Sarcoma entity	1.Genetic mutation	2. Promotor	3. Genetic strategy	4. Mode of activation	5. 2 nd hit	6.Tumor developd? Histology?	7. Mouse strain	8. Penetrance (%) / Latency (Days/weeks)	9. Model details / Significance	10. Reference	
1. UPS/ Undifferentiated	Gli2A (constitutively active) KRASG12D	CAG, PCP2-Cre Rosa-26	Cag-LSL-Gli2A(Germ line transmission) R26 ^{CreER} ; p53 ^{flox} ; K ^{FRT}	Pairing with PCP2-Cre P53 inactivation by systemic Tamoxifen (i.p.), local Kras activation	Pairing with SmoM2 embryonically lethal P53 inactivation, Cardiotoxin	Yes, small round cell (Ewing-like features)	e Mixed C57BL/6/129Sv Mixed	98%/ 8 weeks 100%/8weeks (simultaneous P53 deletion and KRAS	Gli-2induces UPS with Ewing-like-featuresby expression of EWS-ETS target genes, inlcuding Nkx2.2. Muscle injurycooperates with P53 inactivation for sarcomagenesis. P53-/- plus	Fleming et al, Molecular Cancer Research, 2019 Van Mater et al, J Cl Insight, 2018	
pleomorphic sarcoma	KRAS ^{G12D}	Rosa-26	KRAS ^{LSL-G12D} /+- ROSA26 ^{LSL-Cas9eGFP} -mice	(Flipase)	injection i.m. P53 inactivation via CRISPR	Yes, UPS (15% myogenic, 85%	C57BL/6/129Sv Mixed	activation);8%/11 weeks (RRAS expression,3 weeks later P53 deletion);90%/6weeks (RRAS expression,3 weeks later P53 deletion,CTX treatment at day 10) Ad-P-Cre:100%/10 weeks; EPO with Cas9 plasmid: 80%/11 weeks; EPOwith endogenous Cas9: 100%/7weeks	muscle-injury-mediated UPS show chromosomal amplifications including YAP1/ MET. In vivo electroporation can be as effective as lentiviral delivery for	Huang et al, Nature	
	P53 inactivation or hotspot mutation	-	Conditional delection of P53 ^{Lox/Lox} and Pten ^{Lox/Lox}	plasmid Local injection of Ad5-CMVE-Cre (Adenovirus) i.m. or s.c.	: -	non–myogenic) Yes, 93% UPS; 7% pleomorphic RMS	C57BL/6/129Sv C57BL/6	100%/10weeks	conditional tumor induction. First Pten-inactivatedUPS model.	Communications, 2017 Buchakjian et al, PlosOne, 2017	
	P53 and Pten inactivation KRASG12D	SFFV Rosa-26	with KRAS ^{G12D} ,transplanted into hindlimbof P53-/+mice	Constitutive expression Systemic tamoxifen	P53-/-, Cardiotoxin injection	Yes, High grade sarcoma with myofibroblastic differentiation Yes, Spectrum of myogenic and non myogenic arcoma: My77-CreBC-28UPS_248 eMSLyses 1 MiS (748 myogenic, 26	•	85%/ 5 weeks Myf7-CreER: 100%/ 6 weeks,	Mutant KRAS and P53 inactivation cooperate in myoblasts for sarcomagenesis. RMS and UPS have distinct and overlappingcells of origin within the muscle lineage. Pax7+MvoD+	McKinnon et al, Oncotarget, 2015	
	KRAS ^{G12V}	Cyp1A1	Pairing of LSL-KRASCINO-PS3*/-micewith 2 different Tamoxifen-inducibleCre lines: Pax?-CreER(Quiescent & activated satellite cells, myoblasts) & MyoD-CreER(activated MyoD-satellite cells and progenitors) Systemic overexpression/loss(germ L>10	administration Crossing with Ah–Cremice (sceletal	i.m. P53+/R172H, P53flox/R172, P53flox/flox, P53flox/+	My77-CreER-62%UPS,24% eRMS,rest RMS (74% myogenic,26 non-myogenic)MyoD-CreER-100%UPS (60% myogenic,40% non-myogenic); Yes, UPS	C57BL/6/129Sv Mixed	MyOD-CreER: 100% / 21 weeks P53"***R278": 88%/10 weeks: P53"****R578": 100%/7 weeks; P53"************************************	RMS and UPS have distinct and overlappingcells of origin within the muscle lineage. Pax7+MyoD+ quiescent satellite cells can be a cell of origin for RMS, Pax7+MyoD+cells of origin for UPS. Mutant P53 (P53 ^{8172H)} is an even more potent activator of tumorigenesis in Ras-drivenUPS than loss of	Blum et al, Cell Reports, 2013 Doyle et al, Journal of Pathology,	
	KRAS ^{G12D}	Rosa-26	Conditional KRAS ^{G12D} overexpression	muscle) for systemic activation Local injection of Ad-Cre (adenovirus) into leg and uterus	P53 ^{Flox} , Cdkn2a ^{Flox}	Yes, High grade sarcoma with myofibroblastic differentiation	C57BL/6/129Sv C57BL/6	>90%/13 weeks	P53. Low rate spontenous metastasis (13%) was only observed immutant P53,but not loss of P53 group KRAS-drivensarcomagenesis cooperates with Cdkn2a inactivation, but not with inactivation of Bak1 and Bax. Foxm1 expression facilitates UPS metastasis.	2010 Kirsch et al, Nature Medicine, 2007: Mito et al, Plos ONE, 2009;	
	KRAS ^{G12D}	Ryr2	Conditional KRAS ^{G12V} overexpression	Cre vector pCAGnlCre delivered via electroporation of the gastrocnemius muscle of 8–10week old mice	P53 ^{-/-} & P53 ^{+/-}	Yes, pleomorphic RMS (Myogenin, MyoD expression not tested)	Mixed C57BL/10/129S	100%/6 weeks (P53 ^{-/-}) & v 40%/11 weeks (P53 ^{+/-})	Cooperation of oncogenic RAS and P53 inactivation leads to very efficient pleomorphic sarcoma development.	Tsumura et al, Oncogene, 2006	
	P53 inactivation or hotspot mutation Dystrophin	-	Embryonic stem cell alteration, crossing of mice Crossing of mice, additionally Barium		- P53 ^{-/+} & P53 ^{-/-}	Yes,various entities,especially lymphoma and sarcoma (typically UPS,more rarelyalso OS,RMS and Angiosarcoma) Yes, RMS not further specified	Mixed C57BL/6/129Sv C57BL/6		First in vivo models for P53 inactivation, including typical mutations for Li-Fraumeni tumor predisposition syndrome. Duchenne muscular dystrophyseverity facilitates RMS development. Muscle stem cells	Donehower et al,Nature,1992.) acks et al,Current Biology,1994;Lang et al,Cell,2004;Olive et al,Cell, 2004; Boscolo Sesillo et al, Cell Reports,	
2. eRMS/ Embryonal/	(MDX/MtR mouse) Mdm2-ALT1(Splice	promotor),SOX2-Cre,also Cd19	chloride injection to induce muscle damage and regeneration Conditional alleles activated by crossing to Cre line	Constitutive (germline)	background P53+/-background	Yes,eRMS in 50% of Mdm2-ALT1 ^{-/+} and0% Mdm2-ALT1 ^{-/-} ,Restlymphoma,osteosarcoma hemanoinosarcoma terratoma	C57BL/6	MDX/mTR +P53+: 100%/17weeks; MDX/mTR +P53+': 10%/56 weeks; MDX/mTR +P53+': 10%/56 weeks; Mdm2-ALT1-'++P53+': 100%/20weeks (50% of which are eRMS); Mdm2-ALT1-'++P53-'-:	acquire RMS-like gene signature even before transformation. Mdm2-ALT1increases P53-loss-mediatedeRMS tumorigenesis	2019 Comiskey et al, Oncogene, 2018	
Fusion-negative rhabdomyo- sarcoma &	hyap1 ^{S127A}	Cre for lymphoma development Col1a,Cre lines: Pax7-cre/ERT2, Myf5-Cre, Myod1-iCre;	Rosa26-LsL-Col1a-rtTA-TetO-YAP1 ⁵¹²⁷ mice paired with different Cre lines	Constitutive or by systemic Tamoxifen administration	Cardiotoxin and Bariumchloride i.m.	Yes, eRMS	Not reported	100%/27weeks (most of which are lymphoma) 4–8weeks, penetrance not reported	YAP1 hyperactivity in activated, but not quiescent satellite cells, induces eRMS.	Tremblay et al, Cancer Cell, 2014; Slemmons et al, PlosOne, 2015;	
pleomorphic RMS	SmoM2 (Constitutively active) Her-2/neu	Rosa26,Cre lines: aP2-Cre, MCK-Cre, Myog-Cre, Myf5-Cre; MMTV-LTR	different Cre lines via mouse crossing Systemic overexpression(apart from	Constitutive (germline) Constitutive (germline)	Cdkn2a inactivation P53 ^{-/+}	Yes, eRMS Yes, genitorunary eRMS, only in	Mixed	80%/6weeks for adipocyte-specific aP2-Cre; 100%/4 weeks (+Cdkn2a); Close to 100%/17 weeks	SHH activation in adipcyte progenitorscan efficiently induce eRMS, particularly in cooperation with loss of Cdkn2a. <i>Iqf2</i> , <i>p19Arf and p21Cip1 are upregulated in preneoplastic tissue</i> .	Hatley et al, Cancer Cell, 2013 lanzano et al, Oncotarget, 2013	
	Dystrophin (MDX	-	mammary gland),Crossing of mice Crossing of mice,plus Cardiotoxin (CTX) injection to induce muscle	Constitutive (germline)	P53-/-& P53+/-	males Yes, eRMS	Mixed	Mdx/P53***: 9%/43 weeks;Mdx/P53***: 60%/26 weeks; Mdx/P53***: 90%/17 weeks; Mdx/P53** plus CTX: V 100%/13weeks;Mdx/P53** plus CTX: 0%/P53** alone:	P53 inactivation acclerates eRMS induction in dystrophin-inactivatedmice. Muscle damage and regeneration further increases efficiency.	Camboni et al, J ournal of	
	mouse) P53, Ptch and Rb1	Pax7CreER, MCre, Myf5Cre, Myf6Cre;	damage and regeneration; Conditional alleles activated by 4 different Cre lines via mouse crossing	Constitutive and via Tamoxifen (Pax7CreER)	Dackground All tumors P53 inactivated, some also Ptch-, some also Rb1-inactivated	Yes, eRMS,but also UPS,OS and others depending on driver and Cre line	Mixed	20%/26weeks.	eRMS and UPS lie in a continuum(Satellite cells predisposed towards UPS, maturing myoblasts towards eRMS).	Pathology, 2012 Rubin et al, Cancer Cell, 2011	
	Dystrophin (MDX mouse) & Alpha-Sarcoglycan (Sgca)	- EIIA-Cre and	Crossing to hetero-and homo- zygosity of either gene, crossing with mice inactivated for 2nd hits Ptch+/flox & Ptchflox/flox	Constitutive (germline) Systemic tamoxifen	Galgt1, Galgt2, Cmah	Yes, eRMS Yes,RMS and Medulloblastoma upon neonatal	Mixed C57BL/6 & C57BL/10 Mixed Balb/c	9% for mutated Dystrophin, 4% for Sgca ^{-/-} /73 weeks 20% RMS (not specified,probably eRMS-like) & 9%	Muscle dystrophy-relatedmutations in Dystrophinand Sgca can both lead to RMS in aged mice. Tumors exhibitMdm2 & P53 amplification with cancer-associated P53 missense mutations. Prenatally, but not postnatally induced Ptch heterozygosity resulted in the formation of RMS,	Fernandez et al, American J ournal of Pathology, 2010	
	Ptch P53 and Rb1	ROSa26-CreERT2 SM22alpha (Specificity for smooth muscle and embryonic	Microinjection of embryos with SV40-Tumorantigen (TAg) to	administration embryonically and postnatally	-	(Ptch**/flas), basal cell carcinoma upon postnatal Ptch inactivation (Ptch**/flas) Yes, cardiac RMS, not further specified, also smooth muscle proliferation in	and others FVB/N	medulloblatomä/21 weeks,but only upon neonatal induction (Patch*/flax) 12 weeks (8 mice in total, penetrance	accompanied by the silencing of the remaining wild-typePtch allele. Homozygous Ptch loss was embryonically lethal,but led to basal cell carcinoma (no RMS) when induced postnatally.	Zibat et al, Cancer Cell, 2009 et al, PlosOne, 2017 Köbbert et al, Journal of Thoracic	
	inactivation Dystrophin (MDX mouse)	cardiac muscle)	target P53 and RB1 MDX mouse model of Duchenne muscular dystrophy (spontaneous point mutation in exon 23 of the	Constitutive (germline)	-	vessels Yes, late in life aRMS-like(fusion gene presence not analyzed, probably rather eRMS)	C57BL/10	not reported) 6%/87 weeks	Mice of MDX model of Duchenne muscular dystrophy can develop RMS late in life.	and Cardiovascular Surgery, 2008 Chamberlain et al, FASEB J ournal, 2007	
	Sufu+/-(Suppressor of fused)		Dystrophin gene) Systemic Gene trapping	Constitutive (germline)	P53 ^{-/-} background	Yes, eRMS, medulloblastoma	& CD1	lymphoma (probably due to P53 loss alone)	Sufur'+is tumorigenic (including eRMS), but only on P53-'-background. Sufur'- embryonically lethal. Sufu mutations not as tumorigenicas Ptch mutations.	Lee et al, Oncogene, 2007	
	SmoM2(Constitutively active),Ptch Her-2/neu	Rosa26 for SmoM2,CAGGs for Cre-ER(systemic) MMTV-LTR	SmoM2 (Consitutively active form of SMO) Systemic overexpression	Systemic tamoxifen administration Constitutive (germline)	Ptch ^{-/+} background P53 ^{+/-} background	Yes, eRMS, basal cell carcinoma, medulloblastoma, pancreatic mucinous neoplasia Yes, eRMS	Mixed 129Sv/Swiss Webster as main components Balb/C	100%/5 weeks upontamoxifen for RMS (100% for basala cell carinoma,40% for medulloblastoma and pancreas lesions) 100%/14 weeks	Smo-mediatedSonic Hedgehoc signaling can induce multifocal eRMS with high efficiency. Her-2/neuefficiently drives genitourinary eRMS in male mice on	Mao et al, Cancer Research, 2006 Nanni et al, Cancer Research,	
	FOS & P53	-	(apart from mammary gland), Crossing of mice Systemic inactivation,Crossing of mice	Constitutive (germline)	-	Yes, eRMS	Mixed C57BL/6/129Sv	90%/16 weeks for homozygous double knockout, less efficient in heterozygosity	P53 ^{-/-} background, but not in females. Fos/P53 double knockout mice develop eRMS in facial and orbital regions.	2003 Fleischmann et al, Cancer Cell, 2003	
	HGF/HF	MT-1	Crossing of mice	Constitutive (germline)	Ink4a/Arfinactivation	,	C57BL/6	Mixed FVB/C57BL/6	Ink4a/Arfinactivation increases efficiency of c-Met-tumorigenicity.	Sharp et al, Nature Medicine, 2002	
	Ptch (Patch) HGF/HF	- MT-1		Constitutive (germline) Constitutive (germline)	-	Yes, eRMS Yes, RMS not further specified, amelanotic	& CD1 FVB/N	9% in CD1,2% in C57BL/6/latency na 7% RMS/latency na	Germline GEMM of Corlinsyndrome with typical features and eRMS (with Gli1 and Igf2 overexpression). Ptch:/-embryonically lethal. Activation of c-Met-tyrosinekinase via autocrine signaling is	Hahn et al, Nature Medicine, 1998 Takayama et al, PNAS, 1997	
2 - 51464	Pax3-Foxo1	Endogenous Pax3 locus;	Crossing of mice Conditional alleles, targeted via		Inactivation of Cdkn2a, Pax3	melanoma, nepatic and mammary tumors	Mixed		tumorigenic. **Activated Hippo signaling increases tumorigenesis in	Oristian et al, Cancer Research,	
3. aRMS/ Alveolar/	Pax3-Foxo1	Cre line: Myf6Cre Endogenous Pax3 locus;Cre lines: Pax7CreER,Myf5CreER, Myf6CreER,Mcre;	Cre-LoxP-conditional	with Cre lines Systemic Tamoxifen administration at P30	& Stk 3&4 P53,Pax3 inactivation	Yes,aRMS (Pax7CreER tumors showed spindle/pleomorphic morphology)	C57BL/6/129Sv Mixed C57BL/6/129Sv	MCre: 40%/29 weeks;Myf6CreER: 100%/15 weeks; Pax7CreER: 65%/48 weeks;Myf5-Cre: Embryonically lethal except for one (tumor-bearing) mouse;	Cdkn2a-inactivatedaRMS. aRMS can arise from different muscle lineages, particularly efficient in maturing myoblasts.	2018 Abraham et al, Genes and Development, 2014	
Fusion-positive rhabdomyo- sarcoma	Pax3-Foxo1	Endogenous Pax3 locus;Cre lines: Myf6 (differentiateing myogenic cells);	Pax3-Foxo1 knockin allele Systemic overexpressionvia Cre-LoxP-conditional Pax3-Foxo1 knockin allele (partly plus conditional Foxo1-knockoutallele)	Crossing of conditional mice with Cre lines	Ink4a/Arfor P53 inactivation	Yes, aRMS	Mixed C57BL/6/129Sv	0.5%/55weeks (fusion alone);8%/31 weeks (P53**);40%/13 weeks (P53**); 0% (lnk4a/Arf**);29%/10 weeks (lnk4a/Arf**);	Pax3-Foxo1leads to tumorigenesis when expressed in Myf6+cells, efficiency is increased by P53 and Cdkn2a deletion.	Keller et al, Genes and Development, 2004 (2 articles) & Nishijo et al, Cancer Research, 2009	
Jarcoma	Pax3-Foxo1	Endogenous Pax3 locus (PGK-, MyoD-and rat-beta-actin promotor also tried)	Pairing of mice	Constitutive (germline)	-	No	Mixed C57BL/6/NMRI	-	Heterozygous and chimeric mice showed developmental muscle defects and died perinatally from cardiac/respiratory failure. Pax3-Foxo1 expression from PGK-,MyoD-and rat-beta-actinpromotor did not yield any phenotype.	Lagutina et al, Molecular and Cellular Bioloyg, 2002	
4. MYOD1–RMS 5. OS/	No modeling attempts identified in the literature. Only one comprehensively characterized cell line model described (Schleicher et al, Cells, 2020). **Rb, P53** - **Conditional knockdown Crossing with Osx—Cre* - **Yes, OS (osteoblastic subtype)** C57BL/6** **Test Osteoblastic subtype C57BL/6** **In contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the contrast to other osteosarcoma models that used Cre:lox mediated gene deletion, of the con										
Osteosarcoma	SV40 TAg		(shP53) & conditional knockout Rbflox/+, Rbflox/flox SV40 expression under the Og2 promoter	mice	-	Yes, OS	FVB	100%/11-20weeks	osteosarcoma generated through shrNA-based knockdown of p53 results in osteoblastic OS, the most prevalent form sub-type of human osteosarcoma. Prkarla is a bone tumor suppressor gene capable of directing	Molyneux et al, The J ournal of	
	Rb, P53	-		Crossing with Prx-1-Cre	-	Yes, OS (and soft-tissue sarcoma with lower penetrance)	C57BL/6	92%/19 weeks	subclass development in mouse and human OS. The use of Cre in Prx-positive progenitors also leads to development tumors of other mesenchymal linages at high frequencies.	clinical investigation, 2010 Lin et al, Carcinogenesis, 2009; Calo et al, Nature, 2010	
	P53	_	Conditional inactivation of P53 ^{flox/+} , P53 ^{flox/flox}	Crossing with Prx-1-Cre	-	Yes, OS (and soft-tissue sarcoma with lower penetrance)		22%/96 weeks for heterozygous; 61%/50 weeks for homozygous mice;	The use of Cre in Prx positive progenitors also leads to development tumors of other mesenchymal linages at high frequencies.	Lin et al, Carcinogenesis, 2009; Calo et al, Nature, 2010	
	P53 Rb, P53	-	Conditional inactivation of	Crossing with Col1A1–2.3–Cremice Crossing with Osx–Cre	- -	Yes, OS Yes, OS	C57BL/6 Mixed	85%/ 5 weeks 100%/ 23 weeks; 100%/18 weeks;	Deletion of P53 in committed osteoblasts via Col1a1–Crestill results in a high percentage of OS like in Prx1–creprogenitors. Cooperation of P53 and RB1 inactivation accelerates OS tumorigenesis.	Lin et al, Carcinogenesis, 2009 Berman et al, PNAS, 2008; Walkley	
	Rb, P53	_	Conditional inactivation of	mice Crossing with Osx-Cre mice	-	Yes, OS	C57BL/6-J /129 Mixed C57BL/6-J /129	100%/ 41weeks	OS exhibited many of the features characteristic of human OS, including comparable histology, metastatic site preference, karyotypic complexity, and transcriptional profiles.	et al, Gen & Dev, 2008 Berman et al, PNAS, 2008; Walkley et al, Gen & Dev, 2008	
	Rb, P53 P53 ^{R172H/+}		Conditional inactivation of (P53 ^{flox/flox}) Conditional activation of	Crossing with Col1A1-3.6-Cremice		Yes, OS (20% incidence of lymphoma or fibrosarcoma) Yes, OS in 50% (also develop carcinomas,	Not reported	60%/42 weeks 50%/mean survival time 67 weeks	Mice specifically deleted for P53 in osteoblasts developosteosarcomas, wich correlates with increased proliferation, increased expression of RUNX2, increased osteoblast maturation as a result of P53 inactivation.	Lengner et al, The J ournal of Cell Biology, 2006 Lang et al, Cell, 2004; Olive et al,	
	P53	_	P53 ^{R172H/+} mutant Embryonic stem cell alteration, (Constitutive (germline)	53 locus;Prm-Cre (germline)	-	Yes, various entities, especially lymphoma and sarcoma (typically UPS, more rarely		4%/16–20weeks, 25%/39 weeks	mice, the most frequent tumors in the p53 ^{817391/+} mice were osteosarcomas. The unexpected ratio of tumor formation in homozygous vs heterzygous mice is due to the early lethality seen in the homozygous mice (rapid development and higher incidence of other tumors—mostly.)	Cell, 2004; Donehower et al, Nature, 1992; J acks et al, Current Biology, 1994;	
6. EwS/Ewing	A EwS-GEMM has bee	en elusive so far despit	te 16 attempts by 6 differ	ent laboratories to expres		also OS, RMS and Angiosarcoma) ous conditional and tissue–specif	ic strategies and u	sing a plethora of different promotors (Runx2,Col1a2.3,Col1a3.6,Prx1,CAG,Nse, NEFL, Dermo1,P0, Sox9,Cancer Research, 2017		
7. SySo/Synovial		Rosa26	Loxp-PGK-Neo-tPA-Loxp-SS18-		SS18-SSX2 is suficient to drive tumorigenesis in Myf5 positive		Mixed SvJ -129/		SS18-SSX1/2 allele avaiable with and without an IRES-GFP reporter.SS18-SSX1/2 allele can also be activated via a a tamoxifen-inducibleCreER system or by TAT-CREdelivery		
sarcoma	Nf1, P53		One construct with 2 linked	niaction/nairing of mica	progenitors. Accelarated by Pten inactivation or expression of a stabilized version of Beta-Catenin.		C57BL/6 129/SvJ	83%/14 weeks	(focal tumorigenesis albeit with longerlatency) Validation of CRISPR-technology to induce the same tumorigenesis of similar latency	Oncotarget, 2015;Barrott et al, The J ournal of Experimental medicine, 2016; Huang et al, Nature	
8. MPNST/ Malignant peri-	Nf1, P53 Nf1, Ink4a/Arf	for Cas9	sgRNAsvs Nf1 and P53 plus Cas9 packed into adenovirus Systemic Nf1 ^{flox/flox} & Ink4a/Arf ^{flox/flox} , spatially controlled inactivation by	the sciatic nerve Ad-Creinjection into the sciatic nerve or the thigh muscle	-	Yes,MPNST when injected into the sciatic nerve, 50% UPS & 50% RMS when injected	, -	100%/18weeks for MPNST; 100%/26	and molecular makeup as CreLoxP-approach. Conditional MPNST model with wildtype P53 background.	Communications, 2017 Dodd et al, Molecular Cancer	
pheral nerve sheath tumor	Nf1, Ink4a/Arf	-	Cre injection Nf1+/- + Ink4a/Arf-/-or P53+/-; Crossing of mice;	Constitutive (germline)	-	into the muscle (H.E. IHC) Yes, MPNST	C57BL/Ka	weeks for RMS & UPS; Variable, dependent on combination	The tumorigenic potential of MPNST cells with Nf1+/-and Ink4a/Arf-/-ishigher than Nf1+/-and P53+/-, tumorigenisity depends on laminin.	Therapeutics, 2013 Buchstaller et al, Cancer Cell, 2012	
	Nf1, P53 Nf1, P53	-	(conditional knockout)	Crossing with Wnt-1-Cre+/ Periostin-Cre+/ P0a-Cre+mice Constitutive (germline)	-	Yes, Neurofibroma, MPNST, others Yes, sarcoma, MPNST only	Mixed C57BL/6/129Sv Mixed	Variable, dependent on combination 100%/26 weeks for Nf1*/-and P53*/-(NP-CIS); 56	Heterozygosity of Nf1+'-andP53+'-onthe same chromoseome leads to faster MPNST formation than on	J oseph et al, Cancer Cell, 2008 Cichowski et al, Science, 1999	
	Nf1, P53			Constitutive (germline)	-	NP-Cis animals Yes,77% sarcoma(mostly MPNST,but also RMS, MTT,LMS), 14% lymphoma,8%	C57BL/6/129Sv Mixed	weeks for Nf1+/-and P53+/-(NP-Trans); 100%/17 weeks for Nf1-/-and P53+/-; 22 weeks for Nf1+/-and P53+/-;	opposite chromosomes.Chimeras for NT-deletionshow varying developmental defects and plexiform neurofibromas depending on degree of chimerism. Germ line homozygosity for Nf1 deletion leads to embryonic lethality.	Vogel et al, Science, 1999	
9. IFS/Infantile	No holistic GEM mode	eling attempt could be	identified. Nonetheless, b	ooth ETV6-NTRK3 and EM	IL4-NTRK3 tranform N	carcinoma, 3% neuroblastoma	C37BL/0/1293V		e). (Wai et al, Oncogene, 2000 & Tannenbaum–Dvir et al, Cold Sprin	g Harb. Mol. Case Studies, 2015)	
fibrosarcoma 10. MRT /	SmarcB1	-	Smarcb1 ^{flox/flox}	Crossing with Mx-Cre	P53flox/flox	Yes, MRT histology, negative for SNF5 expression. Mice also develop mature	Mixed	100%/11 weeks; 100%/ 3 weeks	Conditional inactivation of SmarcB1 results in highlypenetrant cancer predisposition with 100% of mice developing mature CD8+T cell lymphoma or rabdoid tumors. While	Roberst et al, Cancer Cell, 2002;	
Malignant	SmarcB1		SmarcB1+/-; SmarcB1-/-; C	Systemic inactivation)		expression. Mice also develop mature CD8+T cell lymphoma Yes, MRT, negative for SNF5	C57BL/6/129Sv Mixed	when p53 is co-inactivated 12%/5 weeks	loss of SNF5 predisposes to aggressive cancers, it is also required for survival of virtually all nonmalignant cells in vivo.	Isakoff et al, PNAS, 2005; Roberts et al, PNAS, 2000; Guidi et al, Mol	
				ensulative (germinie) PS		expression	C57BL/6/129Sv		predisposed to MRT. P33 (but not Curnzaor RD inactivation) accelerates turnor formation	Cell Biol, 2001; Klochendler-Yeivinet al, EMBO reports, 2000; DelBove et al, Molecular carcinogenesis, 2009;	
11. CSSK/ Clear cell sarcoma of the kidney	No modeling attemp	ts identified in the liter	ature.								
12.CSS/ Clear	EWSR1-ATF1	Rosa26	Conditional expression cassette	Local injection of Tat-Creinto limbs and various other	-	Yes,CSS resembling human disease	C57BL/6	100%/5weeks (100μM Tat-Cre) to 40	A variety of mouse cells are permissive to EWSR1-ATF1-transformation, embryonic expression is lethal, expression in young mice leads to stunted growth.	Straessler et al, Cancer Cell, 2013	
cell sarcoma of soft tissue	(human cDNA)		(loxP-neo-R+polyA-loxP-	Imbs and various other locations plus systemic Tamoxifen-mediatedactivation (Rosa26-CreER)				weeks (2μM Tat-Cre)	Commented to Stunded Growth.		
13.ASPS/ Alveolar soft part sarcoma	ASPSCR1-TFE3 human cDNA	Rosa26	Rosa26-LSL-AT3 crossed to systemic CreER	Systemic Tamoxifen–mediated activation (Rosa26–CreER)	-	Yes, tumors recapitulated human ASPS histopathologically and in expression analysis	Mixed 129S1/SvImJ and C57BL/6	100%/22 weeks	The completely penetrant tumors were indistinguishable from human ASPS by histology and gene expression. Tumors formed preferentially in an anatomic environment highest in lactate. No concurrent genetic hits were required for tumorigenesis.	Goodwin et al, Cancer Cell, 2014	
14. SBRCT-BCOR No modeling attempts identified in the literature. /Small Blue											
round cell tumor 15. SBRCT-CIC /Small Blue round cell tumor	und cell tumor . SBRCT-CIC No modeling attempts identified in the literature. mall Blue										
16. DSRCT/	No modeling attempts identified in the literature.										
Desmoplastic small round cell											
17. MC/ Mesenchymal	No modeling attemp	No modeling attempts identified in the literature.									
chondrosarcoma											
Inflammatory myofibroblastic tumor	No modeling attempts identified in the literature.										
	figure 1 Ev	dicting CEM	models for no	distric carear	no of 10 foot	us antitios For and	h ontitis (n	ublished CEM medals	are presented in chronological order.s		

Supplemental figure 1. Existing GEM models for pediatric sarcoma of 18 focus entities. For each entitiy published GEM models are presented in chronological order, starting with the most current approach. A color code was applied to highlight groups of models within recurrent genetic pathways/networks: Sonic Hedgehog signaling, RAS signaling, Hippo signaling/Muscle injury/Muscular dystrophy, HER-2/Neusignaling, HGF/HF signaling. P53, Rb1 and Cdkn2a inactivation was so commonly applied that it did not receive a specific color code. For further details on individual models, see text and original articles.