Electronic supplementary information (ESI) for:

Multiplatform metabolomics approach for comprehensive analysis of GIST xenografts with various *KIT* mutations

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GC-MS parameters

Table S1. GC-MS parameters of the applied method.

Parameter	Value
Injector temperature	250 ºC
Detector transfer line temperature	320 ºC
Filament source temperature	230 ºC
Quadrupole temperature	150 ºC
Electron impact voltage	70 eV
Operation mode	scan
Scan speed	2 spectra/s
Mass range	50-600 <i>m/z</i>

LC-MS parameters

The detailed gradient programs for LC-MS analyses in RP and HILIC modes are presented in Table S2. MS parameters during LC-MS runs are presented in Table S3. The mass range for the detection of metabolic features was 61-1200 m/z in both RP and HILIC analyses.

Table S2. Gradient elution conditions during LC-MS analyses in RP and HILIC modes.

RP-LC-MS			HILIC-LC-MS		
Time [min]	Mobile phase A [%]	Mobile phase B [%]	Time [min]	Mobile phase A [%]	Mobile phase B [%]
0.0	18.0	82.0	0.0	5.0	95.0
30.0	4.0	96.0	22.0	45.0	55.0
38.0	4.0	96.0	23.0	45.0	55.0
38.5	0.0	100.0	23.5	5.0	95.0
40.5	0.0	100.0	29.0	5.0	95.0
42.0	18.0	82.0	<i></i>		
50.0	18.0	82.0	-		

Table S3. Mass s	pectrometry	parameters du	ring LC-MS analys	es.
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Parameter	RP	HILIC
Gas temperature	350 ºC	350 ºC
Gas flow	10 L/min	11 L/min
Nebulizer pressure	45 psi	45 psi
Fragmentor voltage	175 V	150 V
Skimmer voltage	65 V	65 V



Figure S1. Relative tumour volume evolution comparing the untreated controls with imatinib twice a day orally at 50 mg/kg in UZLX-GIST1 (A)⁴³, UZLX-GIST4 (B)⁴⁴, UZLX-GIST2 (C)⁴³ and UZLX-GIST9 (D)⁴⁵ patient-derived xenograft models.



Figure S2. PCA score plots for data obtained from GIST extracts GC-MS (A) analysis . RP-LC-TOF-MS analysis in positive (B) and negative (C) ionization modes. and HILIC-LC-TOF-MS analysis in positive (D) and negative (E) ionization modes. Black colour corresponds to quality control samples.



Figure S3. Representation of hierarchical cluster analysis of GIST samples analysed by GC-MS (A) analysis . RP-LC-TOF-MS analysis in positive (B) and negative (C) ionization modes. and HILIC-LC-TOF-MS analysis in positive (D) and negative (E) ionization modes. Imatinib-treated samples from GIST4 model are substantially distinct from the remaining samples.



Figure S4. Heatmaps of the signal intensity of metabolic features detected by RP-LC-TOF-MS analysis in positive (A) and negative (B) ionization modes, and HILIC-LC-TOF-MS analysis in positive (C) and negative (D) ionization modes.

Table S4. Statistically significant metabolites (with 5 lowest FDR values from each analytical technique) differentiating four GIST models used in the study. *p* and FDR values result from ANOVA test. Tukey's test indicates between which groups statistical significance was confirmed.

Metabolite	p value	FDR	Tukey's test	Technique
SM(18:0/16:1)	1.40E-23	3.52E-21	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST4	RP-LC-MS(+)
PE(18:0/16:1)	3.10E-19	1.74E-17	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST4	RP-LC-MS(+)
PC(10:0/20:0)	3.24E-18	1.21E-16	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST2	RP-LC-MS(+)
PE(16:1/22:2)	2.50E-14	3.89E-13	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST4	RP-LC-MS(+)
SM(18:1/22:0)	2.08E-13	2.50E-12	GIST2-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST4	RP-LC-MS(+)
PE(16:0/18:2)	8.69E-18	1.31E-15	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST4	RP-LC-MS(-)
PE(18:0/20:5)	2.01E-15	4.54E-14	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1	RP-LC-MS(-)
PC(14:0/18:1)	6.64E-15	1.25E-13	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST2	RP-LC-MS(-)
PE(18:0/22:4)	4.69E-14	6.04E-13	GIST2-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST2	RP-LC-MS(-)
PE(16:1/22:2)	6.42E-13	6.16E-12	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST2	RP-LC-MS(-)
2-Methylnicotinamide	6.66E-20	8.90E-18	GIST9-GIST1; GIST9-GIST2; GIST9-GIST4	HILIC-LC-MS(+)
PC(14:0/18:1)	5.77E-17	3.31E-15	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1	HILIC-LC-MS(+)
Adenosine	6.23E-09	4.16E-08	GIST4-GIST1; GIST4-GIST2; GIST9-GIST2	HILIC-LC-MS(+)
PC(18:1/0:0)	2.72E-08	1.49E-07	GIST2-GIST1; GIST4-GIST2; GIST9-GIST2	HILIC-LC-MS(+)
Hypoxanthine	3.20E-08	1.67E-07	GIST4-GIST1; GIST4-GIST2; GIST9-GIST4	HILIC-LC-MS(+)
Uridine/Pseudouridine	1.56E-16	4.63E-15	GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST2; GIST9-GIST4	HILIC-LC-MS(-)
Guanosine	6.94E-16	1.55E-14	GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST2; GIST9-GIST4	HILIC-LC-MS(-)
Taurine	6.55E-09	3.44E-08	GIST2-GIST1; GIST4-GIST2; GIST9-GIST2	HILIC-LC-MS(-)
Oxoglutaric acid	2.64E-05	5.76E-05	GIST2-GIST1; GIST9-GIST1	HILIC-LC-MS(-)
Myo-Inositol	5.81E-12	2.60E-10	GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST2; GIST9-GIST4	GC-MS
Uridine	7.97E-12	2.60E-10	GIST4-GIST1; GIST9-GIST1; GIST4-GIST2; GIST9-GIST2	GC-MS
L-Glutamic acid	7.31E-08	7.17E-07	GIST9-GIST1; GIST9-GIST2; GIST9-GIST4	GC-MS
L-Threonine	9.66E-08	8.60E-07	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1	GC-MS
L-Alanine	3.40E-07	2.56E-06	GIST2-GIST1; GIST4-GIST1; GIST9-GIST1	GC-MS



Figure S5. Box plots visualising the differences in metabolite signal intensity between four GIST models. Representative metabolites have been selected from Table S4 (generated in MetaboAnalyst 5.0).