

1 **Supplementary Information**

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3 **Activating Akt1 mutations alter DNA double strand break repair and**
4 **radiosensitivity**

5

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1 **SI Methods**

2 **Reagents.** The inhibitors MK-2206, GDC-0068 and NU7441 were obtained from SelleckChem
3 (Houston, TX, USA). G418 was purchased from Biochrom AG (Berlin, Germany). All other chemicals
4 were from Sigma-Aldrich (St. Louis, MO, USA) and Thermo Fisher Technology (Waltham, MA, USA) if
5 not otherwise indicated.

6
7 **Antibodies.** The γ -H2A.X-AF647 antibody was obtained from Becton Dickinson (#560447, Franklin
8 Lakes, NJ, USA). Mouse monoclonal β -Actin antibody was obtained from Sigma-Aldrich (#A5441, St.
9 Louis, MO, USA). All other antibodies were purchased from Cell Signaling Technology (Danvers, MA,
10 USA): Akt (pan) #4685 (t-Akt), Phospho-Akt (Thr308) XP #13038 (p-T308), Phospho-Akt (Ser473) XP
11 #4060 (p-S473), GSK-3 β XP #12456 (t-GSK3), Phospho-GSK-3 β XP #5558 (p-GSK3).

12
13 **Cell Culture and Treatment.** TRAMP-C1 murine prostatic adenocarcinoma cells were purchased from
14 ATCC (Bethesda, MD, USA). Murine embryonic fibroblasts (MEF) from Akt wildtype (WT) and Akt1
15 knock-out (Akt1 $-/-$) mice were kindly provided by Morris J. Birnbaum (Philadelphia, PA, USA). Cells
16 were cultured in DMEM (Thermo Fisher Technology, Waltham, MA, USA) medium supplemented with
17 10% (v/v) foetal calf serum (Biochrom AG, Berlin, Germany) and maintained in a humidified incubator
18 (Labotect, Goettingen, Germany) at 37°C and 5% CO₂. All cell lines were validated and monitored
19 using microscopy and Western Blots analysis. All cell lines have monthly been tested for mycoplasma
20 contamination. Cells were transduced with viruses produced in Platinum-E cells (ATCC, Bethesda, MD,
21 USA) using the pBABE-retroviral system according to Morgenstern and Morita ^{45,46}. The transduced
22 cells were cultured two weeks before and after eGFP-based cell sorting in DMEM medium containing
23 10% FCS, 1x Pen/Strep, 1 μ g/ml Ciprofloxacin and 4 μ g/ml Puromycin for selection and prevention of
24 contamination.

25
26 **Irradiation.** For radiation treatment cells were exposed 24 h after seeding to 1-10 Gy using an X-RAD
27 320 X-Ray Biological Irradiator with X-RAD MIR-324 x-ray tube (Precision X-Ray Inc, North Branford,
28 CT, USA). Non-irradiated controls were handled in parallel but kept outside of the irradiator during
29 treatment. Cell number and viability was quantified by counting cells with a CASY cell counter from
30 Innovatis (Reutlingen, Germany).

31
32 **Western Blot Analysis.** Cell pellets were lysed for 30 min on ice in RIPA buffer (50 mM HEPES,
33 pH7.5, 1% Triton X-100, 150 mM NaCl, 10 mM NaPP, 1 mM EDTA, 10 mM NaF, 2 mM Na₃VO₄,
34 Protease Inhibitor Cocktail Tablets from Sigma-Aldrich (St. Louis, MO, USA) and lysed with ultrasound
35 (Bandelin Electronic, Berlin, Germany). Lysates were subsequently centrifuged 10 min at 14.000 rpm at
36 4°C and the protein concentration of the supernatants was adjusted to an equal protein amount using
37 the Bradford Protein Assay (BIO-RAD, Munich, Germany). The protein solutions were incubated with
38 4x sample buffer (300 mM TRIS-HCl pH 6.8, 2.4% SDS, 60% Glycerol, 30% β -Mercaptoethanol, 3.6

1 mg/100ml bromphenol blue) for 10 min at 95°C. For Western blot analysis proteins were separated by
2 SDS-PAGE and blotted onto PVDF-membranes (Roth, Karlsruhe, Germany). After blocking with 5%
3 (w/v) non-fat dry milk, membranes were incubated at 4°C overnight with the respective primary
4 antibody (1:20.000 for β -actin, 1:1000 for all other antibodies). Afterwards, the membranes were
5 incubated for 1 h at RT with the secondary antibody (anti-IgG-HRP 1:2000) diluted in 5% (w/v) non-fat
6 dry milk. Detection was performed by chemiluminescence imaging using ECL Prime Western blotting
7 detection reagent (GE Healthcare, Pittsburgh, PA, USA), a FUSION Solo and the Fusion software
8 (peqlab, Erlangen, Germany).

9

10 **Quantification of cell viability and cell proliferation.** The number of living cells was determined
11 using CASY technology based on an electric field multi-channel cell counting system from Innovatis AG
12 (Reutlingen, Germany). For this, cells were harvested by trypsinization, re-suspended in fresh medium,
13 diluted with CASYton solution from Hoffmann-La Roche AG (Basel, Switzerland) and measured.
14 Metabolic activity and cell viability was measured with the Wst-1 assay according to the manufacturer's
15 instructions (Roche Applied Science, Mannheim, Germany) by an ELISA reader (Bio-Tek, Bad
16 Friedrichshall, Germany).

17

18 **Flow cytometry.** Flow cytometric analysis of the cell cycle was based on propidium iodide (PI) staining
19 of the DNA in a hypotonic citrate buffer according to the Nicoletti protocol ⁴⁹. In brief, cells were
20 resuspended in 200 μ L of the staining solution (comprised of 0.05% Triton X-100, 50 μ g/mL propidium
21 iodide and 0.1% sodium citrate in PBS), incubated in the dark for 30 min and measured with a BD
22 Accuri C6 flow cytometer (BD Biosciences, San Jose, CA, USA) was performed. Flow cytometry
23 analyses were performed in BD Accuri C6 Software (BD Biosciences, San Jose, CA, USA).

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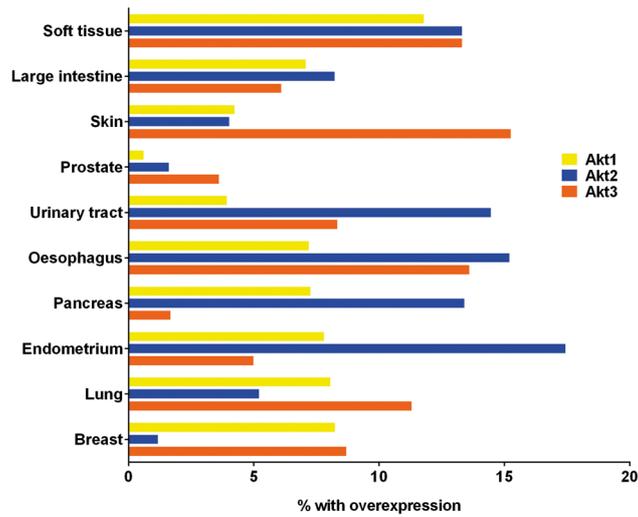
1 **Supplemental Tab. 1: Akt1 constructs and effects of different variants**

VARIANTS	TYPE	EFFECT
Akt1-WT	Wild type	-
Akt1-E17K	Dominant active	Enhanced PIP ₃ coordination
Akt1-R25C	Dominant negative	Impaired PIP ₃ coordination
Akt1-K179A	Dominant negative	Impaired ATP coordination
myrAkt1	Constitutively active	No PH-domain, N-terminal myristoylation
Akt1-T308D/S473D	Constitutively active	T308/S473 phosphorylation mimicked

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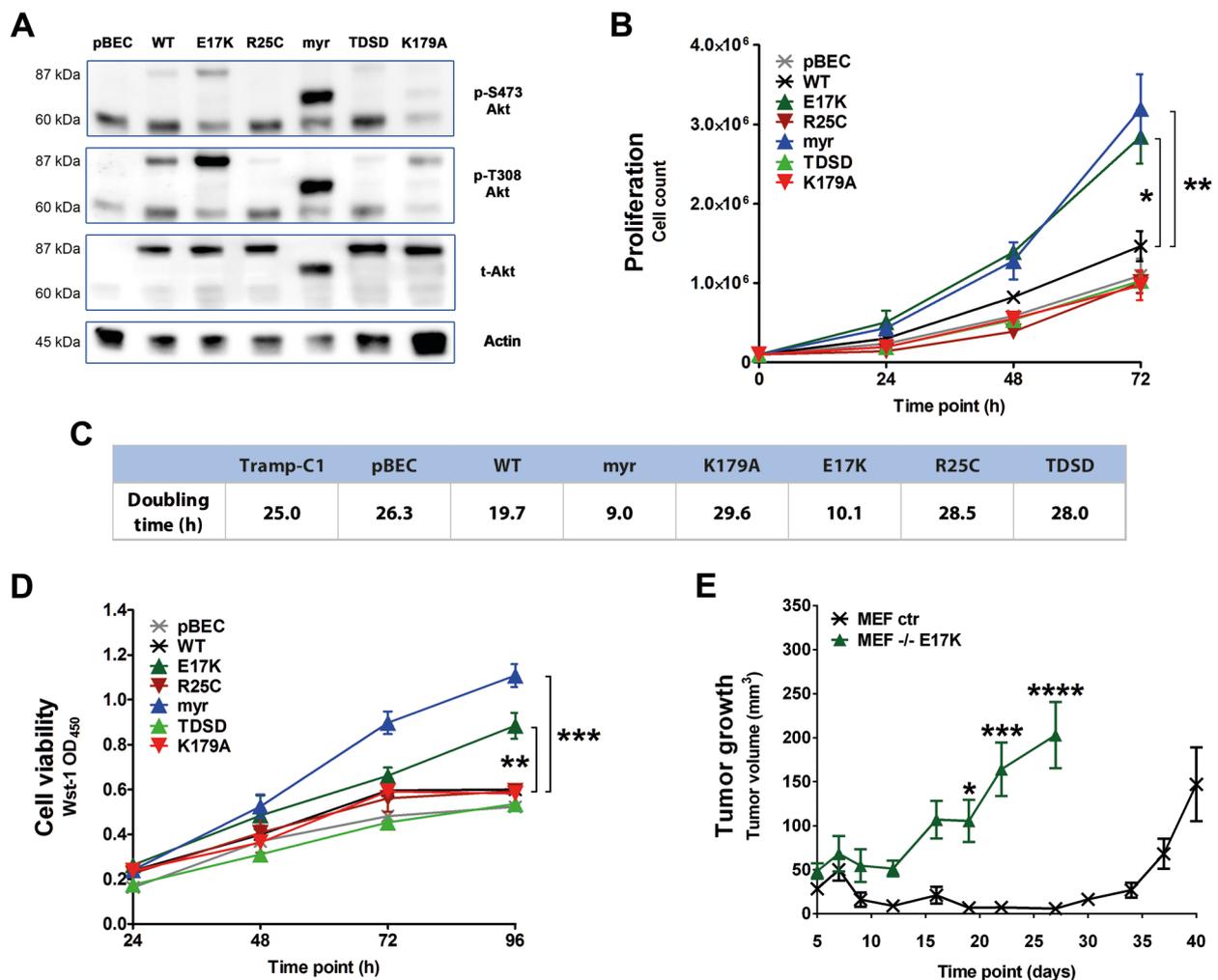
1 Supplemental figures

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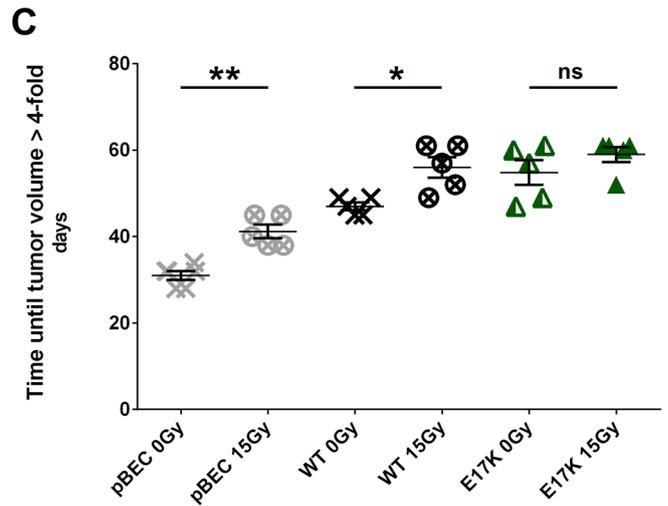
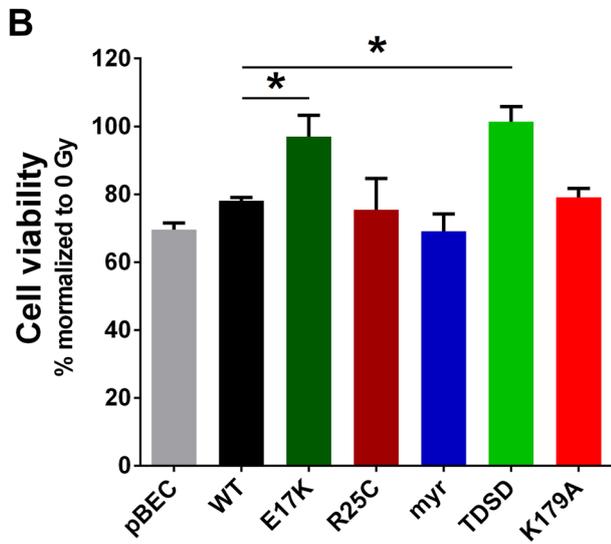
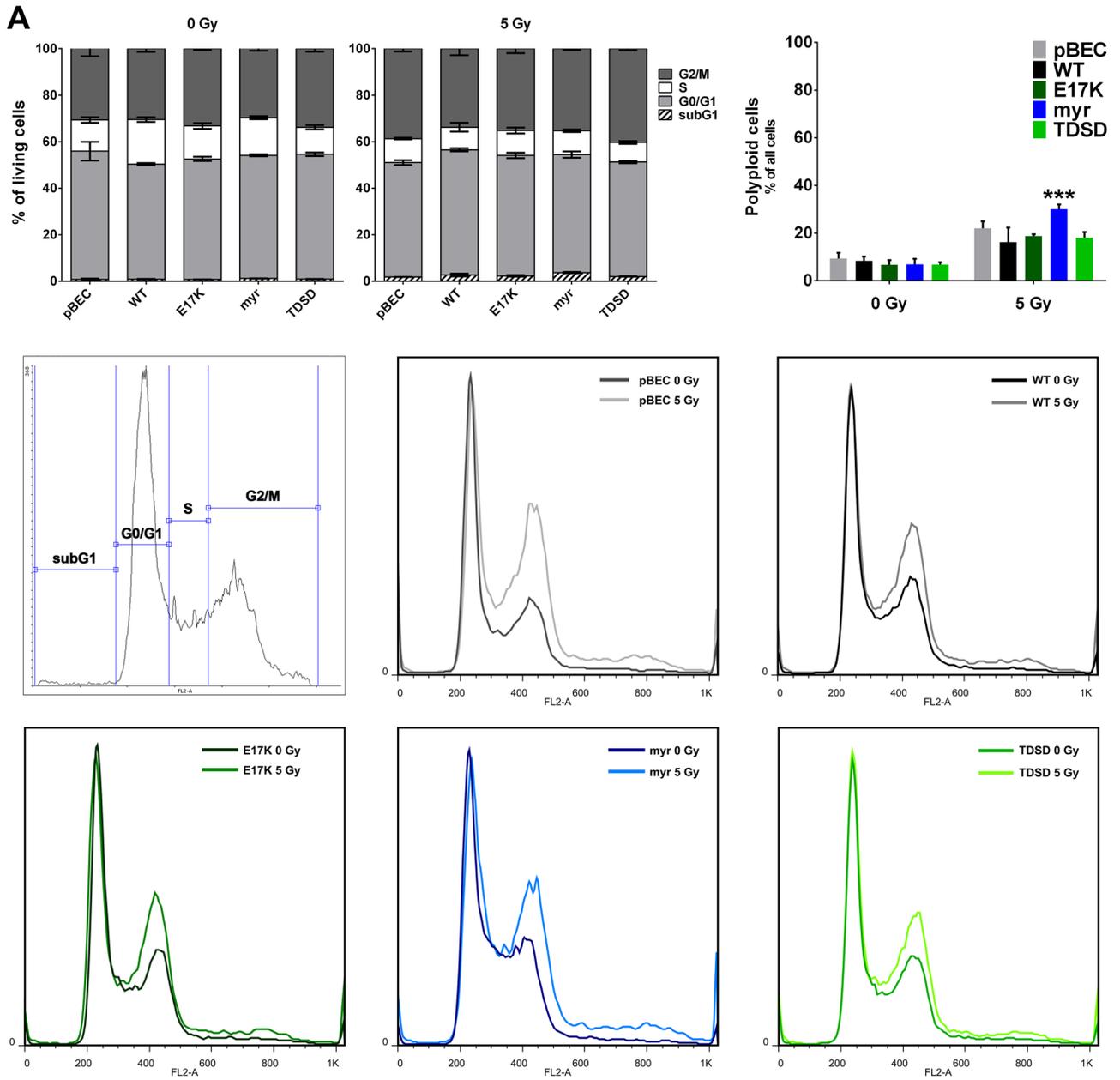
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4 **Supplemental Fig. 1: COSMIC database analysis of tumour tissues with enriched *Akt***
5 **expression.** Percentage of tumour samples expressing at least more than 2-fold levels of the
6 Akt1, Akt2 or Akt3 isoforms, respectively; data represent the percentage of all tumour samples
7 of the respective tissue.

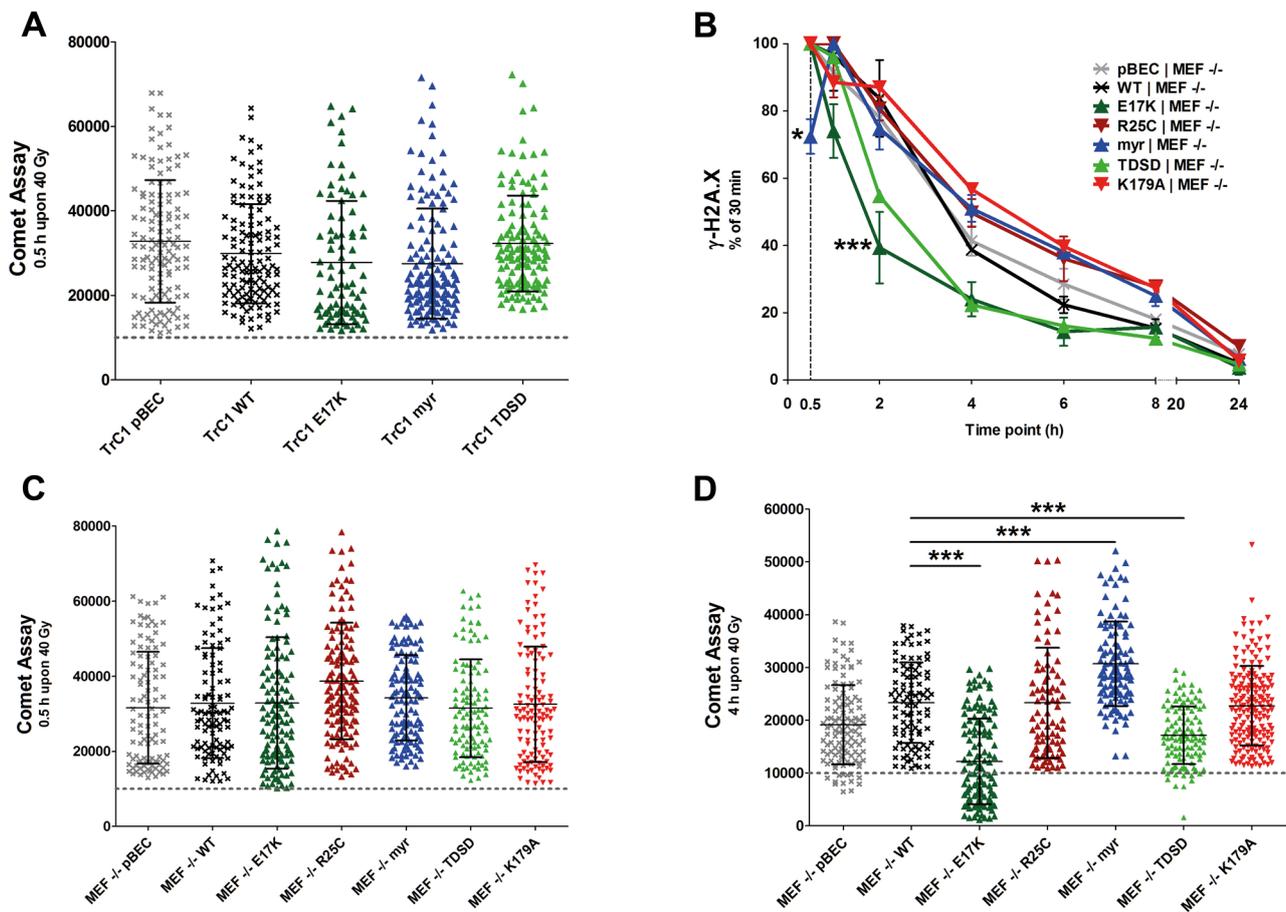


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2 **Supplemental Fig. 2: Akt1 mutations impact on Akt1-phosphorylation and cell**
 3 **proliferation. A** Expression levels and phosphorylation at S473 and T308 of the various Akt1-
 4 mutations expressed in TrC1 cells were analysed by Western Blot. **B** The number of viable
 5 cells was determined at the indicated time points by cell counting using a CASY cell counter.
 6 Data show means \pm SD (n=3). **C** Doubling times were calculated based on cell counting
 7 shown in **B**. **D** Determination of the number of viable cells at the indicated time points by using
 8 the Wst-1 assay. Data show means \pm SEM (n=4). **E** Growth curves of tumours derived from
 9 Akt1-WT MEFs or MEF Akt1 $-/-$ expressing Akt1-E17K in NMRI nude mice was followed by
 10 measuring the tumour volume twice weekly. Data show means \pm SD (n=6 mice per group). *
 11 $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$; ANOVA test with Bonferroni correction).

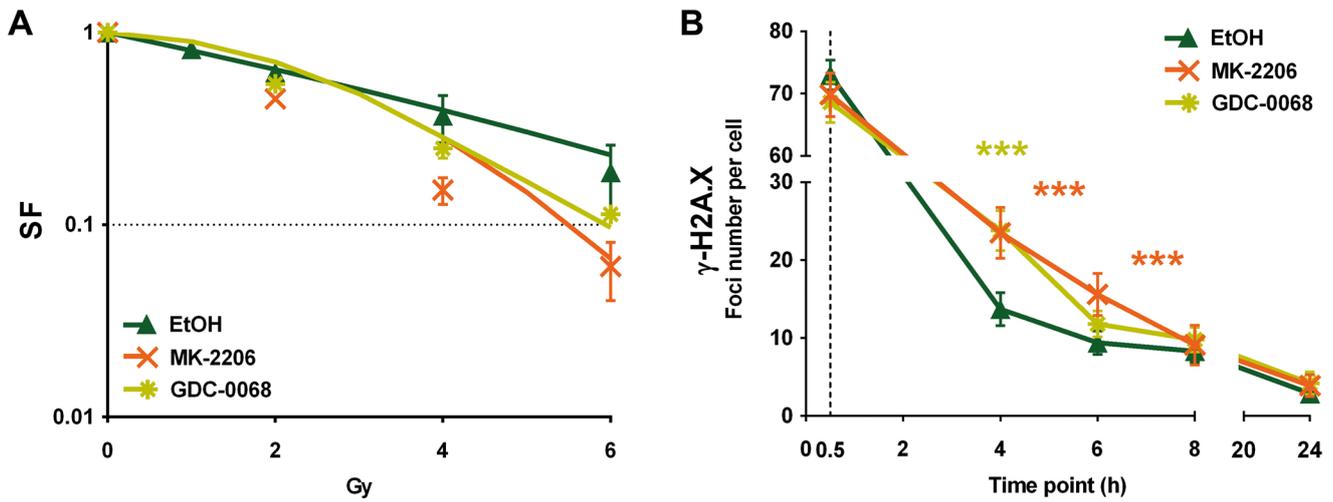


1 **Supplemental Fig. 3: Akt1-E17K mediates increased radiation resistance of TrC1 cells *in***
2 ***vitro* and *in vivo*. A** Cell cycle analysis of the various Akt1-mutants was performed without
3 radiation (upper left panel) and 24 h after a single dose of 5 Gy (upper middle panel). The
4 percentages of subG1, G0/G1, S, G2/M TrC1 cells are displayed. Additionally, polyploid cells
5 were analysed without IR and 72 upon 5 Gy (upper right panel). Data show means \pm SD (n=3).
6 *** p<0.001; ANOVA test with Bonferroni correction. The lower panels show the gating
7 strategy and representative histograms. **B** Effect on short-term survival of TrC1 cells measured
8 72 h after exposure to 5 Gy using the Wst-1 assay. Data show means \pm SD (n=3). * p<0.05, **
9 p<0.01; ANOVA test with Bonferroni correction. **C** Xenograft tumours of TrC1 cells expressing
10 pBEC, Akt1-WT or Akt1-E17K were generated on NMRI nude mice and irradiated 15 days with
11 0 Gy (sham controls) or 15 Gy. Data show the time until xenograft tumours reached 4-fold
12 volume. Data show means \pm SD (n=6). * p<0.05, ** p<0.01, ns: not significant; ANOVA test
13 with Bonferroni correction.



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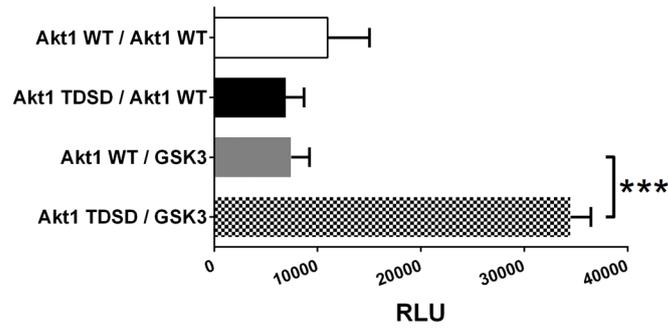
2 **Supplemental Fig. 4: Active Akt1 mutants alter DNA DSB repair in TrC1 and MEF Akt1 -/-**
 3 **cells. A** Determination of DNA fragmentation in TrC1 cells expressing pBEC or the various
 4 Akt1 mutants by neutral comet assay 30 min after irradiation with 40 Gy. **B** γ -H2A.X assay
 5 performed in MEF Akt1 -/- cells expressing the various Akt1 mutants at the indicated times on
 6 irradiation with 3 Gy. Determination of DNA fragmentation in MEF Akt1 -/- cells expressing the
 7 various Akt1-mutants by neutral comet assay performed 30 min (**C**) and 4 h (**D**) on irradiation
 8 with 40 Gy. Data show means \pm SD (n=120). *** p<0.001; ANOVA test with Bonferroni
 9 correction.



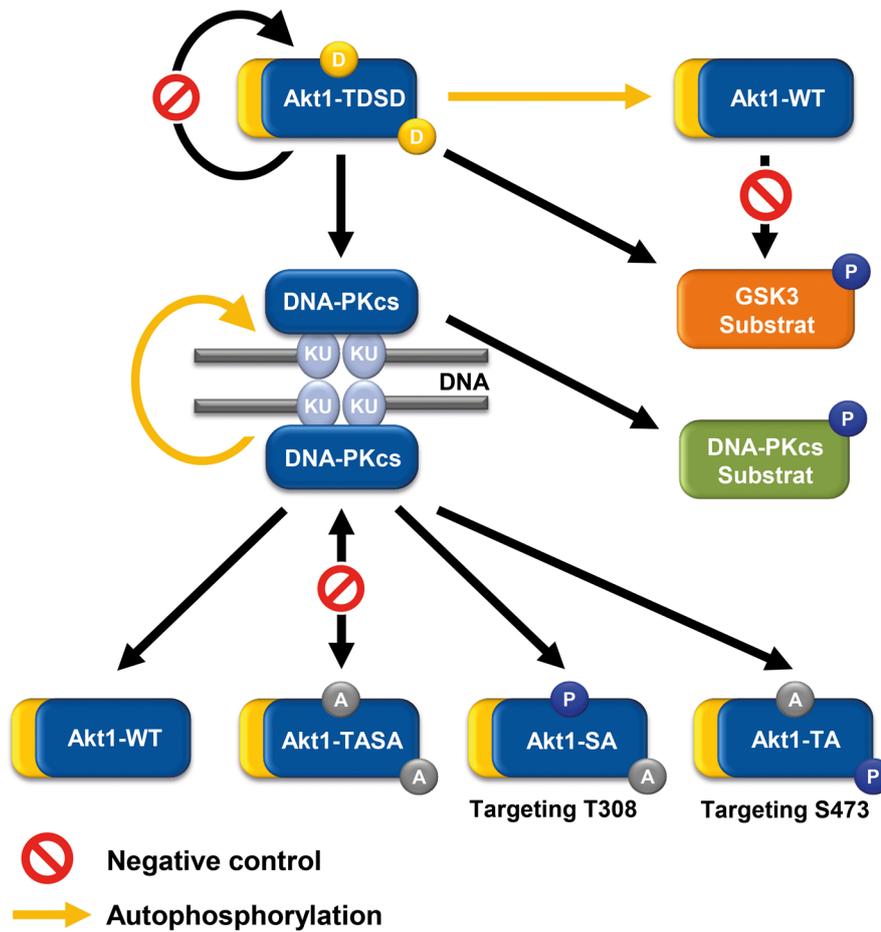
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2 **Supplemental Fig. 5: Clinically relevant Akt inhibitors negate the radioresistance and**
 3 **the DNA repair-promoting effect of Akt1-E17K. A** The effect of pre-treatment with 2 μM MK-
 4 2206 and GDC-0068 on the resistance-promoting effect of Akt1-E17K in TrC1 cells as
 5 determined by colony formation assays upon irradiation (0-6 Gy). EtOH was used as a solvent
 6 control. Data show means ± SEM (n=3). **B** The effect of pre-treatment with 2 μM MK-2206 and
 7 GDC-0068 on the beneficial effects of the active Akt1-E17K on the kinetics of DNA DSB repair
 8 in irradiated TrC1 cells (3 Gy) was analysed by γ-H2A.X assay. The kinetics of DNA DSB
 9 repair were followed by counting the amount of γ-H2A.X foci. Data represent the amount of
 10 foci at the indicated time points. EtOH was used as a solvent control. Data show means ± SD
 11 (n=3). *** p<0.001; ANOVA test with Bonferroni correction.

A



B



1

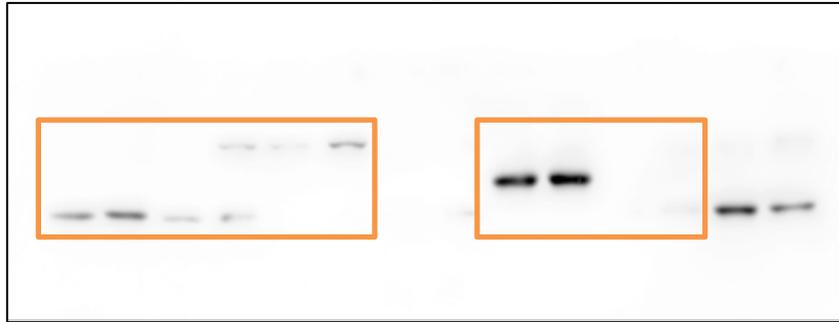
2 **Supplemental Fig. 6: Determination of the phosphorylation hierarchy between Akt1 and**
 3 **DNA-PKcs by an *in vitro* kinase assay. A** Kinase assay with Akt1-WT and TDS variants
 4 with GSK3 target peptide performed to detect the kinase activity. Data show means ± SD
 5 (n=3). *** p<0.001; ANOVA test with Bonferroni correction. **B** Scheme of experimental set-up
 6 of the Akt1/DNA-PKcs kinase assay.

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1 **Supplemental Full Western Blots**

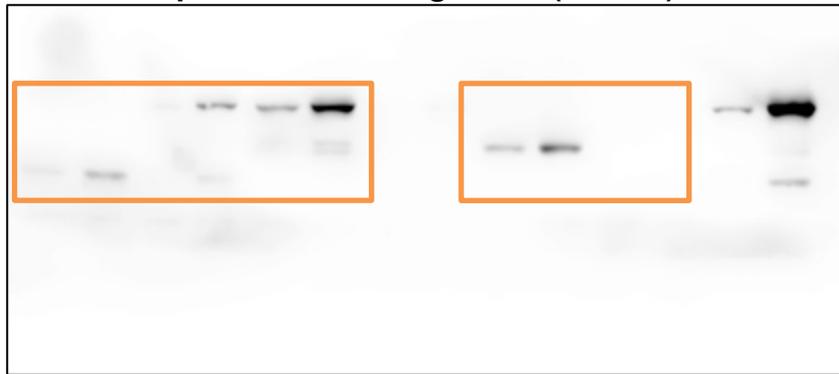
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p-T308 Akt1-eGFP (87 kDa)
p-T308 myrAkt1-eGFP (72 kDa)
p-T308 Akt endogenous (60 kDa)



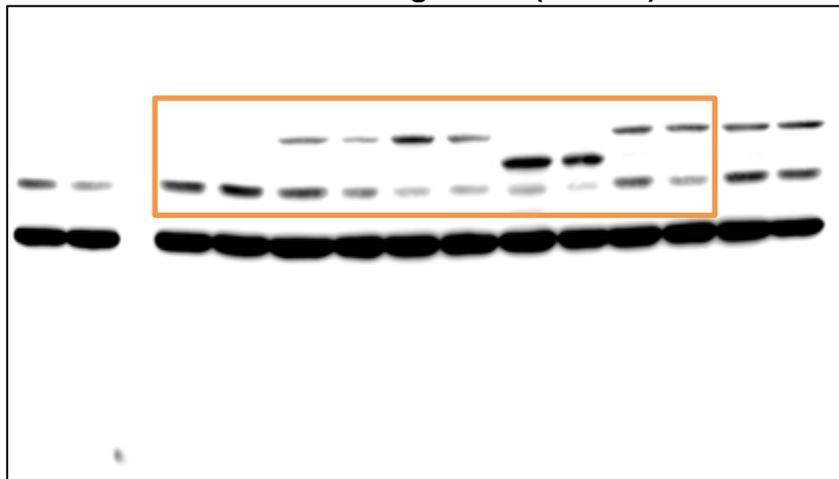
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p-S473 Akt1-eGFP (87 kDa)
p-S473 myrAkt1-eGFP (72 kDa)
p-S473 Akt endogenous (60 kDa)



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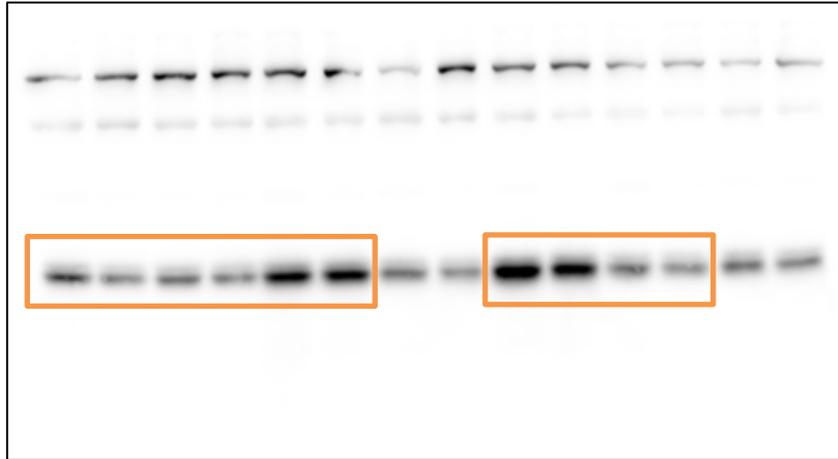
total Akt1-eGFP (87 kDa)
total myrAkt1-eGFP (72 kDa)
total Akt endogenous (60 kDa)



16
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p-GSK3 β (46 kDa)

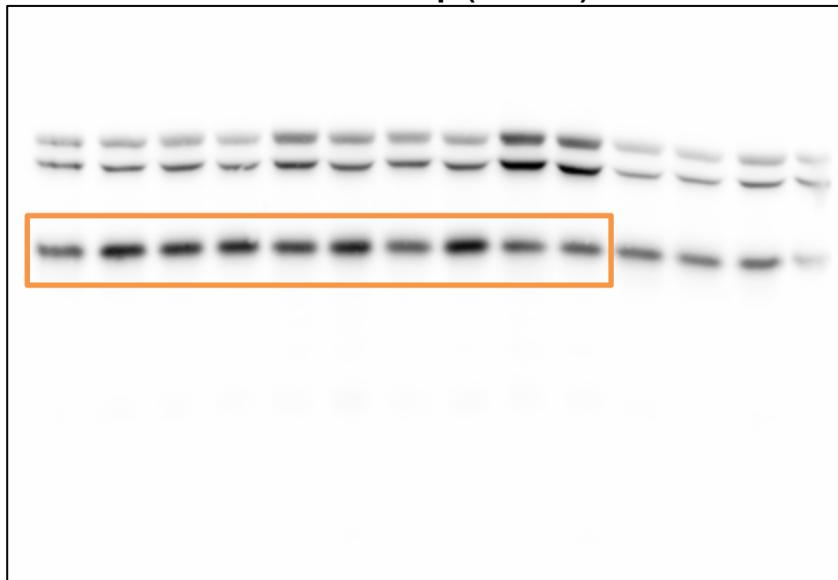


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Total GSK3 β (46 kDa)

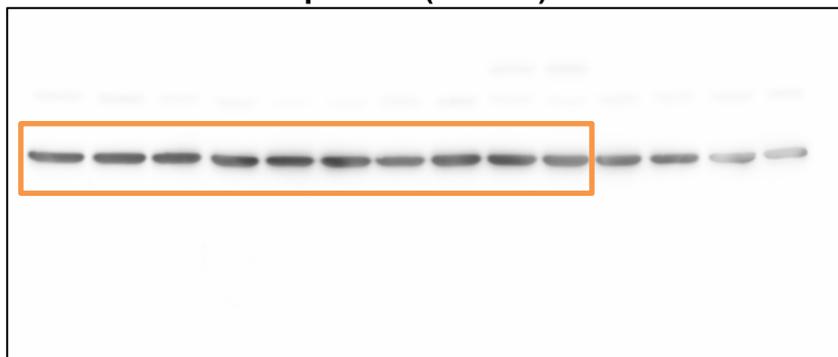


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β -Actin (43 kDa)



8

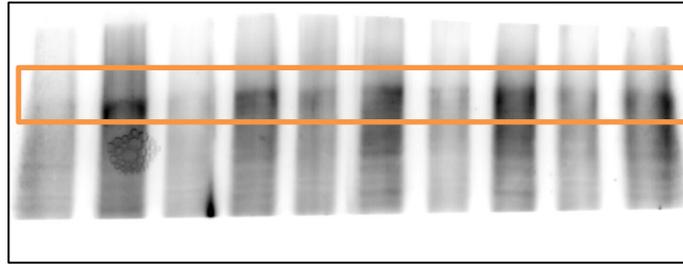
9

10 **Full Western Blots related to Fig. 3B: Akt1-E17K but not myrAkt1 shows pronounced nuclear**
11 **localization and displays higher phosphorylation upon IR.** Indicated proteins were detected
12 with protein-specific antibodies on Western Blots. Equal protein amounts (30 μ g, whole cell
13 lysate) were loaded. β -Actin was used as loading control.

14

1

p-DNA-PK (approx. 450 kDa)

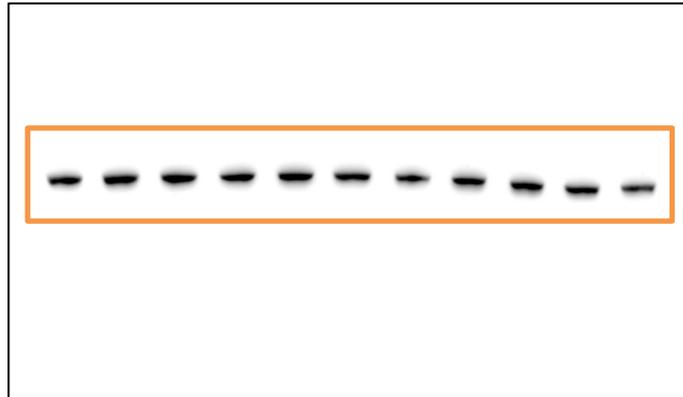


2

3

4

Total DNA-PK (approx. 450 kDa)

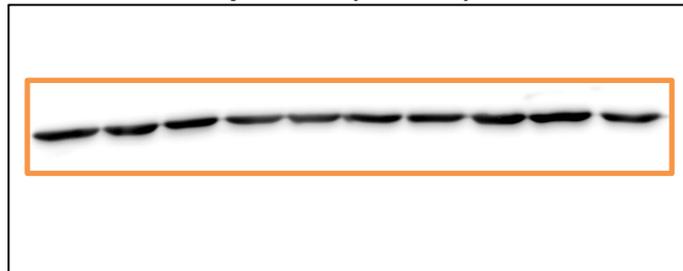


5

6

7

β-Actin (43 kDa)



8

9

10 **Full Western Blots related to Fig. 4C: DNA-PKcs phosphorylates Akt1 at S473 upon DNA**
 11 **damage and is required for radiation resistance mediated by Akt1-TDSD and Akt1-E17K.**
 12 β-Actin (43 kDa), total DNA-PK (approx. 450 kDa), p-DNA-PK (approx. 450 kDa). Indicated
 13 proteins were detected with protein-specific antibodies on Western Blots. Equal protein
 14 amounts (30 μg, whole cell lysate) were loaded. p-DNA-PK levels were normalized to their
 15 related total DNA-PK signals. β-Actin was used as loading control.

16