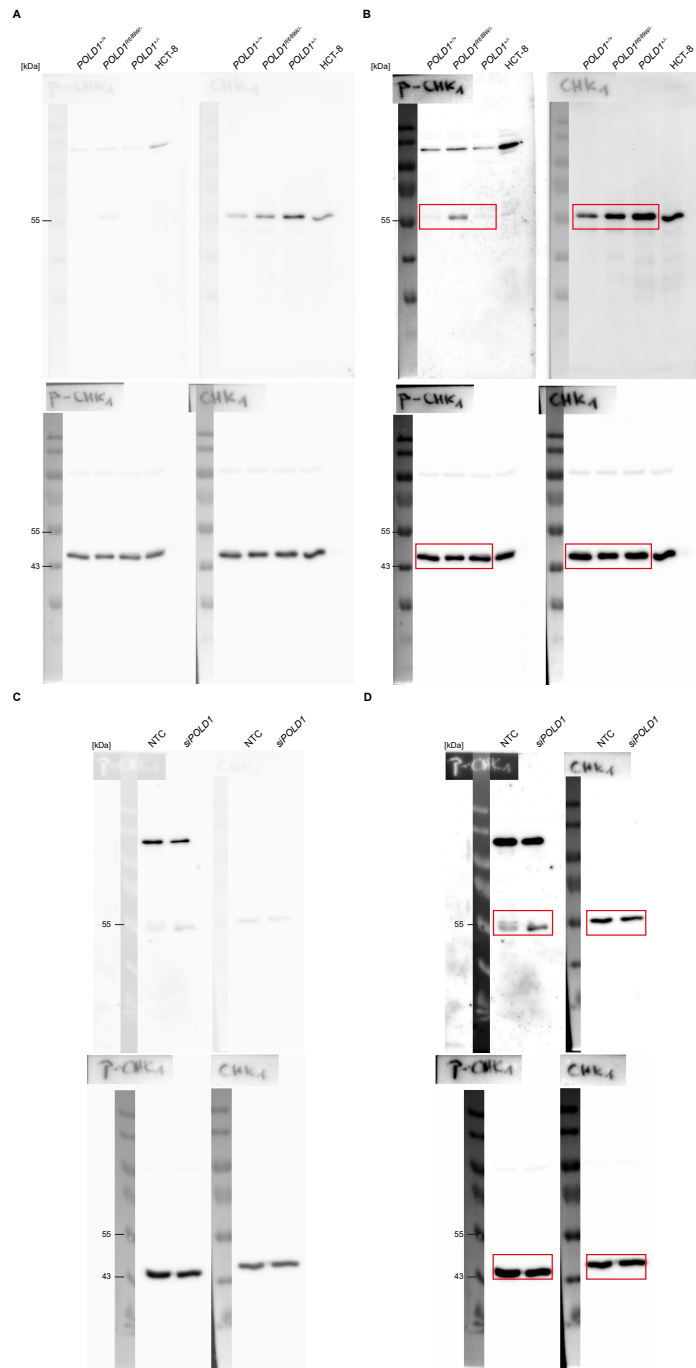
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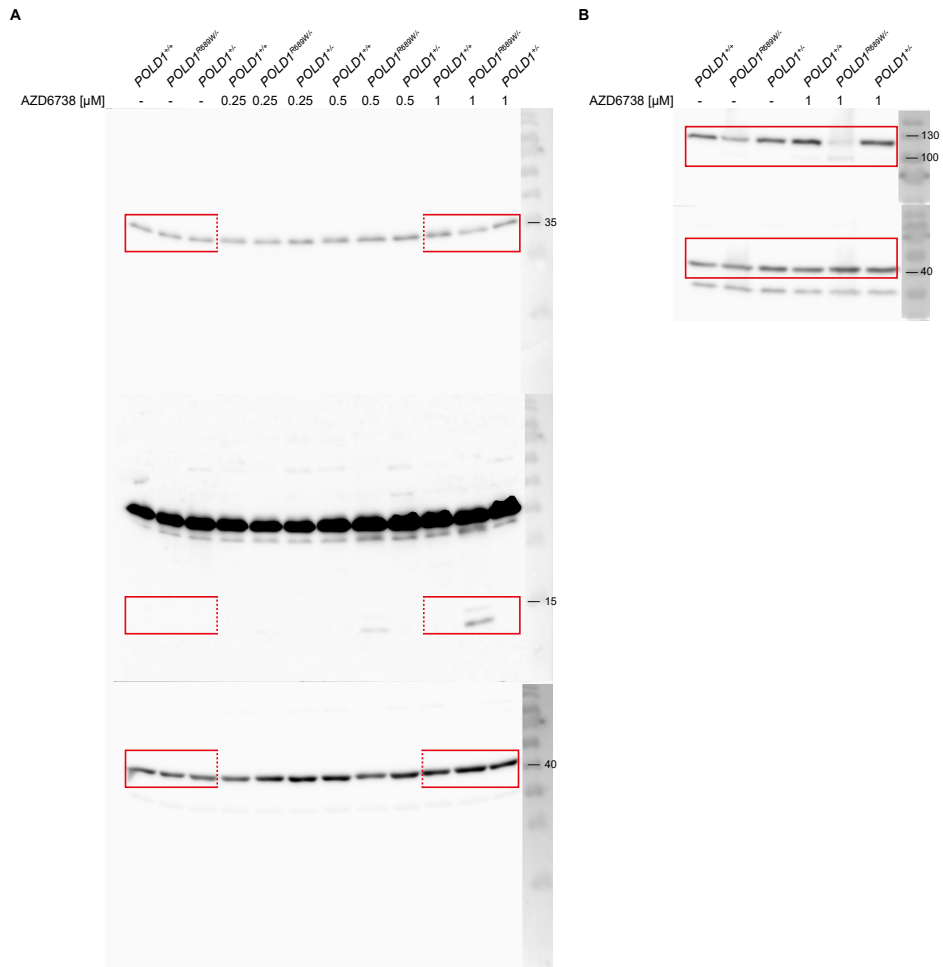
## Supplementary Information



**Figure S1.** (A+B) Original gels of the PCR detecting the *POLD1*-KO and -WT alleles in the *POLD1*<sup>+/+</sup> parental and control clones as well as in the *POLD1*<sup>+/+</sup> clones g1-2 and g2-1. Data which were cropped and depicted in Figure 1B are framed red. (C) Original gels of the PCR detecting the Cre/loxP-mediated excision of the puromycin resistance cassette. Data which were cropped and depicted in Figure 1C are framed red.



**Figure S2.** (A, upper panel) Original immunoblots and (B, upper panel) immunoblots after contrast modification of constitutive protein levels of pCHK1 and CHK1 in *POLD1*<sup>+/+</sup>, *POLD1*<sup>R689W/+</sup> and *POLD1*<sup>+/-</sup> cells. (A+B, lower panels) Original immunoblots and immunoblots after contrast modification, respectively, of β-Actin serving as loading control. (C, upper panel) Original immunoblots and (D, upper panel) immunoblots after contrast modification of protein levels of pCHK1 and CHK1 120 hours after *siPOLD1* transfection in *POLD1*<sup>+/+</sup> cells. (C+D, lower panels) Original immunoblots and immunoblots after contrast modification, respectively, of β-Actin serving as loading control. Data which were cropped and depicted in Figure 3A are framed red.



**Figure S3.** Original immunoblots of protein levels of (A, upper part) caspase 3, (A, middle part) cleaved caspase 3 as well as (B, upper part) PARP and cleaved PARP in *POLD1*<sup>+/+</sup>, *POLD1*<sup>R689W/+</sup> and *POLD1*<sup>+/+</sup> cells 72 hours after treatment with AZD6738. (A+B, lower parts) Original immunoblots of β-Actin serving as loading control. Data which were cropped and depicted in Figure 4C are framed red.



**Figure S4.** Maps of the pCas9 and repair template plasmids.

**Table S1. Primer used in PCR**

<b>Primer</b>	<b>Orientation</b>	<b>Localization</b>	<b>Sequence (5' - 3')</b>
# 1	sense	POLD1 NCS <sup>a</sup> , upstream LHA	GTGAGAGAGCACACACACGAC
# 2	antisense	repair template, GFP	TAGGTGCCGAAGTGGTAGAAGC
# 3	sense	repair template, spacer region	TCTCTTGATTCCCACCTTTGTGGT
# 4	antisense	POLD1 NCS, downstream RHA	CAGATCAACGCTCCAAGCAC
# 5	sense	POLD1, LHA	GAGGTGTCTCCGGTCAGAAC
# 6	antisense	repair template, puromycin resistance	GAGGCCTTCCATCTGTTGCT
# 7	sense	POLD1 NCS, upstream exon 13	CCCAGACCCTGACGACTTGG
# 8	antisense	POLD1 NCS, downstream exon 13	TGGGAGTGGGGAGAAAAAGTG
# 9	sense	POLD1 NCS, upstream exon 17	TGCGTGAATTAGCACAAGGC
# 10	antisense	POLD1 NCS, downstream exon 17	GGACCAATTGCTCAAGCCAC
# 11	sense	POLD1 NCS, upstream exon 18	TCCGCATGATTCTCTCCCCG
# 12	antisense	POLD1 NCS, downstream exon 18	GTGGCTAATGCCAACGGGAC
# 13	sense	POLD1, exon 2a	GGGCCTCTGGGATGATGATG
# 14	antisense	POLD1, exon 21	CTGGGAGATATCGATGCGGT
# 15	sense	POLD1, exon 10	GTATCATGGACCCCGACGTG
# 16	antisense	POLD1, exon 14/15b	CGTAGTACCCTTTGAGGGGC
# 17	sense	POLD1, exon 15b	CTGTGTTACACCACGCTCCT
# 18	antisense	POLD1, exon 20	GTGAGGCAGTGACCAGGTTG

<sup>a</sup> NCS: non coding sequence

**Table S2. Primer used for sequencing**

<b>Primer</b>	<b>Localization</b>	<b>Sequence (5' - 3')</b>
# S1	upstream exon 2a, NCS <sup>a</sup>	TCAGAACCTCCACCAAG
# S2	upstream exon 13, NCS	ACTTCCTTCTCCTGCTC
# S3	upstream exon 17, NCS	TGTGCAGTGCACAGTAC
# S4	upstream exon 18, NCS	GTTCGGACGTCAGATGATC
# S5	upstream exon 13	CTCCTACACGCTCAATG
# S6	upstream exon 17	AGATCCTGGAGAACCTG

<sup>a</sup> NCS: non coding sequence