

Original Article

# Fragmentation of Dicarboxylic and Tricarboxylic Acids in the Krebs Cycle Using GC-EI-MS and GC-EI-MS/MS

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Isotope labeling measurements using mass spectrometry can provide informative insights on the metabolic systems of various organisms. The detailed identification of carbon positions included in the fragment ions of dicarboxylic and tricarboxylic acids in central carbon metabolism is needed for precise interpretation of the metabolic states. In this study, fragment ions containing the carbon backbone cleavage of dicarboxylic and tricarboxylic in the Krebs cycle were investigated by using gas chromatography (GC)-electron ionization (EI)-MS and GC-EI-MS/MS. The positions of decarboxylation in the dicarboxylic and tricarboxylic acids were successfully identified by analyses using position-specific <sup>13</sup>C-labeled standards prepared by *in vitro* enzymatic reactions. For example, carboxyl groups of C1 and C6 of trimethylsilyl (TMS)- and *tert*-butyldimethylsilyl (TBDMS)-derivatized malic and citric acids were primarily cleaved by EI. MS/MS analyses were also performed, and fragment ions of TBDMS-citric and  $\alpha$ -ketoglutaric acids ( $\alpha$ KG) with the loss of two carboxyl groups in collision-induced dissociation (CID) were observed.



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**Keywords:** organic acids, fragmentation, GC-MS, electron ionization, collision-induced dissociation

## INTRODUCTION

Isotope labeling experiments using mass spectrometry (MS) are one of the powerful methods to study metabolic systems. In particular, <sup>13</sup>C-labeling of dicarboxylic and tricarboxylic acids in the Krebs cycle is measured in the fields of biotechnology, systems biology, and medical sciences.<sup>1-4</sup> Since the interpretation of <sup>13</sup>C-labeling data requires positional information about the carbon atoms in the fragment ions, identification of carbon atoms contained in the fragment ions as observed by MS and tandem-MS has been investigated.<sup>5-10</sup> However, the carbon positions in the decarboxylated fragment ions derived from dicarboxylic and tricarboxylic acids have not been experimentally validated, resulting in the loss of rich information. In this study, the electron ionization (EI)- and collision-induced dissociation (CID)-fragmentation of the abundant dicar-

boxylic and tricarboxylic acids in the Krebs cycle, *viz.* citric,  $\alpha$ KG, succinic, fumaric, and malic acids was investigated to maximize the accessible data from a single analysis. The organic acid-derived carbon atoms included in the fragment ions were successfully identified by the analyses of position-specific <sup>13</sup>C-labeled standards synthesized by *in vitro* enzymatic reactions.

## EXPERIMENTAL

### Chemicals

Non-labeled standards were purchased from Sigma-Aldrich (St. Louis, MO, USA). [1,2,3,4-<sup>13</sup>C] $\alpha$ -Ketoglutaric acid (99%), NaH<sup>13</sup>CO<sub>3</sub> (99%), [1-<sup>13</sup>C]pyruvic acid (99%), [2-<sup>13</sup>C]pyruvic acid (99%), [3-<sup>13</sup>C]pyruvic acid (99%), and [1-<sup>13</sup>C]acetic acid (99%) were purchased from Cambridge Isotope Laboratories (Andover, MA, USA). Fully <sup>13</sup>C-labeled

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organic acids were prepared from the extract of yeast cultured in a medium containing [U-<sup>13</sup>C]glucose as the sole carbon source following the previously described method.<sup>11)</sup>

### Preparation of position-specific <sup>13</sup>C-labeled standards

*In vitro* enzymatic reaction was used for the synthesis of position-specific <sup>13</sup>C-labeled standards as described previously with minor modification.<sup>12)</sup> [4-<sup>13</sup>C]Malic acid was prepared *via* the reactions of phosphoenolpyruvate (PEP) carboxylase and malate dehydrogenase. Sixty-six millimolar *Tris*-HCl (pH=9), 10 mM PEP, 10 mM MgCl<sub>2</sub>, 20 mM NaH<sup>13</sup>CO<sub>3</sub>, 0.15 mM NADH, 4 units of PEP carboxylase from *Zea mais* leaves (Wako Pure Chemical Industries, Ltd. Corporation, Osaka, Japan), and 2 units of malate dehydrogenase from porcine heart (Wako Pure Chemical Industries, Ltd. Corporation) were gently mixed and incubated overnight at room temperature (Fig. S1a). [1-<sup>13</sup>C], [2-<sup>13</sup>C], and [3-<sup>13</sup>C]Malic acids were prepared *via* the sequential reactions of pyruvate kinase, PEP carboxylase, and malate dehydrogenase. Hundred millimolar *Tris*-HCl (pH=8), 10 mM ATP, 10 mM <sup>13</sup>C-labeled pyruvic acid, 15 mM MgCl<sub>2</sub>, 20 mM NaHCO<sub>3</sub>, 10 mM NADH, 16 U of pyruvate kinase from rabbit muscle *ca.* 350 U/mg protein suspension (Wako Pure Chemical Industries, Ltd. Corporation), 4 U of PEP carboxylase from *Zea mais* leaves, and 2 U of malate dehydrogenase from porcine heart (Fig. S1a) were gently mixed and incubated overnight at room temperature. [1-<sup>13</sup>C], [2-<sup>13</sup>C], and [3-<sup>13</sup>C]Pyruvic acids were used for the synthesis of [1-<sup>13</sup>C], [5-<sup>13</sup>C], [6-<sup>13</sup>C], and [1,5-<sup>13</sup>C]Citric acids were prepared *via* the multiple reactions of pyruvate kinase, PEP carboxylase, acetyl-CoA synthetase (ACS), and citrate synthase (CS). Hundred millimolar *Tris*-HCl (pH=8), 10 mM pyruvic acid, 20 mM NaHCO<sub>3</sub>, 10 mM MgCl<sub>2</sub>, 10 mM ATP, 10 mM acetate, 10 mM CoA, 16 U of pyruvate kinase, 4 U of PEP carboxylase from *Zea mais* leaves, 2 U of citrate synthase, and 0.2 U of acetyl-CoA synthase were gently mixed and incubated overnight at room temperature. ACS and CS were obtained from the content of F-kit acetate (J.K. International, Tokyo, Japan) (Fig. S1b). Acetic acid, NaHCO<sub>3</sub>, and pyruvic acid were replaced with [1-<sup>13</sup>C]acetic acid, [1-<sup>13</sup>C]pyruvic acid, and NaH<sup>13</sup>CO<sub>3</sub> for the synthesis of [1-<sup>13</sup>C], [5-<sup>13</sup>C], and [6-<sup>13</sup>C]citric acids, respectively.

### Derivatization of organic acids

Standard solution of organic acids was evaporated to dryness by Speed Vac (Thermo Fischer Scientific, Waltham, MA, USA). The dried standards were derivatized by adding 50 μL of 40 mg/mL methoxyamine hydrochloride pyridine solution and incubated for 1 h at 30°C. Subsequently 50 μL of *N*-methyl-*N*-trimethylsilyltrifluoroacetamide containing 1% 2,2,2-trifluoro-*N*-methyl-*N*-(trimethylsilyl)-acetamide, chlorotrimethylsilane (Thermo Fischer Scientific) or *N*-(*tert*-butyldimethylsilyl)-*N*-methyltrifluoroacetamide containing 1% *tert*-butyldimethylchlorosilane (Thermo Fischer Scientific) was added and the mixture was incubated for 1 h at 37 or 95°C for trimethylsilylation or *tert*-butyldimethylsilylation, respectively.<sup>8)</sup> After 1 h of cooling, an aliquot of the supernatant was subjected to the analysis.

### GC-MS and GC-MS/MS analysis

GC-MS (GCMS-QP2010 Ultra, Shimadzu, Kyoto, Japan) and GC-MS/MS (GCMS-TQ8040, Shimadzu) equipped with DB-5MS+DG column ((30 m × 0.25 mm ID × 0.25 μm), Agilent Technologies, Santa Clara, CA, USA) were used.<sup>8)</sup> Analysis conditions were as follows: constant flow rate of helium at 1.0 mL/min; ion source temperature, 230°C; electron impact ionization, 70 eV; injection volume, 1 μL; injection, pulsed split (split ratio, 1:10); oven temperature, 60°C for 3.5 min, increased at a rate of 10°C/min to 325°C, and maintained at that temperature for 10 min or 70°C for 2 min, increased at a rate of 3°C/min to 280°C,

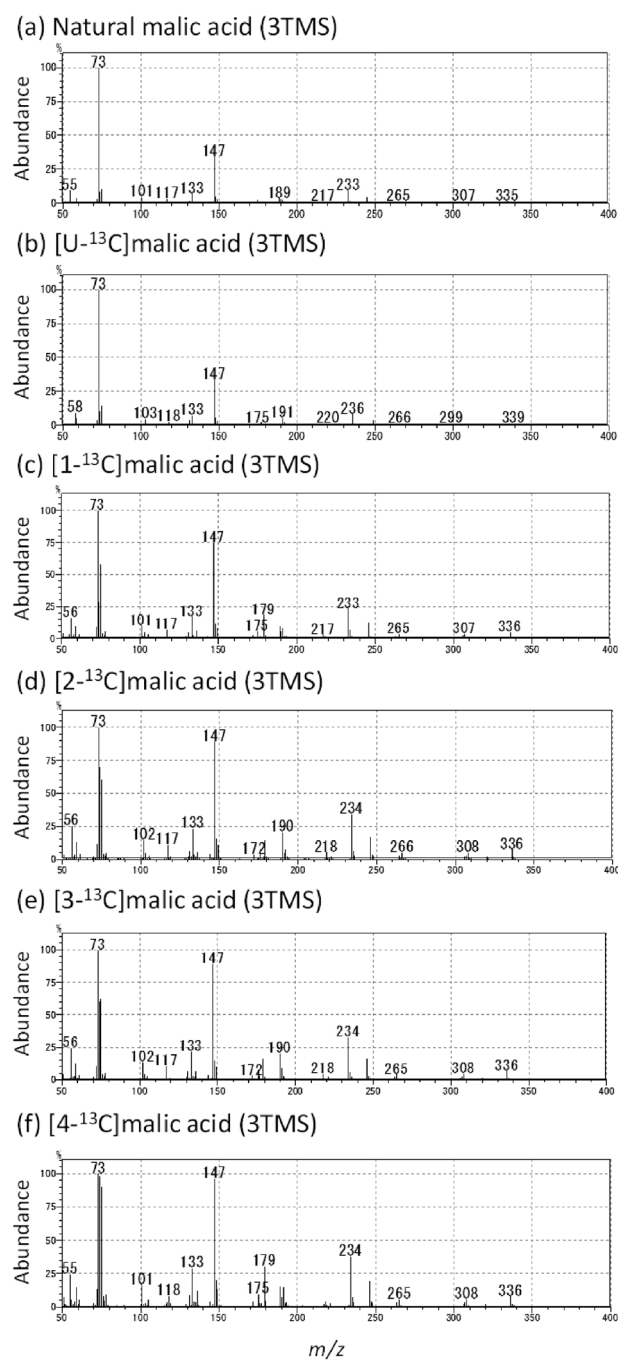


Fig. 1. EI-induced fragmentation of TMS-derivatized malic acid.

(a) natural, (b) [U-<sup>13</sup>C], (c) [1-<sup>13</sup>C], (d) [2-<sup>13</sup>C], (e) [3-<sup>13</sup>C], and (f) [4-<sup>13</sup>C]malic acid.

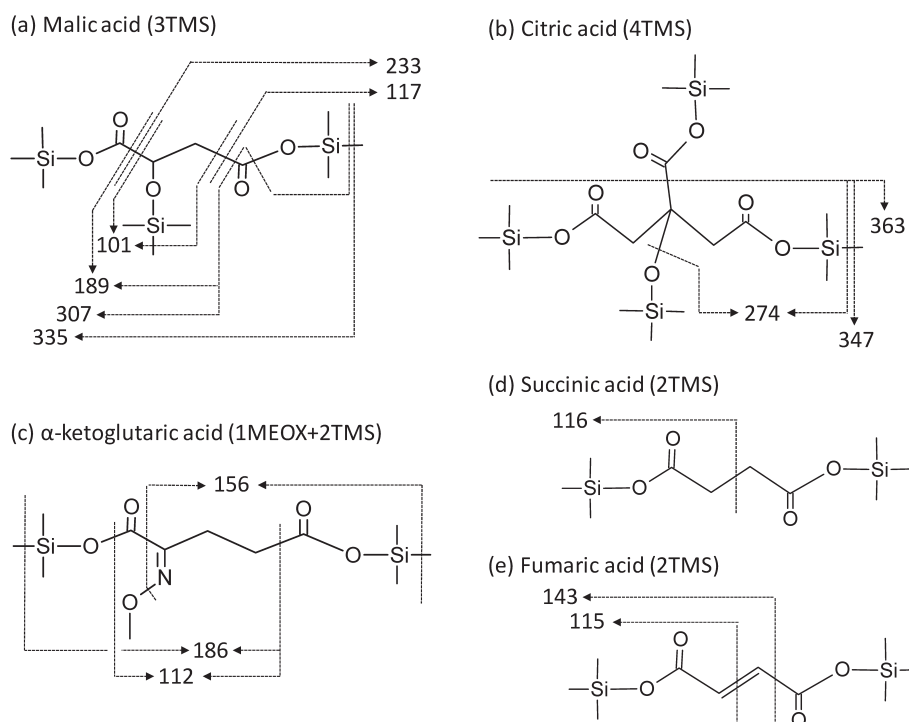


Fig. 2. Estimated EI-fragmentation of TMS-derivatized organic acids.

(a) malic, (b) citric, (c)  $\alpha$ KG, (d) succinic, and (e) fumaric acids. The position of demethylation in TMS group was not determined uniquely.

Table 1. Fragment ions with C-C bond cleavage of TMS- and TBDMS-derivatized organic acids by EI.

Metabolites	<i>m/z</i>	Number of organic acid-derived carbons	Carbon skeleton	Estimated chemical formula	Estimated cleavage group
<i>TMS derivatization</i>					
Citric acid (4TMS)	273	5	C1-2-3-4-5	$C_{11}H_{21}O_4Si_2$	TMS-COO and TMS-OH
Citric acid (4TMS)	347	5	C1-2-3-4-5	$C_{13}H_{27}O_5Si_3$	TMS-COO and $CH_3$
Citric acid (4TMS)	363	5	C1-2-3-4-5	$C_{14}H_{31}O_5Si_3$	TMS-COO
$\alpha$ KG (1MEOX, 2TMS)	112	4	C1-2-3-4	$C_5H_6NO_2$	TMS-COO and TMS-O
$\alpha$ KG (1MEOX, 2TMS)	156	4	C2-3-4-5	$C_6H_{10}NO_2Si$	TMS-COO, $CH_3$ and $CH_3O$
$\alpha$ KG (1MEOX, 2TMS)	186	4	C1-2-3-4	$C_7H_{12}NO_3Si$	TMS-COO and $CH_3$
Succinic acid (2TMS)	116	2	C1-2 or C3-4	$C_4H_8O_2Si$	TMS-COO- $CH_2$
Fumaric acid (2TMS)	115	2	C1-2 or C3-4	$C_4H_7O_2Si$	TMS-COO-CH
Fumaric acid (2TMS)	143	3	C1-2-3 or C2-3-4	$C_6H_{11}O_2Si$	TMS-COO
Malic acid (3TMS)	101	2	C2-3	$C_4H_9OSi$	TMS-COO, TMS-COO and $CH_3$
Malic acid (3TMS)	117	1	C4	$C_4H_9O_2Si$	Except TMS-COO
Malic acid (3TMS)	189	2	C2-3	$C_7H_{17}O_2Si_2$	TMS-COO, $CH_3$ and CO
Malic acid (3TMS)	233	3	C2-3-4	$C_9H_{21}O_3Si_2$	TMS-COO
Malic acid (3TMS)	265	1	C2	$C_9H_{25}O_3Si_3$	Unknown
Malic acid (3TMS)	307	3	C2-3-4	$C_{11}H_{27}O_4Si_3$	CO and $CH_3$
<i>TBDMS derivatization</i>					
Citric acid (4TBDMS)	431	5	C1-2-3-4-5	$C_{19}H_{39}O_5Si_3$	TBDMS-OH, TB and CO
Citric acid (4TBDMS)	357	5	C1-2-3-4-5	$C_{17}H_{33}O_4Si_2$	TBDMS-OH and TBDMS-CO
Citric acid (4TBDMS)	299	5	C1-2-3-4-5	$C_{13}H_{23}O_4Si_2$	TBDMS-OH, TBDMS-CO and TB
$\alpha$ KG (1MEOX, 2TBDMS)	156	4	C2-3-4-5	$C_6H_{10}NO_2Si$	TBDMS-CO, TB and $CH_3O$
$\alpha$ KG (1MEOX, 2TBDMS)	186	4	C1-2-3-4	$C_7H_{12}NO_3Si$	TBDMS-COO and $CH_3$
Malic acid (3TBDMS)	217	2	C2-3	$C_9H_{21}O_2Si_2$	TBDMS-COO, CO, TB and $CH_3$
Malic acid (3TBDMS)	317	3	C2-3-4	$C_{15}H_{33}O_3Si_2$	TBDMS-CO
Malic acid (3TBDMS)	349	1	C2	$C_{15}H_{37}O_3Si_3$	Unknown
Malic acid (3TBDMS)	391	3	C2-3-4	$C_{17}H_{39}O_4Si_3$	TB and CO

and maintained at that temperature for 5 min for the analysis of TMS- and TBDMS-derivatives, respectively. Argon (200 kPa) was used as collision gas for the MS/MS analysis. Collision energy was optimized using Smart MRM (Shimadzu, Kyoto, Japan). The obtained raw MS and MS/MS spectra were deposited in MassBank.<sup>13)</sup>

## RESULTS

### EI-induced fragmentation of TMS- and TBDMS-derivatized organic acids

In this study, the EI- and CID-fragmentation of TMS- and TBDMS-derivatized dicarboxylic and tricarboxylic

acids in the Krebs cycle was investigated by GC-EI-MS and GC-EI-MS/MS. To begin with, the TMS-derivatives of organic acids, one of the major analytes for metabolome analysis, were analyzed using GC-EI-MS. Figure 1 shows the EI-spectra of the TMS-derivatized malic acid. The ions with the highest intensity were 73 and 147, derived from TMS group.<sup>14</sup> The other commonly observed fragment ions in TMS-derivatives were  $[M-15]^+$  and  $[M-117]^+$ , in which the fragments were cleaved at the methyl group in TMS group and TMS-carboxyl group. The EI-spectra of TMS-derivatized malic acid produced intense  $[M-15]^+$  ( $m/z=335$ ) and  $[M-117]^+$  ( $m/z=233$ ) ions with relatively minor ions ( $m/z=101, 117, 189, 217, 265, \text{ and } 307$ ). These data are consistent with the previous study.<sup>14</sup> To identify the carbon atoms included in these fragment ions,  $[U-^{13}C]$ malic acid obtained from yeast cultured in  $[U-^{13}C]$ glucose medium was derivatized and analyzed. As expected, the  $m/z$  of  $[M-15]^+$  and  $[M-117]^+$  ions shifted from 335 to 339, and from 223 to 226, respectively, demonstrating that  $[M-15]^+$  and  $[M-117]^+$  ions contained four and three carbons, respectively (Fig. 1b). Similarly, the fragment ions of  $[f101]^+$ ,  $[f117]^+$ ,  $[f189]^+$ ,  $[f217]^+$ ,  $[f265]^+$ , and  $[f307]^+$  contained 2, 1, 2, 3, 1, and 3 carbons in a malic acid backbone (Fig. 1b). The cleaved TMS-carboxyl group in  $[M-117]^+$  could be C1 or C4 from a malic acid backbone. To determine this, position-specific  $^{13}C$ -labeled malic acid standard was prepared *via in vitro* sequential enzyme reactions of pyruvate kinase, phosphoenolpyruvate (PEP) carboxylase, and malate dehydrogenase supplemented with  $^{13}C$  pyruvic acid and  $^{13}C$  sodium bicarbonate (see experimental section, Fig. S1a). The analyses of TMS-derivatized  $[2-^{13}C]$ ,  $[3-^{13}C]$ , and  $[4-^{13}C]$ malic acids demonstrated the mass shift for  $[M-117]^+$  from 233 to 234 (Fig. 1d–f). However, no shift occurred in the spectrum of  $[1-^{13}C]$ malic acid (Fig. 1c), indicating that the  $[M-117]^+$  loses the C1 in the malic acid backbone. Similarly, the following carbon positions in other fragment ions were identified:  $[f101]^+$  and  $[f189]^+$  had C2-3,  $[f117]^+$  had C4,  $[f265]^+$  had C2, and  $[f217]^+$  and  $[f307]^+$  had C2-3-4 from a malic acid backbone. The estimated structure of each fragment ions is illustrated in Fig. 2a. Although the structure of  $[f265]^+$  cannot be explained in a simple cleavage pattern, it is estimated that three TMS-O groups were possibly linked to the C2 of the malic acid backbone. The same approaches were applied for the survey of fragmentation in the TMS-derivatized citric,  $\alpha$ KG, succinic, and fumaric acids (Supplementary materials 1).  $\alpha$ KG was additionally methoxyaminated before the TMS derivatization. The fragment ions and the estimated structures of these TMS-derivatized organic acids with carbon backbone cleavage are summarized in Table 1 and Fig. 2b–e. The cleavage of TMS-carboxyl group in citric acid was successfully identified as C6 in the carbon backbone (Fig. 2b). The spectrum of  $\alpha$ KG shows the generation of fragment ions with neither C1 nor C5 in the carbon backbone (Fig. 2c). The fragmentation of TMS-derivatized succinic and fumaric acids was estimated as Fig. 2d–e according to the structural symmetry.

The fragmentation of TBDMS-derivatized organic acids was also explored in GC-EI-MS (Supplementary materials 2). The spectra show the relatively abundant  $[M-57]^+$ , which contains all carbon backbones generated by the neutral loss of *tert*-butyl (TB) group. The backbone-derived carbon atoms included in decarboxylated fragment ions were determined by analyzing the synthesized position-specific  $^{13}C$ -labeled

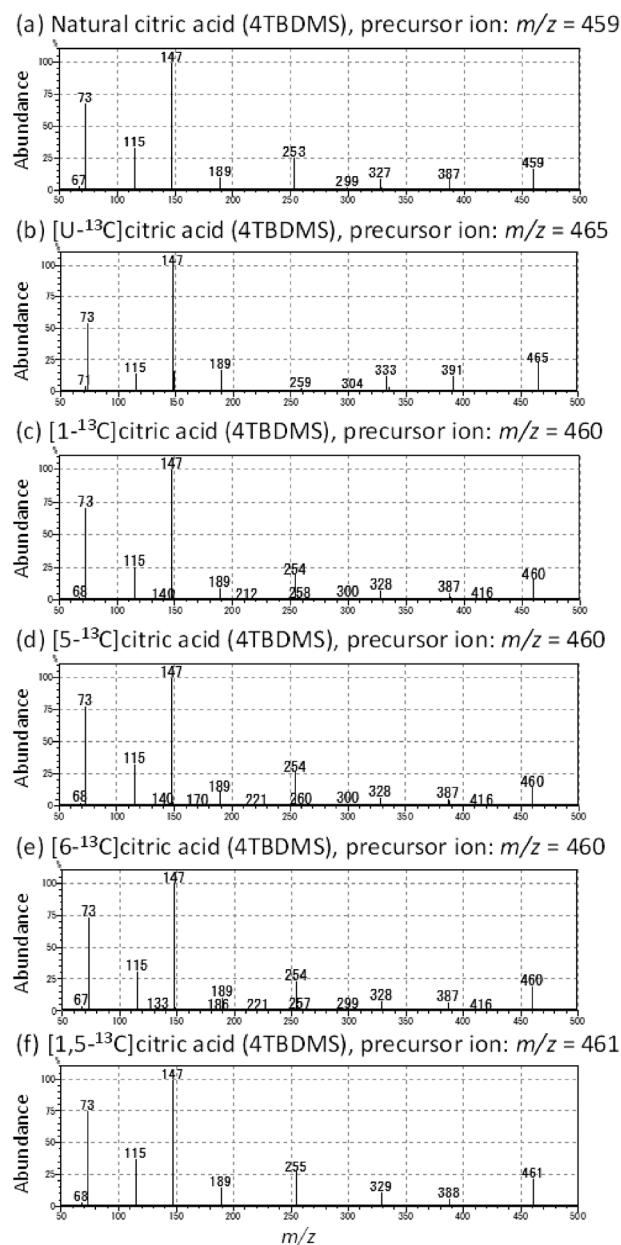


Fig. 3. CID-induced fragmentation of TBDMS-derivatized citric acid.

(a) natural, (b)  $[U-^{13}C]$ , (c)  $[1-^{13}C]$ , (d)  $[5-^{13}C]$ , (e)  $[6-^{13}C]$  and (f)  $[1,5-^{13}C]$ citric acid.

standards in a similar way (Table 1). The decarboxylated position of TBDMS organic acids was similar to that of TMS-derivatized ones. The examples include the cleavage of C6 in citric acid, C1 or C5 of  $\alpha$ KG, and C1 of malic acid.

### CID-induced fragmentation of TMS- and TBDMS-derivatized organic acids

Additionally, CID is a promising fragmentation mode in mass spectrometry. The CID-specific fragmentation was surveyed in TMS and TBDMS-derivatized organic acids using GC-EI-MS/MS (Supplementary materials 3 and 4). Fragment ions containing all carbon backbones, *i.e.*,  $[M-57]^+$  and  $[M-15]^+$  for TBDMS- and TMS-derivatized organic acids, respectively, and relatively intense decarboxylated ions were chosen as precursor ions. The product ion scan of  $[f459]^+$  generated in the EI of TBDMS-derivatized citric acid, which con-

Table 2. Fragment ions with C–C bond cleavage of TMS- and TBDMS-derivatized organic acids by CID.

Organic acids	Precursor ion		Product ion				
	<i>m/z</i>	Carbon skeleton	<i>m/z</i>	Number of organic acid-derived carbons	Carbon skeleton	Estimated chemical formula	Estimated cleavage group from intact molecule
<i>TMS derivatization</i>							
Citric acid (4TMS)	465	C1-2-3-4-5-6	183	5	C1-2-3-4-5	C <sub>8</sub> H <sub>11</sub> O <sub>3</sub> Si	TMS-COO, TMS-OH and TMS-O
Citric acid (4TMS)	465	C1-2-3-4-5-6	257	5	C1-2-3-4-5	C <sub>10</sub> H <sub>17</sub> O <sub>4</sub> Si <sub>2</sub>	CH <sub>3</sub> , TMS-COOH and TMS-OH
Citric acid (4TMS)	465	C1-2-3-4-5-6	347	5	C1-2-3-4-5	C <sub>13</sub> H <sub>27</sub> O <sub>5</sub> Si <sub>3</sub>	CH <sub>3</sub> and TMS-COOH
Citric acid (4TMS)	465	C1-2-3-4-5-6	273	5	C1-2-3-4-5	C <sub>11</sub> H <sub>21</sub> O <sub>4</sub> Si <sub>2</sub>	TMS-OH and TMS-COOH
Citric acid (4TMS)	363	C1-2-3-4-5	183	5	C1-2-3-4-5	C <sub>8</sub> H <sub>11</sub> O <sub>3</sub> Si	TMS-COO, TMS-OH and TMS-O
Citric acid (4TMS)	363	C1-2-3-4-5	273	5	C1-2-3-4-5	C <sub>11</sub> H <sub>21</sub> O <sub>4</sub> Si <sub>2</sub>	TMS-COO and TMS-OH
Citric acid (4TMS)	273	C1-2-3-4-5	183	5	C1-2-3-4-5	C <sub>8</sub> H <sub>11</sub> O <sub>3</sub> Si	TMS-COO, TMS-OH and TMS-O
$\alpha$ KG (1MEOX, 2TMS)	288	C1-2-3-4-5	244	4	C1-2-3-4	C <sub>10</sub> H <sub>22</sub> O <sub>2</sub> NSi <sub>2</sub>	CH <sub>3</sub> O, CO, and CH <sub>3</sub>
$\alpha$ KG (1MEOX, 2TMS)	288	C1-2-3-4-5	170	4	C1-2-3-4	C <sub>7</sub> H <sub>15</sub> NO <sub>2</sub> Si	CH <sub>3</sub> O and TMS-COOH
Fumaric acid (2TMS)	245	C1-2-3-4	217	3	C1-2-3 or C2-3-4	C <sub>8</sub> H <sub>17</sub> O <sub>3</sub> Si <sub>2</sub>	CH <sub>3</sub> and CO
Malic acid (3TMS)	233	C2-3-4	101	2	C2-3	C <sub>4</sub> H <sub>9</sub> OSi	TMS-COO, TMS-OOH and CH <sub>3</sub>
Malic acid (3TMS)	233	C2-3-4	117	1	C4	C <sub>4</sub> H <sub>9</sub> O <sub>2</sub> Si	Except TMS-COO
Malic acid (3TMS)	233	C2-3-4	143	3	C2-3-4	C <sub>6</sub> H <sub>11</sub> O <sub>2</sub> Si	TMS-COO, TMS-OH
Malic acid (3TMS)	233	C2-3-4	189	2	C2-3	C <sub>8</sub> H <sub>21</sub> OSi <sub>2</sub>	TMS-COO, CH <sub>3</sub> and CO
Malic acid (3TMS)	335	C1-2-3-4	307	3	C2-3-4	C <sub>11</sub> H <sub>27</sub> O <sub>4</sub> Si <sub>3</sub>	CH <sub>3</sub> and CO
Malic acid (3TMS)	335	C1-2-3-4	263	2	C2-3	C <sub>10</sub> H <sub>27</sub> O <sub>2</sub> Si <sub>3</sub>	CH <sub>3</sub> , CH <sub>2</sub> , CO and CO
Malic acid (3TMS)	335	C1-2-3-4	217	3	C2-3-4	C <sub>8</sub> H <sub>17</sub> O <sub>3</sub> Si <sub>2</sub>	CH <sub>3</sub> and TMS-COOH
Malic acid (3TMS)	335	C1-2-3-4	117	1	C4	C <sub>4</sub> H <sub>9</sub> O <sub>2</sub> Si	Except TMS-COO
<i>TBDMS derivatization</i>							
Citric acid (4TBDMS)	357	C1-2-3-4-5	225	5	C1-2-3-4-5	C <sub>11</sub> H <sub>17</sub> O <sub>3</sub> Si	TBDMS-OH, TBDMS-COO, TBDMS-O
Citric acid (4TBDMS)	357	C1-2-3-4-5	313	4	C1-2-3-4 or C2-3-4-5	C <sub>16</sub> H <sub>33</sub> O <sub>2</sub> Si <sub>2</sub>	TBDMS-OH, TBDMS-COO, CO and CH <sub>3</sub>
Citric acid (4TBDMS)	431	C1-2-3-4-5	387	4	C1-2-3-4 or C2-3-4-5	C <sub>18</sub> H <sub>39</sub> O <sub>3</sub> Si <sub>3</sub>	TBDMS-OH, TB, CO, CO and CH <sub>3</sub>
Citric acid (4TBDMS)	459	C1-2-3-4-5-6	387	4	C1-2-3-4 or C2-3-4-5	C <sub>18</sub> H <sub>39</sub> O <sub>3</sub> Si <sub>3</sub>	TBDMS-OH, TB, CO, CO and CH <sub>3</sub>
$\alpha$ KG (1MEOX, 2TBDMS)	346	C1-2-3-4-5	186	4	C1-2-3-4	C <sub>7</sub> H <sub>12</sub> O <sub>3</sub> NSi	TB, and TBDMS-COO
$\alpha$ KG (1MEOX, 2TBDMS)	346	C1-2-3-4-5	156	4	C1-2-3-4	C <sub>6</sub> H <sub>10</sub> O <sub>2</sub> NSi	TB, TBDMS-COO and CH <sub>3</sub> O
$\alpha$ KG (1MEOX, 2TBDMS)	156	C1-2-3-4	112	3	C2-3-4	C <sub>5</sub> H <sub>10</sub> NSi	TB, TBDMS-COO CH <sub>3</sub> O, CO and CH <sub>3</sub>
Malic acid (3TBDMS)	419	C1-2-3-4	217	3	C2-3	C <sub>9</sub> H <sub>21</sub> O <sub>2</sub> Si <sub>2</sub>	TB, TBDMS-COO, CO and CH <sub>3</sub>
Malic acid (3TBDMS)	419	C1-2-3-4	391	3	C2-3-4	C <sub>17</sub> H <sub>39</sub> O <sub>4</sub> Si <sub>3</sub>	TB and CO
Malic acid (3TBDMS)	419	C1-2-3-4	403	4	C1-2-3-4	C <sub>17</sub> H <sub>39</sub> O <sub>4</sub> Si <sub>3</sub>	TB and CH <sub>3</sub>
Malic acid (3TBDMS)	317	C2-3-4	273	2	C2-3	C <sub>14</sub> H <sub>33</sub> OSi <sub>2</sub>	TBDMS-COO, CO and CH <sub>3</sub>

tained all carbon backbone, produced the CID-specific ions such as [f253]<sup>+</sup>, [f327]<sup>+</sup>, and [f387]<sup>+</sup> (Fig. 3a). The product ion scan of TBDMS-derivatized [U-<sup>13</sup>C]citric acid showed 6 mass shifts from 253 to 259 and from 327 to 333 and 4 mass shifts from 387 to 391 (Fig. 3b), indicating that [f387]<sup>+</sup> contained 4 citric acid-derived carbon atoms while [f253]<sup>+</sup> and [f327]<sup>+</sup> contained all carbon atoms. The decarboxylated position of [f387]<sup>+</sup> was investigated by the product ion scan of the various position-specific <sup>13</sup>C-labeled citric acids. The *m/z* of product ions of TBDMS-derivatized [6-<sup>13</sup>C]citric acid remained at 387, demonstrating that carboxyl group of C6 was certainly lost in CID. Interestingly, the product ion scan of TBDMS-derivatized [1-<sup>13</sup>C] and [5-<sup>13</sup>C]citric acids produced ions with *m/z* of 387 and 388. These observations suggest that the carboxyl group of either C1 or C5 in TBDMS-citric acid was lost in CID. The assumption was validated by the presence of single ions of *m/z*=388 on the product ion spectra of [1,5-<sup>13</sup>C]-citric acid. These data conclude that [f387]<sup>+</sup> contained either C1-2-3-4 or C2-3-4-5. Similarly, the product ion scan of [f156]<sup>+</sup> containing C1-2-3-4 of methoxyaminated and TBDMS-derivatized  $\alpha$ KG generated in the EI produced [f112]<sup>+</sup> with an additional decarboxylation in C1. Since these fragments were

not observed in EI spectra, CID proved to be a complementary method to measure the <sup>13</sup>C-labeling of decarboxylated fragment ions. The results are summarized in Table 2.

## DISCUSSION

In this study, the EI- and CID-fragmentation of TMS- and TBDMS-derivatized representative dicarboxylic and tricarboxylic acids in the Krebs cycle was explored. The analyses of the position-specific <sup>13</sup>C-labeled standards prepared by *in vitro* enzymatic reactions successfully determined the position of cleaved carbon atom in the organic acid backbone (Tables 1, 2). The findings in our study are summarized as follows: (1) TMS-derivatized organic acids produced more fragment ions with C–C bond cleavage than TBDMS-derivatized ones by EI, (2) the carboxyl group next to hydroxylated carbon was primarily cleaved in EI, and (3) CID generated specific fragment ions with multiple decarboxylations.

As described in Fig. 1, various fragment ions were observed in TMS-derivatized malic acids by EI. In such case, it is possible to calculate the <sup>13</sup>C-labeling of each carbon atom in organic acid backbone based on the <sup>13</sup>C-labeling of sev-

eral fragment ions with C–C bond cleavage.<sup>15)</sup> For example, <sup>13</sup>C-labeling of C1 can be calculated computationally from the <sup>13</sup>C-labeling of fragment ions with C1-2-3-4 and C2-3-4 of TMS-derivatized malic acid. Even the <sup>13</sup>C-labeling of each carbon atom in TMS-derivatized malic acid backbone can be determined by measuring the <sup>13</sup>C-labeling of novel fragment ions identified in this study, leading to maximization of the acquirable information from a single analysis. As compared to TMS-derivatization, TBDMS-derivatized organic acids produced relatively abundant ions with all carbon backbone ( $[M-57]^+$ ) and fewer fragment ions with C–C bond cleavage. For example, fragment ions containing C4 backbone appeared on the spectra of TMS- but not TBDMS-derivatized malic acid. These data demonstrate that TMS-derivatization is beneficial for generating EI-fragment ions with C–C bond cleavage.

Although the fragmentation patterns of TMS and TBDMS-derivatized organic acids were different, a common decarboxylation rule was found in EI. The analyses of the position-specific <sup>13</sup>C-labeled standards validated that C1 of malic acid and C6 of citric acid was cleaved on EI-fragment ions. This suggests that the carboxyl group linked to the hydroxylated carbons in the organic acid backbone is primarily decarboxylated. The decarboxylated fragment ions of C1 and C5 in  $\alpha$ KG were both found in EI, implying that the carboxyl group next to methoxyaminated carbon does not have a prior decarboxylation rule.

This study also highlights the fragmentation specificity of CID in derivatized organic acids as compared to EI. As shown in Table 2, CID can cleave two TBDMS-carboxyl groups. This difference may be explained by the fact that the radical cation could be generated and stabilized even in a single decarboxylation by EI, while the ions of univalent form could be kept stable during two parts of neutral loss of carboxyl group in CID. This insight sheds light on multiple CID events by ion-trap mass spectrometry for in-depth profiling of labeling information.

In this study, the EI- and CID-fragmentation of dicarboxylic and tricarboxylic acids was surveyed. The positions of organic acid-derived carbons contained in each fragment ion were successfully identified. These findings can contribute to the development of a fundamental theory of fragmentation in derivatized organic acids as well as the improvement of <sup>13</sup>C-labeling experiments for biological system.

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## REFERENCES

- 1) J. D. Young, A. A. Shastri, G. Stephanopoulos, J. A. Morgan. Mapping photoautotrophic metabolism with isotopically nonstationary <sup>13</sup>C flux analysis. *Metab. Eng.* 13: 656–665, 2011.
- 2) C. M. Metallo, P. A. Gameiro, E. L. Bell, K. R. Mattaini, J. Yang, K. Hiller, C. M. Jewell, Z. R. Johnson, D. J. Irvine, L.

- Guarente, J. K. Kelleher, M. G. Vander Heiden, O. Iliopoulos, G. Stephanopoulos. Reductive glutamine metabolism by IDH1 mediates lipogenesis under hypoxia. *Nature* 481: 380–384, 2012.
- 3) W. S. Ahn, M. R. Antoniewicz. Metabolic flux analysis of CHO cells at growth and non-growth phases using isotopic tracers and mass spectrometry. *Metab. Eng.* 13: 598–609, 2011.
- 4) J. M. Buescher, M. R. Antoniewicz, L. G. Boros, S. C. Burgess, H. Brunengraber, C. B. Clish, R. J. DeBerardinis, O. Feron, C. Frezza, B. Ghesquiere, E. Gottlieb, K. Hiller, R. G. Jones, J. J. Kamphorst, R. G. Kibbey, A. C. Kimmelman, J. W. Locasale, S. Y. Lunt, O. D. Maddocks, C. Malloy, C. M. Metallo, E. J. Meuillet, J. Munger, K. Noh, J. D. Rabinowitz, M. Ralsler, U. Sauer, G. Stephanopoulos, J. St-Pierre, D. A. Tennant, C. Wittmann, M. G. Vander Heiden, A. Vazquez, K. Vousden, J. D. Young, N. Zamboni, S. M. Fendt. A roadmap for interpreting <sup>13</sup>C metabolite labeling patterns from cells. *Curr. Opin. Biotechnol.* 34: 189–201, 2015.
- 5) T. P. Mawhinney, R. S. R. Robinett, A. Atalay, M. A. Madson. Analysis of amino acids as their *tert*.-butyldimethylsilyl derivatives by gas–liquid chromatography and mass spectrometry. *J. Chromatogr. A* 358: 231–242, 1986.
- 6) M. Rühl, B. Rupp, K. Nöh, W. Wiechert, U. Sauer, N. Zamboni. Collisional fragmentation of central carbon metabolites in LC-MS/MS increases precision of <sup>13</sup>C metabolic flux analysis. *Biotechnol. Bioeng.* 109: 763–771, 2012.
- 7) D. McCloskey, J. D. Young, S. Xu, B. O. Palsson, A. M. Feist. MID Max: LC-MS/MS method for measuring the precursor and product mass isotopomer distributions of metabolic intermediates and cofactors for metabolic flux analysis applications. *Anal. Chem.* 88: 1362–1370, 2016.
- 8) N. Okahashi, S. Kawana, J. Iida, H. Shimizu, F. Matsuda. GC-MS/MS survey of collision-induced dissociation of *tert*-butyldimethylsilyl-derivatized amino acids and its application to <sup>13</sup>C-metabolic flux analysis of *Escherichia coli* central metabolism. *Anal. Bioanal. Chem.* 408: 6133–6140, 2016.
- 9) J. Kappelmann, B. Klein, P. Geilenkirchen, S. Noack. Comprehensive and accurate tracking of carbon origin of LC-tandem mass spectrometry collisional fragments for <sup>13</sup>C-MFA. *Anal. Bioanal. Chem.* 409: 2309–2326, 2017.
- 10) J. Choi, M. R. Antoniewicz. Tandem mass spectrometry for <sup>13</sup>C metabolic flux analysis: Methods and algorithms based on EMU framework. *Front. Microbiol.* 10: 1–8, 2019.
- 11) S. Nishino, N. Okahashi, F. Matsuda, H. Shimizu. Absolute quantitation of glycolytic intermediates reveals thermodynamic shifts in *Saccharomyces cerevisiae* strains lacking PFK1 or ZWF1 genes. *J. Biosci. Bioeng.* 120: 280–286, 2015.
- 12) L. Peng, K. Shimizu. Global metabolic regulation analysis for *Escherichia coli* K12 based on protein expression by 2-dimensional electrophoresis and enzyme activity measurement. *Appl. Microbiol. Biotechnol.* 61: 163–178, 2003.
- 13) H. Horai, M. Arita, S. Kanaya, Y. Nihei, T. Ikeda, K. Suwa, Y. Ojima, K. Tanaka, S. Tanaka, K. Aoshima, Y. Oda, Y. Kakazu, M. Kusano, T. Tohge, F. Matsuda, Y. Sawada, M. Y. Hirai, H. Nakanishi, K. Ikeda, N. Akimoto, T. Maoka, H. Takahashi, T. Ara, N. Sakurai, H. Suzuki, D. Shibata, S. Neumann, T. Iida, K. Tanaka, K. Funatsu, F. Matsuura, T. Soga, R. Taguchi, K. Saito, T. Nishioka. MassBank: A public repository for sharing mass spectral data for life sciences. *J. Mass Spectrom.* 45: 703–714, 2010.
- 14) G. Petersson. Mass spectrometry of hydroxy dicarboxylic acids as trimethylsilyl derivatives. Rearrangement fragmentations. *Org. Mass Spectrom.* 6: 565–576, 1972.
- 15) J. Choi, M. T. Grossbach, M. R. Antoniewicz. Measuring complete isotopomer distribution of aspartate using gas chromatography/tandem mass spectrometry. *Anal. Chem.* 84: 4628–4632, 2012.