

Supplementary Materials

Characterization of mTOR Activity and Metabolic Profile in Pediatric Rhabdomyosarcoma

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Table S1. The relevant clinicopathological characteristics of the studied patients.

Case No.	Primary	After Preoperative Therapy	Relapse	Fusion Status	Histological Subtype	Risk Group	Gender
1	x			positive	alveolar	HR	male
2	x			negative	spindle cell	HR	male
3	x		x	negative	embryonal	HR	female
4	x			negative	embryonal	HR	female
5	x			negative	embryonal	HR	female
6		x		negative	embryonal	HR	male
7	x			negative	embryonal	HR	female
8	x			negative	embryonal	MR	female
9	x	x		negative	embryonal	HR	female
10	x		x	negative	embryonal	HR	female
11	x		x	negative	embryonal	HR	male
12	x			negative	embryonal	HR	male
13	x		x	negative	embryonal	HR	male
14	x		x	negative	embryonal	MR	female
15	x			negative	embryonal	MR	male
16	x			negative	embryonal	HR	male
17	x			negative	embryonal	MR	male
18	x			negative	embryonal	MR	male
19	x	x		negative	botryoid	HR	female
20	x			negative	embryonal	HR	female
21	x			negative	embryonal	SR	female
22	x			positive	alveolar	HR	female
23	x			negative	embryonal	MR	male
24	x			negative	embryonal	MR	male
25	x			negative	embryonal	HR	male
26	x			negative	embryonal	HR	male
27			x	negative	embryonal	HR	male
28	x		x	positive	alveolar	HR	male
29	x	x	x	negative	embryonal	HR	female
30	x	x		positive	alveolar	HR	female
31	x			negative	embryonal	MR	male
32	x			negative	embryonal	MR	male
33	x		x	negative	embryonal	HR	male
34	x			negative	embryonal	HR	female
35	x			negative	embryonal	HR	male
36	x			negative	spindle cell	HR	male
37	x	x	x	negative	embryonal	HR	female
38	x			negative	embryonal	HR	male
39	x	x		negative	embryonal	HR	female
40	x			negative	embryonal	MR	male
41	x			negative	embryonal	HR	male
42	x	x		positive	alveolar	HR	male
43	x	x		negative	embryonal	HR	male
44	x			negative	spindle cell	HR	female
45	x			negative	embryonal	HR	female
46	x			negative	embryonal	HR	male
47	x			negative	embryonal	MR	male
48			x	negative	embryonal	HR	female

Table S2. The mean and median H-score results and the belonging probability values of the primary and recurrent samples according to PAX-FOXO fusion status, risk stratification and survival. .

	Total no.	pmTOR			pS6			Rictor			PFK			LDHA			G6PDH			ATPB			GLS		
		mean	median	<i>p</i>	mean	median	<i>p</i>	mean	median	<i>p</i>	mean	median	<i>p</i>	mean	median	<i>p</i>	mean	median	<i>p</i>	mean	median	<i>p</i>	mean	median	<i>p</i>
Primary																									
Fusion negative	40	107	120	0.366	51	38	0.148	140	140	0.02	43	8	0.214	110	110	0.985	139	146	0.597	100	98	0.485	120	137	0.448
Fusion positive	5	137	143		16	13		189	193		79	103		104	80		125	124		99	63		101	108	
Low risk	12	133	153	0.095	49	58	0.418	149	143	0.668	30	3	0.185	132	149	0.126	142	151	0.626	65	54	0.626	122	129	0.626
High risk	33	101	115		46	20		145	150		53	18		100	94		135	133		83	75		116	135	
Survived	32	108	121	0.634	48	32	0.773	146	150	0.599	44	12	0.723	114	125	0.475	141	146	0.288	83	70	0.79	115	129	0.867
Progressed	13	115	122		44	30		138	135		54	10		99	78		124	124		78	95		124	135	
Recurrent																									
Fusion negative	9	170	178	0.099	73	76	0.637	111	105	0.068	73	58	0.458	138	148	0.906	86	82	0.813	88	84	0.239	127	143	0.637
Fusion positive	2	10	10		36	36		189	189		3	3		131	131		67	67		21	21		86	86	
Low risk	3	202	202	0.838	8	8	0.014	140	140	0.794	58	58	1	144	144	0.608	67	67	0.609	84	85	0.414	143	143	0.759
High risk	8	122	125		80	76		123	121		59	3		135	148		86	82		73	43		113	106	
Survived	3	90	90	0.133	80	80	0.776	179	179	0.4	60	60	0.63	115	115	0.497	108	108	0.776	83	83	0.497	155	155	0.921
Progressed	8	150	178		62	60		114	105		58	13		142	148		76	82		73	49		110	90	

Table S3. The immunopositivity and the statistical results of pmTOR, pS6, Rictor, PFKP, LDHA, G6PDH, ATPB and GLS stainings in the studied cases according to PAX-FOXO fusion status, risk stratification and survival. (H-score < 100 – negative (neg.), H-score > 100 – positive (pos.)).

	Total no.	pmTOR			pS6			Rictor			PFKP			LDHA			G6PDH			ATPB			GLS		
		Pos. (%)	Neg. (%)	<i>p</i>	Pos. (%)	Neg. (%)	<i>p</i>	Pos. (%)	Neg. (%)	<i>p</i>	Pos. (%)	Neg. (%)	<i>p</i>	Pos. (%)	Neg. (%)	<i>p</i>	Pos. (%)	Neg. (%)	<i>p</i>	Pos. (%)	Neg. (%)	<i>p</i>	Pos. (%)	Neg. (%)	<i>p</i>
Primary																									
Fusion negative	40	55	45	0.286	18	82	0.411	80	20	0.357	13	87	0.033	53	47	0.411	87	13	0.643	35	65	0.592	72	28	0.467
Fusion positive	5	80	20		0	100		100	0		60	40		40	60		80	20		40	60		80	20	
Low risk	12	75	25	0.142	0	100	0.094	92	8	0.302	8	92	0.302	67	33	0.26	92	8	0.302	25	75	0.3	83	17	0.223
High risk	33	52	48		21	79		79	21		21	79		48	52		79	21		79	21		39	71	
Survived	32	57	48	0.745	16	84	0.68	81	19	0.58	16	84	0.672	58	42	0.347	84	16	0.42	34	66	0.528	71	29	0.589
Progressed	13	62	38		15	85		85	15		23	77		46	54		77	23		38	62		69	31	
Recurrent																									
Fusion negative	9	78	22	0.109	11	89	0.818	50	50	0.333	33	67	0.509	89	11	0.345	22	78	0.491	56	44	0.273	56	44	0.273
Fusion positive	2	0	100		0	100		100	0		0	100		50	50		50	50		0	100		0	100	
Low risk	3	67	33	0.721	0	100	0.727	67	33	0.333	0	100	0.339	100	0	0.509	0	100	0.339	67	33	0.424	33	67	0.576
High risk	8	63	37		13	87		57	43		38	62		75	25		38	62		38	62		50	50	
Survived	3	33	67	0.279	0	100	0.727	100	0	0.333	33	67	0.661	67	33	0.491	33	67	0.661	67	33	0.424	67	33	0.424
Progressed	8	75	25		13	88		50	50		25	75		88	13		25	75		38	63		38	63	



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