

# Abundant Extraterrestrial Amino Acids in the Primitive CM Carbonaceous Chondrite Asuka 12236

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**Abstract.** The Asuka (A) 12236 meteorite has recently been classified as a CM carbonaceous chondrite of petrologic type 2.9 and is among the most primitive CM meteorites studied to date (Kimura *et al.* 2020). We report the concentrations, relative distributions, and enantiomeric ratios of amino acids in a hot water extract of the A 12236 meteorite determined by ultra-high-performance liquid chromatography time-of-flight mass spectrometry (Glavin *et al.* 2020). A wide diversity of two- to six-carbon aliphatic primary amino acids were identified in A 12236, which had a total amino acid abundance of  $360 \pm 18$  nmol g<sup>-1</sup>, with most amino acids present in a free form (Table 1). The amino acid concentrations of A 12236 were double those previously measured in the CM2.7 Paris meteorite (Table 1), consistent with A 12236 being a highly primitive and unheated CM chondrite. The high relative abundance of  $\alpha$ -amino acids in A 12236 are also consistent with formation by a Strecker-cyanohydrin dominated synthesis during a limited early aqueous alteration phase on the CM meteorite parent body. The presence of predominantly free glycine, a near racemic mixture of alanine (D/L ~0.93 to 0.96), and elevated abundances of several terrestrially rare non-protein amino acids including  $\alpha$ -aminoisobutyric acid ( $\alpha$ -AIB) and racemic isovaline, indicate that these amino acids in A 12236 are extraterrestrial in origin. Given a lack of evidence for biological amino acid contamination in A 12236, it is possible that some of the L-enantiomeric excesses ( $L_{ee}$  ~34 to 64%) of the protein amino acids aspartic and glutamic acids and serine are indigenous to the meteorite; however, isotopic measurements are needed for confirmation. In contrast to more aqueously altered CMs of petrologic types  $\leq 2.5$ , no L-isovaline excesses were detected in A 12236. This observation strengthens the hypothesis that extensive parent body aqueous activity is required to produce or amplify the large L-isovaline excesses that cannot be explained solely by exposure to circularly polarized radiation or other chiral symmetry breaking mechanisms prior to incorporation into the asteroid parent body. The fact that only L-enantiomeric excesses (and no D-excesses) have been observed in amino acids with a single asymmetric carbon in carbonaceous meteorites suggests that the origin of life on Earth or elsewhere in our solar system may have been biased toward L-amino acid homochirality from the very beginning. The return of pristine materials from the surface of carbonaceous asteroids to Earth for detailed laboratory organic analyses will be essential to advance our understanding of the origin and evolution of amino acid chiral asymmetry in the early Solar System given that all meteorites on Earth have been compromised, to some degree, by terrestrial contamination. JAXA's Hayabusa2 mission and NASA's OSIRIS-REx mission will return the first pristine samples collected from the surfaces of the carbonaceous asteroids Ryugu and Bennu. These missions will provide a unique opportunity to evaluate the effects of parent body processing on the distributions and enantiomeric abundances of amino acids and other prebiotic molecules in carbon-rich asteroids.

**Table 1.** Summary of the average concentrations (nmol g<sup>-1</sup>) of the two- to six-carbon amino acids in the non-hydrolyzed (free) and 6M HCl acid-hydrolyzed (total) water extracts of the CM meteorites Asuka 12236 and Paris.

Amino Acids	Asuka 12236 (CM2.9)		Paris (CM2.7)	
	free	total	free	total
<b>Dicarboxylic Amino Acids</b>				
D-aspartic acid	1.7 ± 0.2	2.4 ± 0.1	0.67 ± 0.03	1.01 ± 0.04
L-aspartic acid	1.8 ± 0.1	4.9 ± 0.2	0.90 ± 0.05	1.50 ± 0.04
D-glutamic acid	0.63 ± 0.07	3.1 ± 0.2	4.1 ± 0.3	5.6 ± 0.4
L-glutamic acid	0.9 ± 0.1	8.4 ± 0.5	4.9 ± 0.3	6.7 ± 0.5
<b>Hydroxy Amino Acids</b>				
D-serine	1.23 ± 0.05	1.1 ± 0.2	n.r.	n.r.
L-serine	2.78 ± 0.07	5.0 ± 0.2	n.r.	n.r.
D-threonine	n.d.	n.d.	n.r.	n.r.
L-threonine	n.d.	n.d.	n.r.	n.r.
<b>C2 Amino Acid</b>				
Glycine	155 ± 3	160 ± 5	45 ± 2	110 ± 5
<b>C3 Amino Acids</b>				
$\beta$ -alanine	13.1 ± 0.6	21 ± 1	7.0 ± 0.3	14 ± 2
D-alanine	25 ± 1	28 ± 2	3.3 ± 0.1	8.0 ± 0.4
L-alanine	26 ± 1	30 ± 2	4.9 ± 0.5	10.8 ± 0.5

<i>C4 Amino Acids</i>				
D,L- $\alpha$ -amino- <i>n</i> -butyric acid	15 $\pm$ 1 <sup>a</sup>	16 $\pm$ 2 <sup>a</sup>	1.17 $\pm$ 0.04 <sup>b</sup>	2.97 $\pm$ 0.07 <sup>b</sup>
D- $\beta$ -amino- <i>n</i> -butyric acid	3.35 $\pm$ 0.07	4.5 $\pm$ 0.1	< 0.01	< 0.01
L- $\beta$ -amino- <i>n</i> -butyric acid	3.2 $\pm$ 0.1	4.4 $\pm$ 0.1	< 0.01	< 0.01
$\alpha$ -aminoisobutyric acid	17.0 $\pm$ 0.2	17.9 $\pm$ 0.4	0.84 $\pm$ 0.06	3.0 $\pm$ 0.1
D,L- $\beta$ -aminoisobutyric acid <sup>a</sup>	n.d.	n.d.	< 0.01	< 0.01
$\gamma$ -amino- <i>n</i> -butyric acid	2.0 $\pm$ 0.1	12.1 $\pm$ 0.6	1.2 $\pm$ 0.1	1.8 $\pm$ 0.2
<i>C5 Amino Acids</i>				
D-norvaline (D-2-apa)	2.42 $\pm$ 0.05	2.7 $\pm$ 0.2	< 0.3	1.32 $\pm$ 0.03
L-norvaline (L-2-apa)	2.48 $\pm$ 0.05	2.5 $\pm$ 0.1	< 0.4	1.52 $\pm$ 0.05
D-isovaline (D-2-a-2-mba)	3.9 $\pm$ 0.3	4.2 $\pm$ 0.3	0.22 $\pm$ 0.02 <sup>a</sup>	1.9 $\pm$ 0.1
L-isovaline (L-2-a-2-mba)	4.1 $\pm$ 0.2	4.0 $\pm$ 0.2		1.9 $\pm$ 0.1
D-valine (D-2-a-3-mba)	5.1 $\pm$ 0.1	5.6 $\pm$ 0.9	1.86 $\pm$ 0.07	4.9 $\pm$ 0.1
L-valine (L-2-a-3-mba)	5.7 $\pm$ 0.2	6.8 $\pm$ 0.4	1.9 $\pm$ 0.2	5.2 $\pm$ 0.2
D,L-3-apa <sup>b</sup>	1.8 $\pm$ 0.1	1.8 $\pm$ 0.2	n.r.	n.r.
D,L- and allo-3-a-2-mba <sup>b</sup>	1.40 $\pm$ 0.04	1.4 $\pm$ 0.2	n.r.	n.r.
3-a-3-mba	0.92 $\pm$ 0.04	0.8 $\pm$ 0.1	n.r.	n.r.
3-a-2,2-dmpa	0.98 $\pm$ 0.05	2.48 $\pm$ 0.03	n.r.	n.r.
D,L-3-a-2-epa <sup>a</sup>	< 0.3 <sup>c</sup>	< 0.3 <sup>c</sup>	n.r.	n.r.
D,L-4-apa <sup>a</sup>	0.46 $\pm$ 0.07	1.46 $\pm$ 0.05	n.r.	n.r.
D,L-4-a-2-mba <sup>a</sup>	n.d.	2.6 $\pm$ 0.2	n.r.	n.r.
D,L-4-a-3-mba <sup>a</sup>	0.05 $\pm$ 0.01	0.28 $\pm$ 0.02	n.r.	n.r.
5-apa	0.64 $\pm$ 0.04	2.7 $\pm$ 0.3	n.r.	n.r.
<i>C6 Amino Acids</i>				
$\epsilon$ -amino- <i>n</i> -caproic acid (EACA)	1.16 $\pm$ 0.05	2.00 $\pm$ 0.04	< 0.01	< 0.01
<b>Sum (nmol g<sup>-1</sup>)</b>	<b>300 <math>\pm</math> 9</b>	<b>360 <math>\pm</math> 18</b>	<b>78 <math>\pm</math> 4</b>	<b>182 <math>\pm</math> 10</b>

<sup>a</sup>Enantiomers could not be separated under the chromatographic conditions.

<sup>b</sup>Enantiomers could be separated, but not identified individually due to the lack of optically pure standards.

<sup>c</sup>Poor chromatographic resolution prevented accurate quantification, therefore only upper limits are reported.

n.d. = amino acids identified, but concentration not determined due to interfering peaks or other analytical issues.

n.r. = concentration not reported.

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