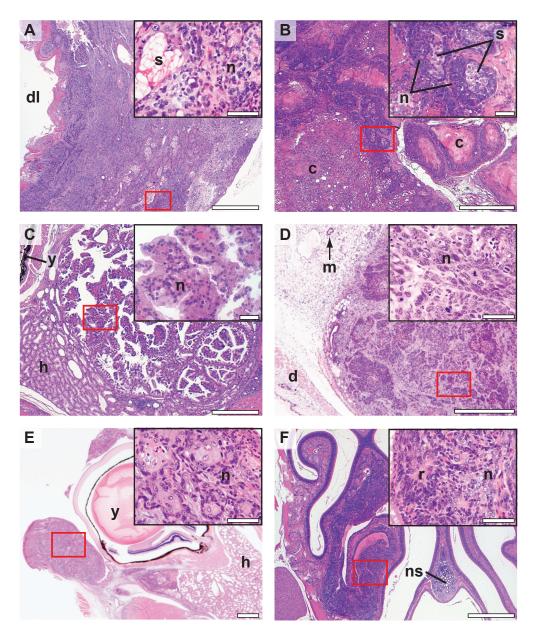
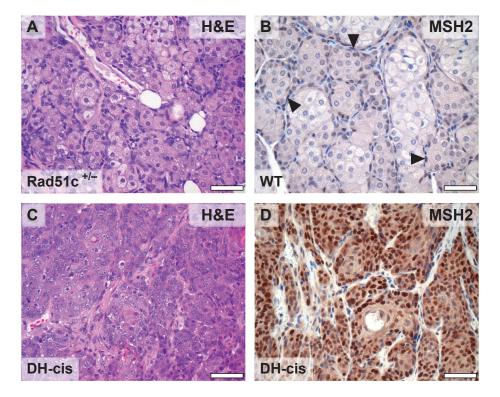
## **Supplementary Figure 1**



Supplementary Figure 1. Histology of tumors characteristic for Rad51c mutants. (A) Preputial gland carcinoma in a DH-cis male. (B) Zymbal's gland carcinoma in a DH-cis male; (C) Harderian gland adenoma in a  $Rad51c^{ko/+}$  male. (D) Mammary gland carcinoma in a DH-cis female. (E) Basal cell carcinoma, skin/subcutis, possibly Meibomian gland, in a DH-cis female; (F) Neuroepithelial carcinoma in a DH-cis male; Bowman's gland area, nasal section. Area shown in insets is marked with a red frame in each respective figure. s, normal sebaceous cells; n, neoplastic epithelial cells; dl, duct lumen; c, sebaceous cyst lumen; y, eye; h, Harderian gland; ns, nasal septum; r, neural rosettes. Scale bar corresponds to 500  $\mu$ m in the main figures and to 50  $\mu$ m in the insets.



Supplementary Figure 2. Msh2 is not downregulated in preputial gland carcinomas in DH-cis mice. ( $\bf A$  and  $\bf B$ ) normal control preputial gland stained with hematoxylin and eosin (H&E) and with antibodies against Msh2 (MSH2), respectively. Arrowheads indicate Msh2-positive basal cells. ( $\bf C$  and  $\bf D$ ) Preputial gland carcinoma from a DH-cis male stained the same way as the controls. Note the number of cells staining strongly positive for Msh2 in ( $\bf D$ ). Genotype is indicated in the lower left corner for each image. Scale bar corresponds to 50  $\mu m$ .

## **Supplementary Table 1.** *Rad51c* is essential for embryo viability.

Developmental stage	Number of litters dissected	Total number of embryos	Portion of Phenotypically abnormal embryos		Portion of Phenotypically normal embryos		Empty decidua, lost, or genotype not determined	Deviation from expected mendelian distribution <sup>†</sup> , <i>P</i> -value (χ2-test)		
			ko/ko	ko/+	ko/+	+/+				
E5.5	2	22	5 (2	3%)	17 (77%)		0	0.805		
E6.5	5	57	12 (2	21%)	44 (79%)		44 (79%) 1		1	0.531
E7.5*	12	132	38 (2	38 (29%) 94 (71%)		1	0.236			
	3	31	5 (18%)	0 (0%)	21 (68%)	5 (16%)	1	0.272		
E8.5*	12	126	49 (39%)		77 (61%)		0	0.0004		
	5	50	16 (32%)	2 (13%)	18 (36%)	14 (28%)	6	0.540		
E10.5*	5	51	14 (27%) <sup>§</sup>		37 (73%)		0	0.686		
	2	22	5 (23%)§	0 (0%)	8 (36%)	9 (39%)	1	0.213		
Newborn	59	395	0 (0%)	0 (0%)	239 (61%)	256 (39%)	0	2.723E-30 (0.009) <sup>‡</sup>		

<sup>\*</sup>Shaded fields indicate litters, from which embryos have been genotyped.

<sup>&</sup>lt;sup>†</sup>Deviation from the Mendelian 1:2:1 genotype and 1:3 phenotype ratio was tested. Statistically significant differences are highlighted in bold.

†The number of *Rad51c*<sup>ko/+</sup> and wild-type progeny was tested against 2:1 ratio.

<sup>§</sup>Embryos were almost resorbed.

## **Supplementary Table 2.** Complete list of observed neoplasms grouped by target organ, gender, and genotype.

	Females					Males				
Target organ and tumor type	WT	Rad51cko/+	Trp53 <sup>ko/+</sup>	DH-trans	DH-cis	WT	Rad51c <sup>ko/+</sup>	Trp53 <sup>ko/+</sup>	DH-trans	DH-cis
Adrenal										
Pheochromocytoma  Bone	-	-	-	-	-	-	-	-	1	-
Osteosarcoma	_	2	5	11	7	_		_	1	
Cecum										
Adenoma	-	-	-	-	-	-	1	-	-	-
Epididimis   Hemangioma	_	_				_			_	1
Fat pads, gonadal										
Hemangiosarcoma	-	-	-	-	-	-	-	-	-	1
Hematopoietic neoplasm Lymphoma, small cell	_	_		1	2	_	_	1	_	_
Lymphoma, follicular center cell (B-cell)	-	1	1	4	1	-	-	1	-	2
Lymphoma, diffuse large cell (B-cell)	-	-	-		-	-	-	-	1	-
Lymphoma, splenic marginal zone Lymphoma, NOS	-	-	 1	1	-	-	-	1	1	1
Histiocytic sarcoma	-	-	-	2	-	-	-	2	1	
Mast cell sarcoma	-	-	-	1	-	-	-	-	-	2
Liver										
Adenoma, hepatocellular Cholangiocarcinoma	-	-	-	-	-	-	-	-	1 1	-
Lung	_	_	-	-	_	_	_	-		
Adenoma, alveolar Type II cell		2	-	2	-	-	2	3	1	1
Carcinoma, alveolar Type II cells	-	2	-		-	1	1	3	-	-
Mammary gland Adenocarcinoma	_	2	3	1	7	_	_	_	_	_
Muscle										
Rhabdomyosarcoma	-	-	-	-	2	-	•	5	5	1
Sarcoma, NOS Hemangiosarcoma	-	-	- 1	•	-	-	1	6	8	5
Mediastinum	-	-		-	-	-	-	-	-	-
Myxosarcoma	-	-	-	1	-	-	-	-	-	-
Mesentery										
Hemangiosarcoma Ovary	-	-	-	•	-	-	-	•	-	1
Cystadenoma	_	-	-		-	-	-	-	-	-
Granulosa cell tumor	-	-	-	-	-	-	-	-	-	-
Hemangiosarcoma	-		-	1	2	-	-	-	-	-
Pituitary Adenoma, pars distalis	2	2	_		1	_		_		_
Salivary gland	_									
Myoepithelioma	-	-	-	-	1	-	-	-	-	-
Sebaceous glands Preputial gland carcinoma	_	_							1	10
Zymbal's gland adenocarcinoma	-	-	-	2	-	-		1		5
Spinal cord										
Neurofibrosarcoma	-	-	-	-	-	-		-	-	1
Spleen Hemangiosarcoma	_	_	_	_	1	_	_	_	_	_
Skin					•					
Hemangiosarcoma	-	-	1	-	1	-	-	-	-	-
Myxosarcoma / myxoma Squamous cell papilloma	-	-	-	-	2	-	-	-	1	-
Squamous cell carcinoma	_	-	-	:	1	-			1	2
Sarcoma, NOS	-	-	-	-	-	-	-	-	1	-
Mast cell tumor	-	-	-	-	-	1	-	-	-	-
Trichoepithelioma Testis	-	-	-	•	-	-	-	-	-	-
Neoplasm, NOS	-					1			-	
Thyroid										
Adenoma, follicular cells	-	-	-	-	1	-		-	-	-
Urinary bladder Submucosal mesenchymal tumor	_	_	1		_	_		_	_	_
Uterus										
Polyp, endometrial	-	-	-	:	2	-	-	-	-	-
Leiomioma Hemangiosarcoma	-	-	-	1 -	-	-	-	-	-	-
Muzzle area		-	-	-	-		-	_	-	-
Harderian gland adenoma / carcinoma	1	1	-	-	-	-	2	1	1	1
Bowman's gland neuroepithelial carcinoma	-	-	-	;		-	•	1	-	1
Olfactory neuroblastoma  Meibomian gland, squamous cell carcinoma	-	•	-	1	1 1	-		-	-	1 1
Vomeronasal neuroepithelial carcinoma	-	-			1	-	-	-	-	
Squamous cell carcinoma	-	-	-	-	1	-		-	-	1
Carcinoma, NOS	-	-	-			-	-	-	1	-
Total number of animals	17	38 42	11 10	26 25	23	21	32	20 16	24	24
Number of animals that died by 600 days Total number of animals with neoplasms	2 2	12 7	10 8	25 21	23 22	3 2	9 6	16 16	23 22	23 22
Total number of primary neoplasms	3	12	13	30	35	2	7	25	27	38

## **Supplementary Table 3.** Quantification of Rad51 and γH2AX foci in *Rad51c*-deficient and control MEFs.

Genotype	Genotype Treatment		Average number of Rad51 foci per cell	Average number of γH2AX foci per cell	P-value*	
wt	none	62	2.3	4.5	n.a.	
Rad51c <sup>ko/+</sup>	none	64	2.0	3.4	0.380	
Trp53 <sup>ko/ko</sup>	none	55	1.9	3.1	0.350	
Rad51c <sup>ko/ko</sup> ; Trp53 <sup>ko/ko</sup>	none	50	0	10.6	0.001	
wt	IR	55	16.2	48.3	n.a.	
Rad51c <sup>ko/+</sup>	IR	52	18.7	49.0	0.507	
Trp53 <sup>ko/ko</sup>	IR	50	29.1	67.3	0.007	
Rad51c <sup>ko/ko</sup> ; Trp53 <sup>ko/ko</sup>	IR	51	0	37.5	<0.001	

<sup>\*</sup>Statistically significant differences from wild-type cells are highlighted in bold (Wilcoxon text).