

Supporting materials for

Nuclear spin hyperpolarization of pyruvate enables longitudinal monitoring of treatment response in intestinal tumor organoids

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Metabolic kinetic parameters of control and rapamycin-treated tumor organoids

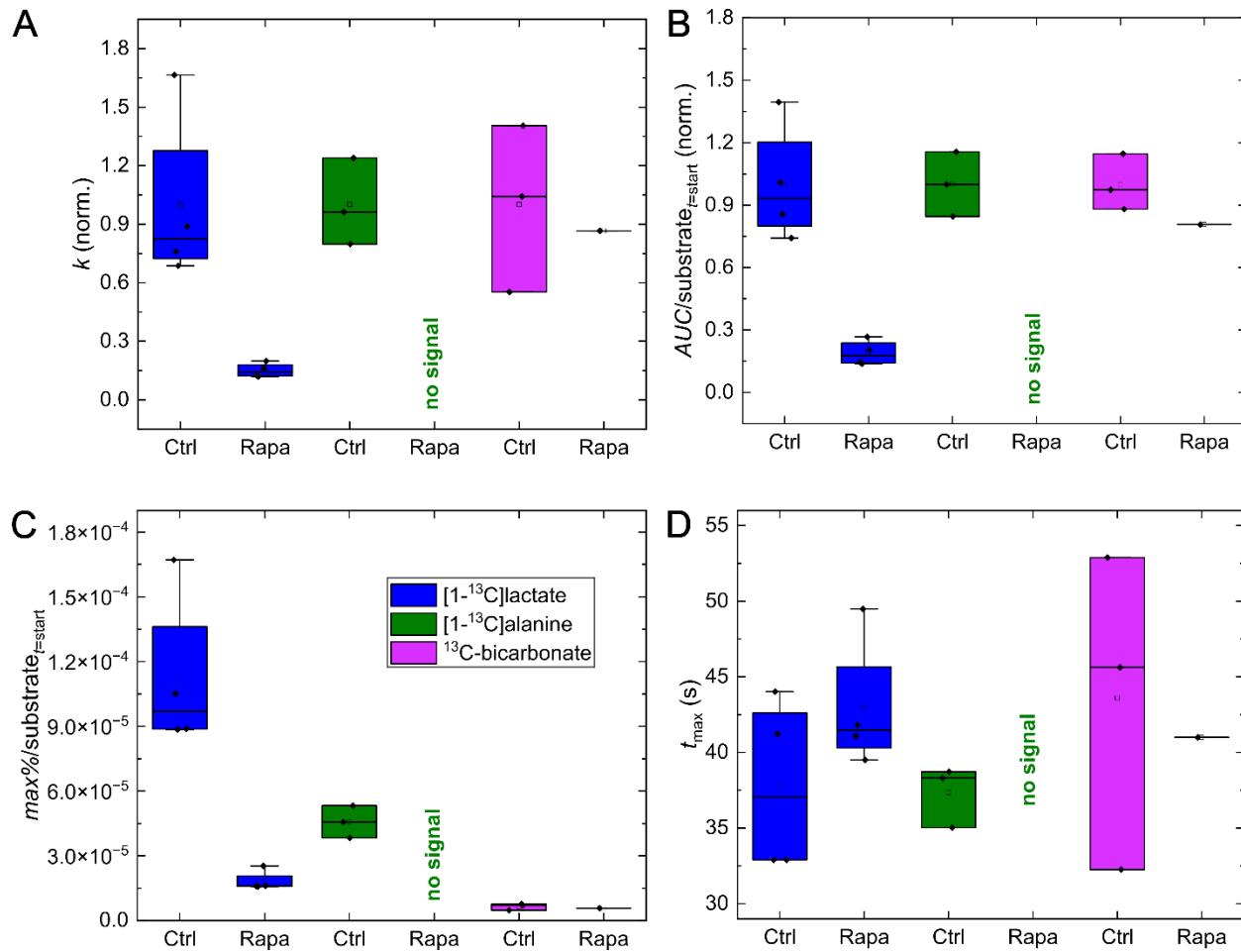


Figure S1: Strong metabolic effect of rapamycin treatment revealed by hyperpolarization enhanced NMR. Four datasets were evaluated in both cases (4 control, 4 treated with rapamycin). (A) The k -rate for lactate as anaerobe metabolism marker is significantly lower in the treated tumor organoids (6.6-times, effect size $d=2.7$). No alanine signal was observed in the treated tumor organoids suggesting that alanine experiences also about the same difference in signal or more between control and treated when compared to lactate, since the alanine in the treated group is hidden within the noise floor (SNR of alanine of about 5 in control group). Bicarbonate was only found in one instance for the treated organoids. This suggests a higher average metabolism for the control group in all cases. (B) The areas under kinetic curve (AUC, **Figure 2**) closely matches the results (treated 5.3-times lower, $d=3.9$) from A, as does maximum signal intensity (C, treated 6.2-times lower, $d=3.5$). The maximum bicarbonate signal was found to be just above the noise floor, making analysis less accurate. (D) A difference in t_{max} for lactate was observed between the treated and untreated organoids (5.2 seconds, $d=1.0$), though the difference is not significant. The big deviation between bicarbonate data points (SNR of only about 1.7) is not sufficient to generate quantitative results. The differences in A-C are significant to $p<0.005$ with and $p<0.05$ without equal variance assumed. The polarization of all experiments was $(27.0 \pm 5.7)\%$ after (29.4 ± 12.2) s following dissolution.

Lactate production rate in WT, tumor-control, and tumor-rapamycin-treated organoids

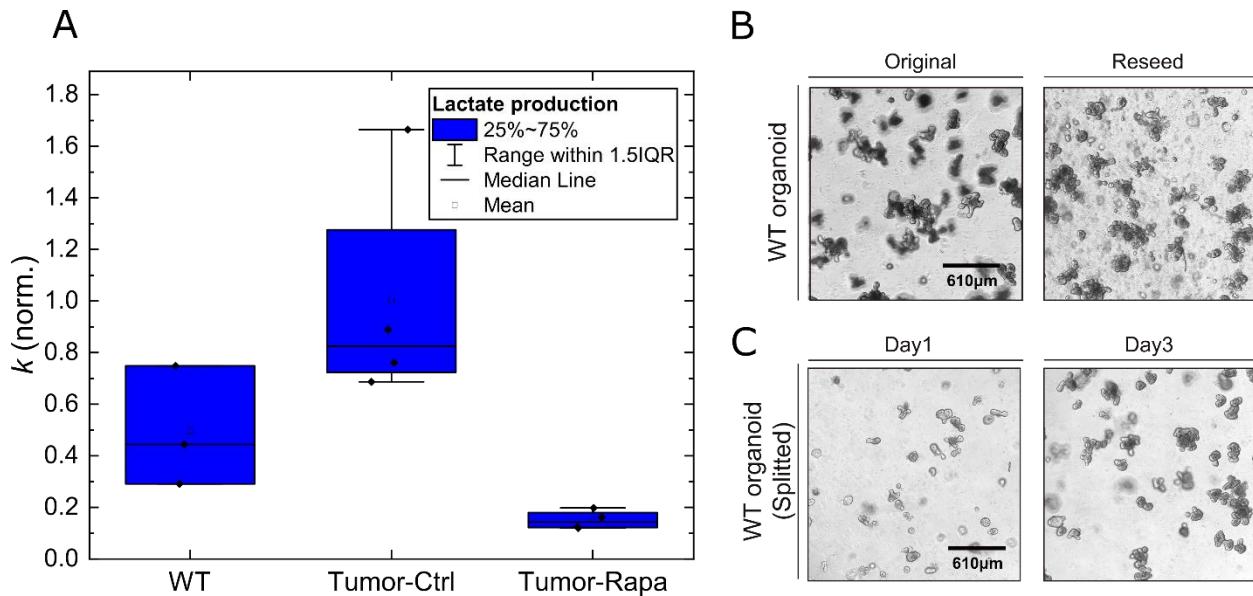


Figure S2. Comparing pyruvate to lactate conversion rate, k , in wild-type, tumor control, and tumor rapamycin-treated organoids and their morphology. (A) We compared the metabolism of tumor organoids to WT organoids. The WT organoids were enriched to increase the density of cells in the Matrigel. Without enriching, we were not able to readily observe their metabolism using hyperpolarization. The densities inside the Matrigel were 18.4% (WT), 34.8% (control), and 21.7% (stimulated). When comparing the conversion rates of Rapa and WT groups, the k of the WT organoids was higher despite its lower density. This indicates that the rapamycin treatment was efficiently suppressing metabolism below the level of the WT organoids, indicating the success of the treatment. Polarization across all experiments was $(28.6 \pm 6.0)\%$ and pH in the NMR tube was 7.4 ± 0.4 . (B) The WT organoids were reseeded after testing and representative pictures were taken on the next day of reseeding. Scale bar = $610\mu\text{m}$. The reseeded WT maintained its initial morphological characteristics. (C) After 2 days of reseeding, the WT organoids were passaged to demonstrate its sustainable culturability. Representative pictures were taken on Day 1 and Day 3, scale bar = $610\mu\text{m}$.

Phenotype behavior of organoids

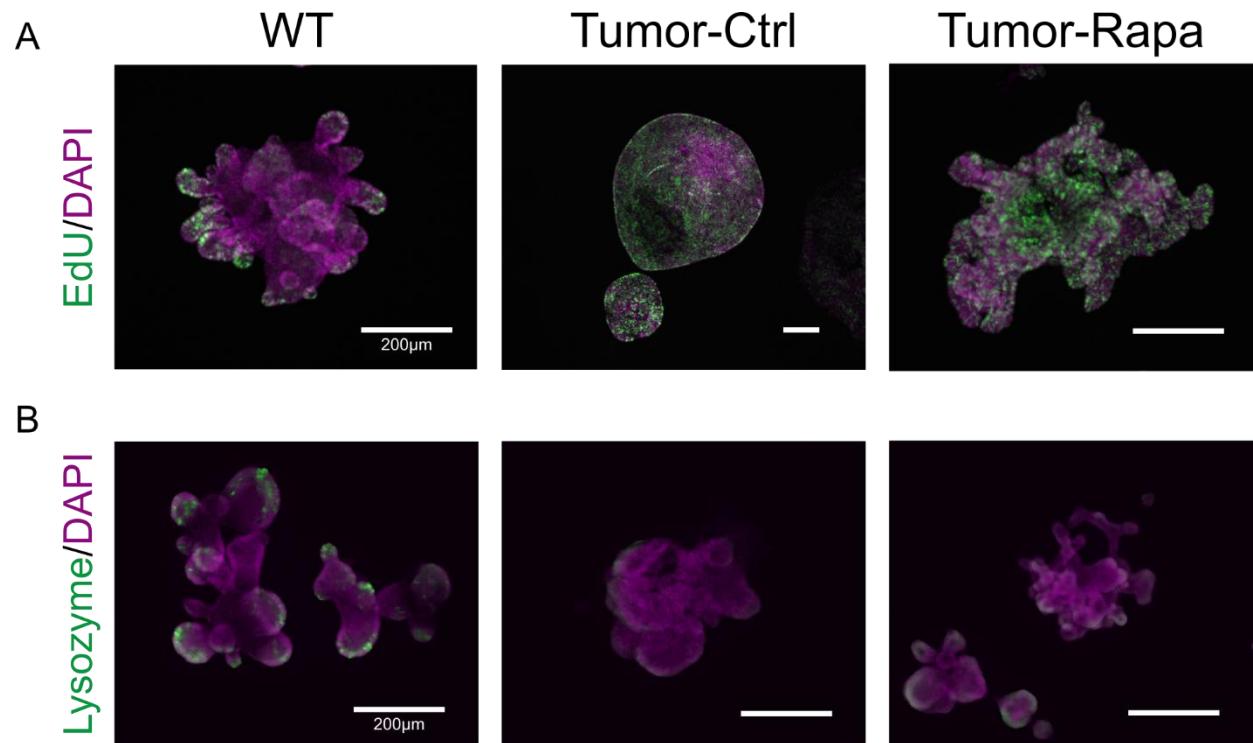


Figure S3. Rapamycin treated tumor organoid did not change tumor Sub-cell types. The sub-cell type of organoids was further characterized by fluorescent staining. (A) Stem cells were labeled via the EdU assay (see methods). In WT organoid, stem cells are confined to the crypt-like domain, whereas in both Ctrl and Rapa treated tumor groups, they are scattered throughout the organoid. (B) Paneth cells were labeled using lysozyme staining. There were Paneth cells apparent in WT, whereas no Paneth cells were found in both tumor-Ctrl and Rapa treatment groups. Scale bar = 200 μ m.

Estimated parameters of hyperpolarization and metabolic conversion

The tables below refer to the data presented in **Figure 2**, **Figure S1**, and **Figure S2**.

Table S1. All measured estimated parameters for conversion rate (k), area-under-the-curve (AUC/substrate $_{t=0}$), maximum signal intensity (max%), and time to reach the maximum signal intensity (t_{max}) of **Figure 2** in the main text and **Figures S1** and **S2** in SI. The sample pH value was measured inside the tube following the acquisition. The organoids coverage area was determined before the experiment. Polarization estimates of each sample were calculated from measured in parallel polarization at different system without administration to organoids (detailed in Table S2).

Sample	Coverage area (%)	Sample pH	Polarization estimates (%)	$k * 10^6$ (1/s)	AUC $*10^3$ /substrate $_{t=0}$	max% $*10^2$ (%)	t_{max} (s)
Control1	35.1	7.49	24.9	14.3	20.4	1.67	32.9
Control2	33.1	6.87	37.5	5.91	12.5	0.89	41.2
Control3	<i>not performed</i>	7.70	20.1	6.55	14.7	1.05	44.0
Control4	24.5	7.70	27.9	7.65	10.8	0.89	32.9
Rapa1	30.6	7.85	25.3	1.07	2.11	0.16	41.1
Rapa2	35.2	6.89	20.5	1.70	3.02	0.16	39.5
Rapa3	21.4	7.24	25.3	1.39	3.91	0.25	49.5
Rapa4	<i>not performed</i>	7.61	34.0	1.03	2.02	0.16	41.8
WT1	18.0	7.51	33.0	6.45	7.83	0.69	30.4
WT2	20.1	6.87	38.0	3.82	4.28	0.38	28.6
WT3	17.1	7.66	27.6	2.50	3.44	0.28	32.1

Table S2. Measured polarization (P_{SS}) and $T_{1,SS}$ on 1 T SpinSolve ^{13}C machine with transfer times to the SpinSolve (t_{SS}) and the Bruker 9.4 T machine (t_{Bruker}) after injection of the hyperpolarized solution to the cells) of Figure 2 in the main text and Figures S1 and S2 in SI. Using the $P_{Bruker} = P_{SS} * \exp\left(\frac{t_{SS} - t_{Bruker}}{T_{1,SS}}\right)$ equation, the polarization at the time of injection to the organoids was estimated.

Sample	Polarization SpinSolve (%)	T_1 SpinSolve (s)	Transfer SpinSolve (s)	Transfer Bruker (s)	Polarization estimates Bruker (%)
Control1	27.71	54.6	16.5	22.4	24.87
Control2	40.37	78.6	15.1	21.0	37.45
Control3	35.40	75.9	16.0	58.8	20.14
Control4	32.05	74.4	15.2	25.4	27.94
Rapa1	31.04	41.0	16.2	24.5	25.35
Rapa2	24.33	49.2	15	23.5	20.47
Rapa3	34.19	78.4	14.6	38.2	25.30
Rapa4	36.38	75.7	16.5	21.6	34.01
WT1	34.8	74.8	16	20.1	32.95
WT2	no acquisition	no acquisition		21.6	38.04
WT3	29.5	76.8	16	21.1	27.64

The tables below refer to the data presented in **Figure 3**.

Table S3. All measured estimated parameters for conversion rate (k), area-under-the-curve (AUC/substrate_{t=0}), maximum signal intensity (max%), and time to reach the maximum signal intensity (t_{max}) of Figure 3 in the main text. The sample pH value was measured inside the tube following the acquisition. The organoids coverage area was determined before the experiment. Polarization estimates of each sample were calculated from measured in parallel polarization at different system without administration to organoids (detailed in Table S2).

Sample	Coverage area (%)	Sample pH	Polarization estimates (%)	k *10 ⁶ (1/s)	AUC*10 ³ /substrate _{t=0}	max%*10 ² (%)	t _{max} (s)
Control1	33.7	7.42	38.5	10.9	14.0	1.16	30.4
Control1_reseed	36.4	7.6	37.5	6.5	8.8	0.72	31.7
Rapa1	19.8	7.38	31.1	4.3	3.3	0.34	22.4
Rapa1_reseed	37.7	7.74	36.6	3.8	5.8	0.45	32.9
Control2	38.5	7.74	33.0	10.5	16.4	1.29	34.4
Control2_reseed	30.6	7.39	35.7	17.7	16.2	1.46	24.4
Rapa2	19.9	none	33.9	3.5	5.3	0.43	34.7
Rapa2_reseed	9.4	7.99	38.3	1.24	2.1	0.15	34.3

Table S4. Measured polarization (P_{SS}) and T_{1,SS} on 1 T SpinSolve ¹³C machine with transfer times to the SpinSolve (t_{SS}) and the Bruker 9.4 T machine (t_{Bruker}) after injection of the hyperpolarized solution to the cells of Figure 3 in the main text. Using the $P_{\text{Bruker}} = P_{\text{SS}} * \exp\left(\frac{t_{\text{SS}} - t_{\text{Bruker}}}{T_{1,SS}}\right)$ equation, the polarization at the time of injection to the organoids was estimated.

Sample	Polarization SpinSolve (%)	T ₁ SpinSolve (s)	Transfer SpinSolve (s)	Transfer Bruker (s)	Polarization estimates Bruker (%)
Control1	39.9	75.7	17	19.7	38.48
Control1_reseed	37.6	76.7	21	21.2	37.49
Rapa1	33.6	75.9	17	22.9	31.11
Rapa1_reseed	38.2	73.3	21	20.1	36.61
Control2	35.0	76.4	17	21.3	33.04
Control2_reseed	37.7	76.8	15.5	19.6	35.68
Rapa2	34.0	75.4	24	24.3	33.91
Rapa2_reseed	40.1	77.6	16.5	20.0	38.32