## **Supplementary information**

**Supplementary table 1.** List of detected metabolites. Five metabolites (in bold) were assigned with high confidence by their accurate mass (MS1) and their MS/MS information. Eleven (\*) were assigned based on their accurate mass (MS1). HMDB and METLIN software were used for molecular assignments with a tolerance up to 1 ppm. Drugs and exogenous compounds were excluded during the search.

Assignment	Designation	Theoretical <i>m/z</i> value	Experimental m/z value	PPM error	Chemical Formula	MS/MS fragments
CMP*	[M-H] <sup>-</sup>	322.0446	322.0448	0.70	$C_9H_{14}N_3O_8P$	-
UMP*	[M-H] <sup>-</sup>	323.0286	323.0288	0.65	$C_9H_{13}N_2O_9P$	-
AMP*	[M-H₂O-H] <sup>-</sup>	328.0452	328.0458	1.70	$C_{10}H_{14}N_5O_7P$	-
АМР	[M-H] <sup>-</sup>	346.0558	346.0558	0.02	$C_{10}H_{14}N_5O_7P$	78.95 96.96 134.05 210.99
GMP*	[M-H] <sup>-</sup>	362.0507	362.0507	0.06	$C_{10}H_{14}N_5O_8P$	-
UDP*	[M-H₂O-H] <sup>-</sup>	384.9844	384.9848	1.15	$C_9H_{14}N_2O_{12}P_2$	-
UDP	[M-H] <sup>-</sup>	402.9949	402.9948	0.30	$C_9H_{14}N_2O_{12}P_2$	78.95 158.91
ADP*	[M-H <sub>2</sub> O-H] <sup>-</sup>	408.0116	408.0118	0.55	$C_{10}H_{15}N_5O_{10}P_2$	-
UDP*	[M+Na-2H] <sup>-</sup>	424.9769	424.9778	2.20	$C_9H_{14}N_2O_{12}P_2$	-
ADP	[M-H] <sup>-</sup>	426.0221	426.0228	1.50	$C_{10}H_{15}N_5O_{10}P_2$	78.95 96.96 134.04 158.92 290.96 328.05
GDP*	[M-H] <sup>-</sup>	442.0171	442.0170	0.12	$C_{10}H_{15}N_5O_{11}P_2$	-
ADP*	[M+Na-2H] <sup>-</sup>	448.0041	448.0048	1.60	$C_{10}H_{15}N_5O_{10}P_2$	-
UTP*	[M-H] <sup>-</sup>	482.9612	482.9620	1.66	$C_9H_{15}N_2O_{15}P_3$	-
АТР	[M-H] <sup>-</sup>	505.9885	505.9888	0.65	$C_{10}H_{16}N_5O_{13}P_3$	78.95 96.96 134.04 158.92
GTP*	[M-H] <sup>-</sup>	521.9834	521.9840	1.15	$C_{10}H_{16}N_5O_{14}P_3$	-
ATP*	[M+Na-2H] <sup>-</sup>	527.9704	527.9708	0.73	$C_{10}H_{16}N_5O_{13}P_3$	-
ADP-Ribose	[M-H <sub>2</sub> O-H] <sup>-</sup>	540.0538	540.0538	0.06	$C_{15}H_{23}N_5O_{14}P_2$	158.91 408.01
Sodium ATP*	[M+Na-2H]	549.9524	549.9528	0.80	$C_{10}H_{15}N_5NaO_{13}P_3$	-
UDP-Glucose/Galactose*	[M-H] <sup>-</sup>	565.0477	565.0484	1.20	$C_{15}H_{24}N_2O_{17}P_2$	-
UDP-N-Acetyl Glucosamine/Galactosamine*	[M-H] <sup>-</sup>	606.0743	606.0754	1.80	C <sub>17</sub> H <sub>27</sub> N <sub>3</sub> O <sub>17</sub> P <sub>2</sub>	-

Supplementary table 2. List of TM-SPRAYER<sup>™</sup> settings. According to the experiments performed in step 1, the matrix concentration, the temperature, the number of layers and the flow rate were defined as non-significant parameters. Only the percentage of solvent used to dissolve the matrix was defined as a significant setting. Drying time was added to step 2 to confirm the non-significance of this setting.

			STEP 1		STEP 2		
TM-SPRAYER™ Settings	Unit	Lower value	Center value	High value	Lower value	Center value	High value
Matrix Concentration	mg/mL	7	8.5	10	fixed value: 10		
Solvent	%	60	65	70	50	75	100
Temperature	°C	65	72.5	80	fixed value: 80		
Number of layers	N	4	5	6	fixed value: 10		
Flow rate	mL/min	0.12	0.135	0.15	fixed value: 0.1		
Drying time	Sec	22	26	30	2	31	60

**Supplementary figure 1. Validation of the experimental design model**. Intensities of ATP, ADP and UDP were normalized by root mean square (RMS) for each experiment performed in duplicate.



Supplementary figure 2. Spatial distribution of GTP and UTP.

