

## Supporting Information

### SI.1

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#### SI.1 FAs monitored in plasma

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	Ions pair		Retention time/min	
	Normal tagged	Heavy tagged	Normal tagged	Heavy tagged
<b>AT1</b>	548.0 -> 254.0	568.0 -> 274.0	7.72	7.41
<b>AT2</b>	546.0 -> 254.0	566.0 -> 274.0	6.91	6.64
<b>AT3</b>	542.0 -> 254.0	562.0 -> 274.0	5.66	5.37
<b>AT4</b>	540.0 -> 254.0	560.0 -> 274.0	5.37	5.08
<b>AT5</b>	540.0 -> 254.0	560.0 -> 274.0	5.97	5.75
<b>AT6</b>	538.0 -> 254.0	558.0 -> 274.0	4.69	4.39
<b>AT7</b>	520.0 -> 254.0	540.0 -> 274.0	6.67	6.39
<b>AT8</b>	518.0 -> 254.0	538.0 -> 274.0	5.94	5.65
<b>AT9</b>	518.0 -> 254.0	538.0 -> 274.0	6.66	6.41
<b>AT10</b>	516.0 -> 254.0	536.0 -> 274.0	5.204	4.92
<b>AT11</b>	516.0 -> 254.0	536.0 -> 274.0	6.05	5.82
<b>AT12</b>	514.0 -> 254.0	534.0 -> 274.0	4.504	4.20
<b>AT13</b>	512.0 -> 254.0	532.0 -> 274.0	3.87	3.40
<b>AT14</b>	506.0 -> 254.0	528.0 -> 274.0	6.13	5.864
<b>AT15</b>	492.0 -> 254.0	512.0 -> 274.0	5.60	5.322
<b>AT16</b>	492.0 -> 254.0	512.0 -> 274.0	6.43	6.18
<b>AT17</b>	490.0 -> 254.0	510.0 -> 274.0	4.80	4.49
<b>AT18</b>	478.0 -> 254.0	498.0 -> 274.0	5.03	4.69
<b>AT19</b>	464.0 -> 254.0	484.0 -> 274.0	4.36	4.00
<b>AT20</b>	458.0 -> 254.0	478.0 -> 274.0	2.47	2.056
<b>AT21</b>	436.0 -> 254.0	456.0 -> 274.0	2.4	2.07
<b>BT1</b>	684.0 -> 254.0	704.0 -> 274.0	2.58	2.204

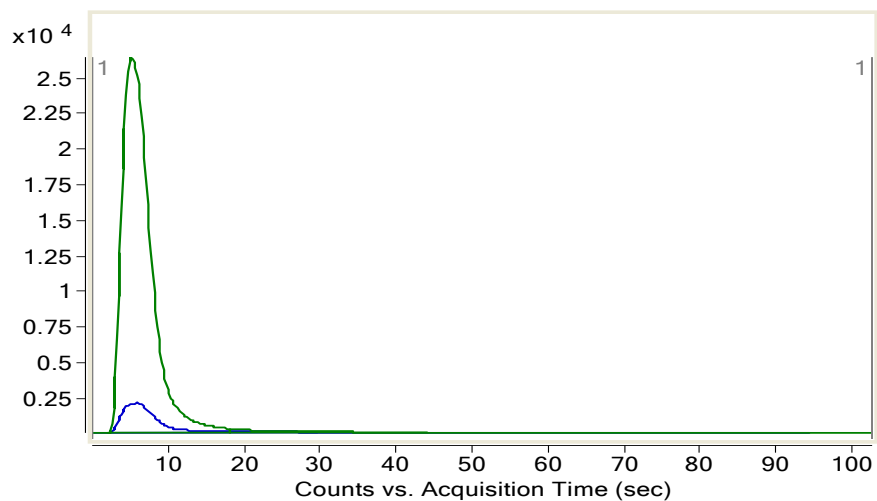
<b>BT2</b>	630.0 -> 254.0	650.0 -> 274.0	5.146	4.803
<b>BT3</b>	602.0 -> 254.0	622.0 -> 274.0	8.75	8.54
<b>BT4</b>	600.0 -> 254.0	620.0 -> 274.0	3.54	2.98
<b>BT5</b>	592.0 -> 254.0	612.0 -> 274.0	5.71	5.4
<b>BT6</b>	590.0 -> 254.0	610.0 -> 274.0	4.97	4.68
<b>BT7</b>	588.0 -> 254.0	608.0 -> 274.0	4.12	3.76
<b>BT8</b>	576.0 -> 254.0	596.0 -> 274.0	8.63	8.33
<b>BT9</b>	574.0 -> 254.0	594.0 -> 274.0	7.88	7.56
<b>BT10</b>	568.0 -> 254.0	588.0 -> 274.0	6.08	5.79
<b>BT11</b>	566.0 -> 254.0	586.0 -> 274.0	5.40	5.07
<b>BT12</b>	566.0 -> 254.0	586.0 -> 274.0	5.962	5.73
<b>BT13</b>	564.0 -> 254.0	584.0 -> 274.0	4.53	4.21
<b>BT14</b>	544.0 -> 254.0	564.0 -> 274.0	6.20	5.92
<b>IS1</b>	454.0 -> 254.0		3.223	
<b>IS2</b>	342.0 -> 324.0		7.22	

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## SI.2 Derivatization yield

As shown in bellowing, 1mg/mL PA was analyzed before and after derivatizaion.

More than 90% of PA were converted to hydrazides.



EIC (extracted ions chromatogram) of PA. 1mg/mL PA before derivatization (green line) and after derivatization (blue line) were analyzed with HPLC-MS. After derivatization, less than 10% PA were left.

**SI.3 LOD, dynamic range, linearity, and reproducibility for individual Fatty acids standards**

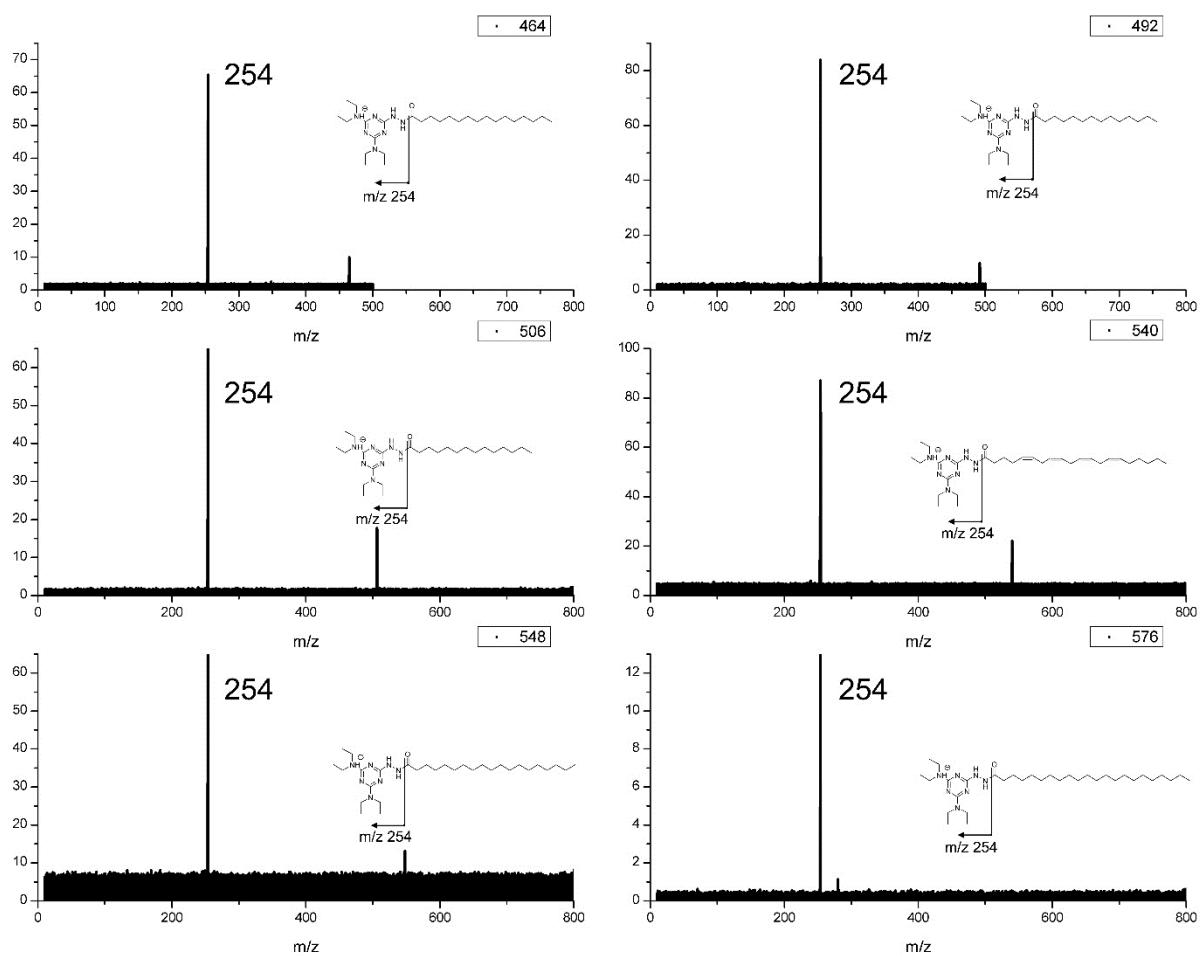
**SI.3 LOD, dynamic range, linearity, and reproducibility for individual Fatty acids standards**

	<b>LOD</b>	<b>Dynamic range</b>	<b>R</b>	<b>Reproducibility</b>
	<b>ng/mL</b>	<b>ng/mL</b>		<b>RSD</b>
<b>MA</b>	$1.00 \times 10^{-1}$	$2.00 \times 10^{-1} - 5.00 \times 10^3$	0.9998	4.89
<b>PA</b>	1.00	$5.00 - 5 \times 10^5$	0.9990	10.7
<b>HA</b>	$2.00 \times 10^{-1}$	$5.00 \times 10^{-1} - 5.00 \times 10^3$	0.9991	6.31
<b>AA</b>	$5.00 \times 10^{-1}$	$1.00 - 5.00 \times 10^3$	0.9996	2.00
<b>ARA</b>	$5.00 \times 10^{-1}$	$1.00 - 5.00 \times 10^4$	0.9991	4.05
<b>BA</b>	$1.00 \times 10^{-2}$	$1.00 \times 10^{-1} - 5.00 \times 10^3$	0.9995	3.00

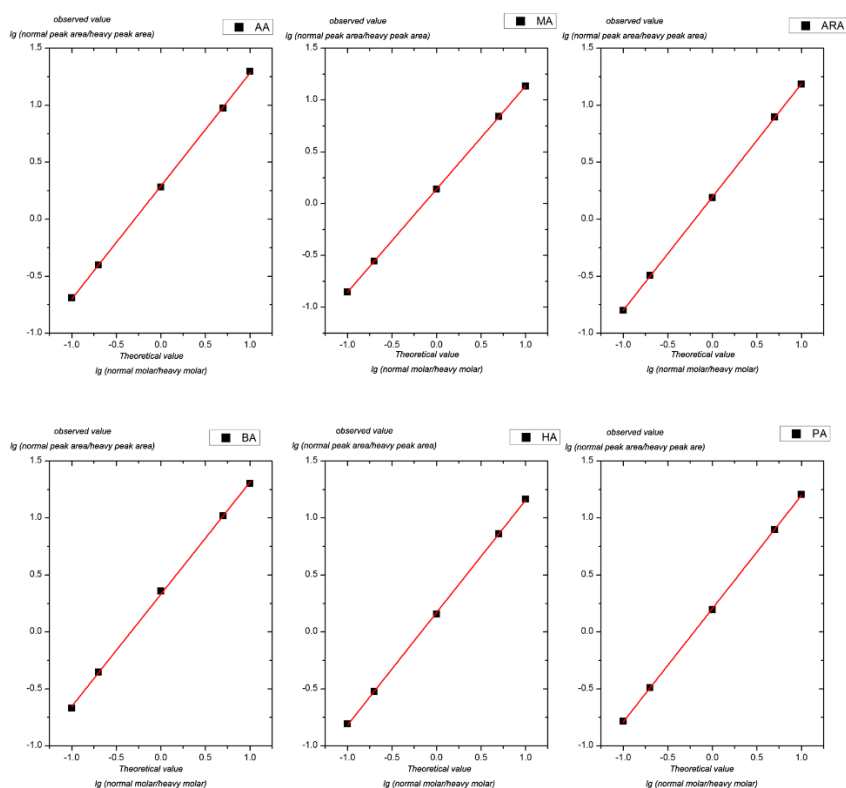
The reproducibility of each FA were calculated under the concentration of 500ng/mL (n=6).

## SI.4 MS/MS spectra of different FAs.

The 254 m/z fragments were observed in the MS/MS spectra of six different FA derivatives. These fragments are generated through the fragmentation process, as shown below.



## SI.5 Relative quantification analysis



Relative quantification based on ratios of “normal” and “heavy” tagged FAs showed good linear relationship. This indicated this strategy showed good agreement with the expected ratio.

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### Quantification results comparison between ISTD and ESTD

ID	Peak Area	Linear relationship	ESTD	ISTD	Ratio
			ng/mL	ng/mL	%
1	63738.32	$y=40.7206x+1008.038$	$1.54 \times 10^3$	$1.50 \times 10^3$	97.4
2	128685.9	$R=0.9999$	$3.14 \times 10^3$	$3.00 \times 10^3$	95.7

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The comparison was carried out with BA. The quantification results of ISTD is close to ESTD. This indicated the relative quantification is comparable with the classical ESTD method.

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**SI.6 Statistical analysis of FAs in plasma**

Compound	OPLS-DA analysis		
	VIP(>1)	Jack-knife(>0)	Pcorr(>0.58)
<b>AT10**</b>	1.15047	0.1705981	0.872339
<b>AT12**</b>	1.13208	0.1399155	0.849407
<b>AT13**</b>	1.14578	0.1524128	0.85804
<b>AT15***</b>	1.22366	0.1317099	0.890092
<b>AT16*</b>	1.21917	0.096167	0.897197
<b>AT17**</b>	1.10697	0.0943761	0.798911
<b>AT19****</b>	1.30934	0.1742674	0.966637
<b>AT2**</b>	1.18897	0.1585336	0.867255
<b>AT21***</b>	1.27173	0.1615318	0.905519
<b>AT3***</b>	1.22503	0.1493259	0.907871
<b>AT4***</b>	1.16241	0.1264132	0.887541
<b>AT8**</b>	1.10556	0.1251008	0.827057
<b>BT1</b>	1.07861	0.047673	0.790033
<b>BT11**</b>	1.1523	0.1503661	0.854977
<b>BT13***</b>	1.21637	0.1417778	0.889454
<b>BT3**</b>	1.05307	0.1111463	0.796992
<b>BT6**</b>	1.04131	0.0941234	0.771279
<b>BT7**</b>	1.09343	0.1510907	0.82858
<b>BT9****</b>	1.26696	0.1640355	0.931002

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The number of \* indicates the significances between aged and young groups

\*:p < 0. 05

\*\* :p < 0. 01

\*\*\*:p < 0. 005

\*\*\*\*:p < 0. 001