

## **Appendix to Biedermann et al.**

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## Appendix Tables

**Appendix Table S1 – Significance table for Fig. 4I**

	WT -B -R	WT +B -R	WT -B +R	WT +B +R	rbr1 -B -R	rbr1 +B -R	rbr1 -B +R	rbr1 +B +R
WT -B -R		*	-	*	-	**	-	**
WT +B -R			*	-	-	*	-	*
WT -B +R				*	-	***	-	***
WT +B +R					*	-	*	-
rbr1 -B -R						**	-	**
rbr1 +B -R							**	-
rbr1 -B +R								**
rbr1 +B +R								

- No significant differences

\* P value < 0.05

\*\* P value < 0.01

\*\*\* P value < 0.001

**Appendix Table S2 – Primer sequences**

### Genotyping

Wildtype	Designation	Sequence
<i>CDKA;1</i>	A03	CAGATCTCTCCTGGTTATTCA
	A04	TGTACAAGCGAATAAAGACATTGA
<i>E2FA</i>	E2Fa FW	ATGTCGGTGTGTCGTACGATCTC
	E2Fa RW	TCATCTGGGTTGAGTCAAC
<i>KU70</i>	A17	AACCCTTACTTAGATATGATTAC
	A18	AGGGTGTTATTCCGAGGCTTACT
<i>RAD51</i>	A302	GGTTCCATCACAGAGTTATATGG

	A303	AGCCATGATATTCCCACCAATC
<i>RBR1</i>	M206	CTTCCACAGCCCGGTCGTTTC
	M207	GATTACCGCAGCATTCTAGTTGAACGC
<i>WEE1</i>	A06	TCAATAAGGCTGGTTCTTCAGT
	A07	AGGCATGTAACGTGCATCTC

T-DNA insertion allele	Designation	Sequence
<i>cdka;1</i>	A01	GCGTGGACCGCTTGCTGCAACTCTCTCAGG
	A02	CCAGATTCTCCGTGGAATTGCG
<i>e2fa-2</i>	GABI-Kat LB	ATATTGACCATCATACTCATTGC
	E2Fa RW	TCATCTGGGGTTGAGTCAAC
<i>ku70</i>	A18	AGGGTGTATTCCGAGGCTTACT
	LBd1	TCGGAACCACCATCAAACAG
<i>rad51-1</i>	A302	GGTTCCATCACAGAGTTATATGG
	GABI-Kat LB	ATATTGACCATCATACTCATTGC
<i>rbr1-2</i>	M206	CTTCCACAGCCCGGTCGTTTC
	J504	GCGTGGACCGCTTGCTGCAACTCTCTCAGG
<i>wee1-1</i>	A05	CCCATTGGACGTGAATGTAGACAC
	A06	TCAATAAGGCTGGTTCTTCAGT

## qPCR

Target	Forward primer	Reverse primer
<i>AHP2</i>	GCGCAAGCGTAAGAGGGATGTTC	CCAAGCTCCTCCTTAGTTCCCTC
<i>ATM</i>	TTGGTCTTGGTGACCGACATGC	TCTTGTCACTCTGAACGGAACCC
<i>ATR</i>	GTGCCATTGAGATTGACCCAGAAC	TGCCCTCATATCCAGTGATGCC
<i>BRCA1</i>	TGCATCCATTAAGTTGCCCTGTG	TAGGCTGAGAGTGCAGTGGTTC

<i>FAN1</i>	AGTGAAGGACGCAAATGAACCTG	TGAGGTGCGACTGAACACCTTG
<i>KU70</i>	AGCAGTCGATTATGGCGATGACC	CCACAGTCAAGTCCTCAGCTTC
<i>LIG4</i>	TTGGCTTCAAGTGAGAACAGAGC	TGACCCACTTCATCTCCTGAGC
<i>MLH1</i>	ACGACGGTCACGGTATTAGACG	AGCTTCGATGTTATGTCTCTCG
<i>MRE11</i>	AACAAATCTCAGCCTCGGGTTAC	AGAAGTTGTTCCGCTTGAGAGGTC
<i>MSH4</i>	TGAGAACAGAGCCAAGAGGTTGG	AGATGCAGTGAAGCCGATCTTAGG
<i>MSH5</i>	ACGAGAGCTGCTGCCTGTTTC	TCTGTGTCTGGCCTAAGAACGC
<i>MUS81</i>	TTGCCATTCACTGCTAATGCAGGTC	TCTTCTTGCGCCGAGACATCTG
<i>NBS1</i>	CTTCACTGATAACCACCATCCGTTG	GCTTCAGAACCGCTACCACTG
<i>PARP2</i>	ATGCTACTCTGGCACGGTTAC	AGGAGGAGCTATTGCAGACCTTG
<i>PCNA1</i>	CGGTGACATTGGAACCGCTAAC	TCACAATTGCATCTCCGGCTTG
<i>RAD51</i>	TTCCGCTCTGAAAGACTCAGC	ACCTCCTGATCCATGGAAAGTTG
<i>RECQL</i>	ATGCTTCGTGGTGGACAACTC	GCAGCCAAACCCATAACCTGATCC
<i>SOG1</i>	CCATGAGGTTCTTGCCGAGAC	TCAGGCCAGAACATTGGTCTTC
<i>TSO2</i>	TTTCGCTGCGTCGAAGGTATC	AGGCATGAGTCCTCGTTCTCAG

Reference genes	Forward primer	Reverse primer
At1g02410	ATCTGGTACCGTCACTGAAAGGG	TGCATCCCCTTGCAACATCAGC
At4g26410	TCGTCTTGCTAAAGTCCGTCCAC	GCCGAAGTCCGTCCATCAAATCAG
At4g30520	TTCCCAGCAAGGACTTCAACG	CAGAACAAATCTCAGGTGGTTGC
At5g36210	AGAGCGGTTGTTAAGGCAGTGG	TCTGCCTGCCAGAACGATACCAAG

## ChIP

Designation	Sequence
RAD51_+1209_F	ATGCTGAGGGAACATTCAAGG
RAD51_+1330_R	TCCAAACCTGCATCAATTCA
RAD51_+228_F	CGCGATGTGAAATAGGTGTG
RAD51_+323_R	ACCCCCAAAAAGTGTGATT

RAD51_-558_F	TGTTGGAATTGTGGTGGTC
RAD51_-489_R	TTTGACCGCCGAGTGATAC
RAD51_-1063_F	GGGTCCATAGCTCAGTGATAGAG
RAD51_-1002_R	GAAAACATATAAGGCCAACC
RAD51_E2F(-113)_F	GTTGTGGCGCTTCTTCAAT
RAD51_E2F(-17)_R	GATTCTGGTCACCAGAGAAA
ACT7(At5g09810)_F	GTTGCCATTAGGCCGTTCTTC
ACT7(At5g09810)_R	CAGAATCGAGCACAATACCGGTTG
MCM_F4	TCCCGCCAAACTCATAGTC
MCM_R4	TGACATCGTTGCTTCGTCTC
ORC3_F	CCAATTCCGGTCTAGTCTGG
ORC3_R	TGGAGCAATCGAAAACGACG

## Appendix Figure Legends

### Appendix Figure S1 – The *rbr1* mutant line is hypersensitive to BLM and cisplatin.

- **A** Root tip phenotypes of eight day-old seedlings germinated and grown on medium containing no supplement and transferred to medium containing BLM (6 $\mu$ g/ml) for two days.
- **B** Comparison of root growth of wild-type and *rbr1* plants on cisplatin (cisPt). Plants were germinated on MS medium containing no supplement and transferred to medium with 15 $\mu$ M cisPt three days after germination. Daily root growth was measured one, two and three days after transfer (dat). Error bars signify the standard deviation in three independent experiments. Two asterisks indicate significance higher than 99% as calculated by Student's T test.
- **C** Root tip phenotypes of eight day-old seedlings germinated and grown on medium containing no supplement and transferred to medium containing cisPt (15 $\mu$ M) for one or two days respectively.

### Appendix Figure S2 – Cell death in root tips upon drug treatment.

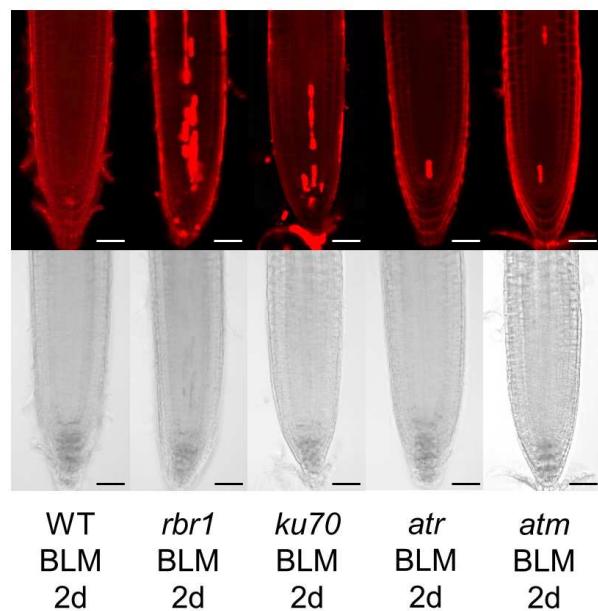
- Root tip phenotypes of ten day-old seedlings germinated and grown on medium containing HU (1mM) or no supplement (-). Upper rows show cell death visualized by propidium iodide staining, lower rows show brightfield microscopic images of root tips. Scale bars: 50 $\mu$ m.

### Appendix Figure S3 – Co-localization of RBR1 and RAD51 to $\gamma$ H2AX foci.

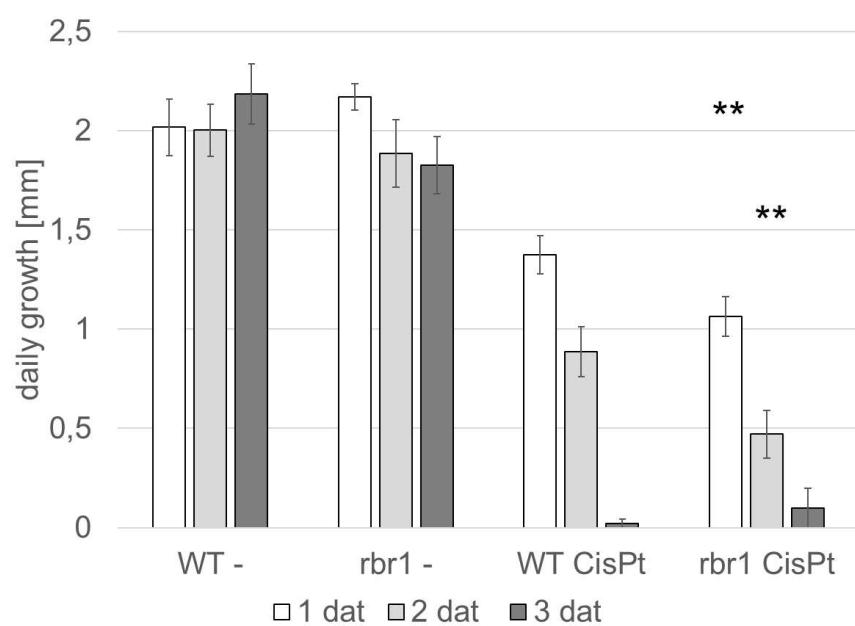
- Z-stack image of co-localization of RBR1 (green) and RAD51 (red) to  $\gamma$ H2AX foci (grey) in immuno-stained spreads of BLM-treated root tips of *PRO<sub>RBR1</sub>mCherry:RBR1 rbr1-3* plants, counter-stained with DAPI (DNA, blue).

## Appendix Figure S1

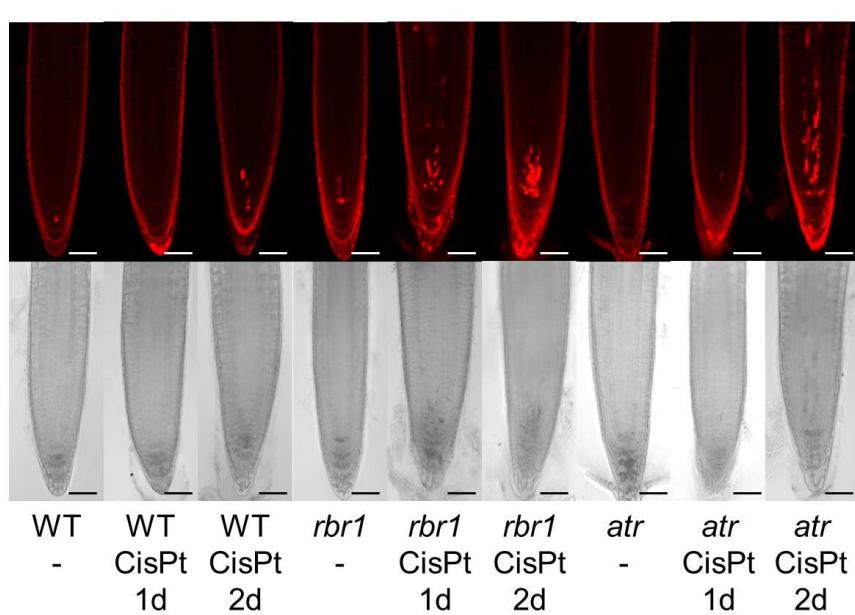
A



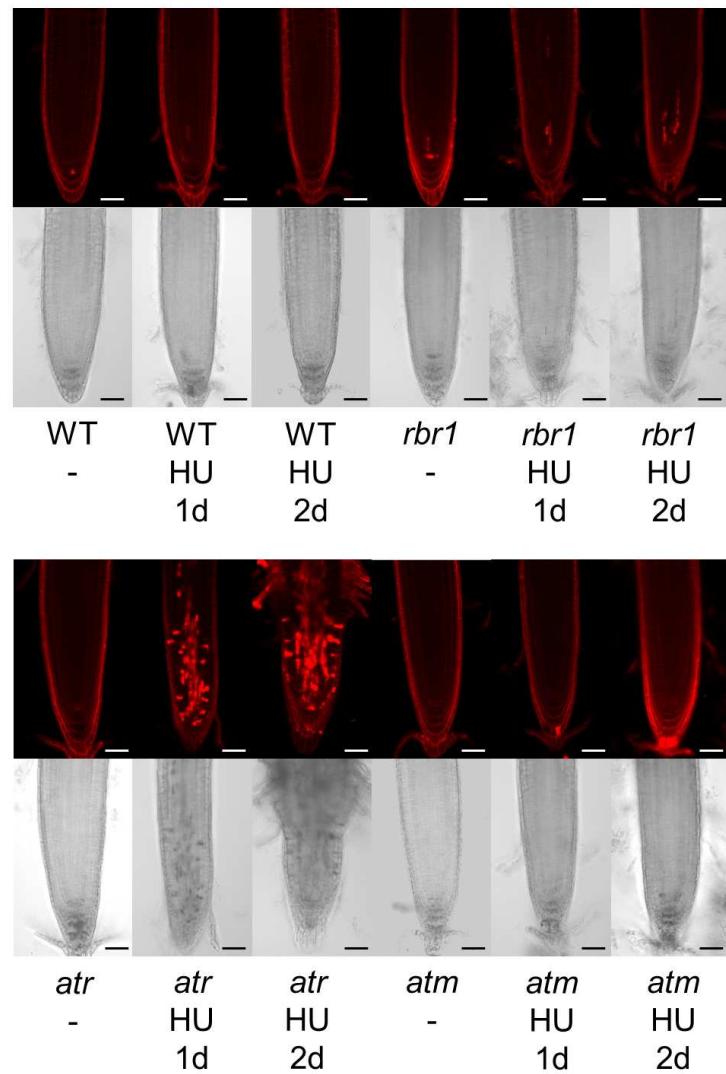
B



C



## Appendix Figure S2



**Appendix Figure S3**

