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**The Effects of Parent Body Processes on Amino Acids in Carbonaceous  
Chondrites**

Daniel P. Glavin\*, Michael P. Callahan, Jason P. Dworkin, and Jamie E. Elsila

NASA Goddard Space Flight Center, Greenbelt, MD 20771

\*Corresponding author. Email: [daniel.p.glavin@nasa.gov](mailto:daniel.p.glavin@nasa.gov), Tel: 301-614-6361, FAX: 301-614-6406

## ABSTRACT

To investigate the effect of parent body processes on the abundance, distribution, and enantiomeric composition of amino acids in carbonaceous chondrites, the water extracts from nine different powdered CI, CM, and CR carbonaceous chondrites were analyzed for amino acids by ultrahigh performance liquid chromatography-fluorescence detection and time-of-flight mass spectrometry (UPLC-FD/ToF-MS). Four aqueously altered type 1 carbonaceous chondrites including Orgueil (CI1), Meteorite Hills (MET) 01070 (CM1), Scott Glacier (SCO) 06043 (CM1), and Grosvenor Mountains (GRO) 95577 (CR1) were analyzed using this technique for the first time. Analyses of these meteorites revealed low levels of two- to five-carbon acyclic amino alkanolic acids with concentrations ranging from ~1 to 2,700 parts-per-billion (ppb). The type 1 carbonaceous chondrites have a distinct distribution of the five-carbon ( $C_5$ ) amino acids with much higher relative abundances of the  $\gamma$ - and  $\delta$ -amino acids compared to the type 2 and type 3 carbonaceous chondrites, which are dominated by  $\alpha$ -amino acids. Much higher amino acid abundances were found in the CM2 chondrites Murchison, Lonewolf Nunataks (LON) 94102, and Lewis Cliffs (LEW) 90500, the CR2 Elephant Moraine (EET) 92042, and the CR3 Queen Alexandra Range (QUE) 99177. For example,  $\alpha$ -aminoisobutyric acid ( $\alpha$ -AIB) and isovaline were ~100 to 1000 times more abundant in the type 2 and 3 chondrites compared to the more aqueously altered type 1 chondrites. Most of the chiral amino acids identified in these meteorites were racemic, indicating an extraterrestrial abiotic origin. However, non-racemic isovaline was observed in the aqueously altered carbonaceous chondrites Murchison, Orgueil, SCO 06043, and GRO 95577 with L-isovaline excesses ranging from ~11 to 19%, whereas the most pristine, unaltered carbonaceous chondrites analyzed in this study had no detectable L-isovaline excesses. These results are consistent with the theory that aqueous alteration played an important role in amplification of small initial left handed isovaline excesses on the parent bodies.

**Keywords:** Amino acids, Antarctic carbonaceous chondrites, parent body aqueous alteration, EET 92042, enantiomeric excess, GRO 95577, isovaline, LEW 90500, liquid chromatography time-of-flight mass spectrometry, LEW 90500, LON 94102, MET 01070, Murchison, Orgueil, SCO 06043, Strecker synthesis, QUE 99177

## INTRODUCTION

Meteorites provide a record of the chemical processes that occurred in the early solar system before life began on Earth. The delivery of organic compounds by carbonaceous chondrites to the early Earth and other planetary bodies could have been an important source of prebiotic compounds required for the emergence of life as we know it (Chyba and Sagan, 1992). The carbonaceous chondrite class represents roughly 4% of all meteorite falls (Grady, 2000) and is sub-divided into separate groups (CI, CM, CR, CV, CO, CH, CB, and CK) based on differences in elemental composition and mineralogy (Sears and Dodd, 1988). In addition, a number is given after the carbonaceous chondrite group based on their petrologic histories (i.e., degree of mineralogical or textural modification of the meteorite by aqueous and/or thermal alteration on the parent body), where type 1 chondrites represent the most altered by low temperature aqueous alteration and type 3 the least aqueously altered (McSween, 1979; Van Schmus and Hayes, 1974). Petrographic types 4 to 6 refer to carbonaceous chondrites with increasing thermal alteration. Carbonaceous chondrites represent a very primitive class of meteorite that generally contain 2 to 5 wt.% of carbon, most of which is from organic matter (Sephton, 2002). More than 70% of the organic carbon is in the form of an insoluble kerogen-like macromolecule that is structurally complex (Cody and Alexander, 2005; Sephton et al., 2000), while the remaining solvent extractable organic carbon is comprised of a complex mixture of organics that includes amino acids.

The amino acid compositions of a variety of carbonaceous chondrites, in particular the CMs, have been studied extensively because these prebiotic molecules are essential components of life as the monomers of proteins and enzymes. Over 80 different amino acids have been named in the CM chondrites Murchison and Murray, comprising a mixture of 2- to 8-carbon cyclic and acyclic monoamino alkanolic and alkanedioic acids of nearly complete structural diversity, many of which are rare or absent in the terrestrial biosphere (Botta and Bada, 2002; Cronin and Chang, 1993; Cronin and Pizzarello, 1983; Cronin and

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3 Pizzarello, 1986; Sephton, 2002). Recent studies of organic compounds in solvent extracts  
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5 of the Murchison meteorite by ultrahigh resolution Fourier transform ion cyclotron resonance  
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7 mass spectrometry have detected thousands of different molecular compositions and likely  
8  
9 millions of diverse structures (Schmitt-Kopplin et al., 2010). Therefore, it is likely that  
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11 previous amino acid analyses of Murchison have greatly underestimated the chemical  
12  
13 complexity of this meteorite due to a lack of sensitivity and standards. Amino acids have  
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15 also been identified in the more aqueously altered C11 carbonaceous chondrites Orgueil and  
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17 Ivuna, and these two meteorites have a relatively simple distribution of amino acids  
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19 (predominantly  $\beta$ -alanine and glycine) that is distinct from the complex mixture of amino  
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21 acids found in the CMs Murchison and Murray (Ehrenfreund et al., 2001). CR chondrites  
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23 are believed to contain the most primitive insoluble organic matter (IOM) of any  
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25 carbonaceous chondrite group (Cody and Alexander, 2005), and it has recently been  
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27 discovered that two Antarctic CR2 chondrites, Elephant Moraine (EET) 92042 and Graves  
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29 Nunataks (GRA) 95229 have the highest amino acid abundances of any meteorite analyzed  
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31 to date (Martins et al., 2007; Pizzarello et al., 2008). One of the most primitive CR  
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33 chondrites, the CR3 Queen Alexandra Range (QUE) 99177 has also been observed to have  
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35 a similarly high abundance and distribution of five-carbon amino acids compared to EET  
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37 92042 (Glavin and Dworkin, 2009).  
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42 Most amino acids found in carbonaceous chondrites are structurally chiral, meaning they  
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44 possess two non-superimposable mirror image structures or enantiomers (by convention: S,  
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46 left, L and R, right, D). With a few very rare exceptions, only L-amino acids are found in  
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48 biology, whereas amino acids formed by abiotic processes are racemic (equal mixtures of L-  
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50 and D-enantiomers). Therefore, the molecular structure of these compounds can be a very  
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52 useful tool to help discriminate between biotic and abiotic (non-biological) origins of amino  
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54 acids in meteorites. The first analyses of interior pieces of the Murchison meteorite shortly  
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56 after its fall found that the chiral amino acids were racemic ( $D/L = 1$ ), indicating that the  
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58 amino acids were indigenous to the meteorite and very little, if any, terrestrial amino acid  
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3 contamination of the meteorite had occurred (Kvenvolden et al., 1970; Kvenvolden et al.,  
4 1971). One major discovery is the finding of non-racemic  $\alpha$ -dialkyl amino acids in the CM  
5 chondrites Murchison and Murray, with slight to significant L-isovaline excesses ranging  
6 from 0 to 15.2% (Cronin and Pizzarello, 1997; Pizzarello and Cronin, 2000; Pizzarello et al.,  
7 2003). These results are difficult to explain since the abiotic formation of isovaline and other  
8  $\alpha$ -dialkyl amino acids (e.g. by Strecker synthesis) on the CM parent body should produce  
9 racemic mixtures. In contrast to the  $\alpha$ -hydrogen protein amino acids common to all life on  
10 Earth, non-biological  $\alpha$ -dialkyl amino acids such as isovaline found in meteorites are not  
11 prone to geologically rapid racemization (conversion of one enantiomer to the other) under  
12 aqueous or radiogenic conditions (Pollock et al., 1975). Therefore, the initial enantiomeric  
13 ratios of these amino acids are more likely to have been preserved since the time of their  
14 formation. The finding of L-amino acid excesses in Murchison and Murray could point  
15 toward a possible prebiotic contribution to the origin of biological homochirality by the  
16 delivery of extraterrestrial organic material from asteroids and comets to the early Earth.

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34 Amino acid analyses by high performance liquid chromatography with fluorescence  
35 detection (HPLC-FD) of the Antarctic type 1 CM carbonaceous chondrites MET 01070, Allan  
36 Hills (ALH) 88045, and LaPaz Icefield (LAP) 02277, and the only known type 1 CR chondrite  
37 GRO 95577 have been reported (Botta et al., 2007; Martins et al., 2007). The very low  
38 amino acid abundances in these meteorites compared to CM2 and CR2 chondrites was  
39 suggested to be a result of chemical oxidation during an extended aqueous alteration phase  
40 on the parent bodies (Martins et al., 2007). It has been suggested that aqueous alteration  
41 on the parent body may also play an important role in the enrichment of L-isovaline  
42 observed in the CI1 Orgueil and CM2 Murchison meteorites (Glavin and Dworkin, 2009) by  
43 asymmetric amplification via crystallization (Blackmond, 2004; Klussmann et al., 2006) or  
44 asymmetric autocatalytic reactions (Soai et al., 1995). Additional measurements of the  
45 enantiomeric ratio of isovaline in other aqueously altered type 1 chondrites are necessary to  
46 test this hypothesis. Precise abundances and enantiomeric ratios for isovaline and other  
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3 five-carbon ( $C_5$ ) amino acids were not determined in the previous analyses of CM1 and CR1  
4 chondrites (Botta et al., 2007; Martins et al., 2007). In addition, the compound assignments  
5 of the amino acids extracted from these chondrites were based solely on derivatization  
6 chemistry and chromatographic retention time without confirmation by high-resolution mass  
7 spectrometry. To our knowledge, amino acid analyses of any kind have not been previously  
8 reported for the CM1 chondrite Scott Glacier (SCO) 06043.  
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10 To continue to investigate the potential role of aqueous alteration in amino acid  
11 decomposition, amplification of L-isovaline enantiomeric excess, and changes in the relative  
12 distributions of amino acids in carbonaceous chondrites, we analyzed the type 1  
13 carbonaceous chondrites Orgueil (CI1), SCO 06043 (CM1), MET 01070 (CM1) and GRO  
14 95577 (CR1) and compared our measurements with data from several less altered type 2  
15 and type 3 CM and CR carbonaceous chondrites including Murchison (CM2), LEW 90500  
16 (CM2), LON 94102 (CM2), EET 92042 (CR2), and QUE 99177 (CR3), using the identical  
17 extraction and analytical technique. The CR3 chondrite QUE 99177 was originally  
18 classified as a CR2, but analyses of presolar grains and the carbon isotopic composition of  
19 organic matter in this meteorite indicate that QUE 99177 is less altered than other CR2  
20 chondrites and is the most primitive CR chondrite analyzed to date (Floss and Stadermann,  
21 2009). Although LON 94102 has been classified as a CM2, it is a relatively unaltered  
22 meteorite, with oxygen isotope values that fall at the low end (i.e. least hydrated) of the CM  
23 chondrites (Clayton and Mayeda, 1999). Amino acid abundances and enantiomeric ratios  
24 were determined using a highly sensitive and selective ultrahigh performance liquid  
25 chromatography fluorescence detection and time of flight mass spectrometry (UPLC-  
26 FD/ToF-MS) technique (Glavin and Dworkin, 2009; Johnson et al., 2008) coupled with *o*-  
27 phthaldialdehyde/*N*-acetyl-L-cysteine (OPA/NAC) derivatization. All of the UPLC-FD/ToF-  
28 MS results presented here for the CR1 and CM1 meteorites and one sample of the CM2  
29 Murchison meteorite (USNM 5453) have not previously been reported. For comparison,  
30 some of the amino acid data for the CI Orgueil, the CM's Murchison (USNM 6650), LEW  
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3 90500, and LON 94102 and the CR's EET 92042 and QUE 99177 shown in Tables 2-5 were  
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5 taken from previous studies (Glavin and Dworkin, 2009; Glavin et al., 2006).  
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## 8 **MATERIALS AND METHODS**

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### 10 **Meteorite Samples and Controls**

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12 The amino acid extracts of the CI, CM, and CR carbonaceous chondrites analyzed in  
13 this study were obtained from interior chips that did not contain any visual evidence of fusion  
14 crust. All of the meteorite samples were received as chips and were separately crushed into  
15 fine powders (estimated < 150  $\mu\text{m}$ ) and manually homogenized by using a mortar and pestle  
16 in a positive pressure HEPA filtered laminar flow hood (AirClean). We analyzed only one CI  
17 chondrite Orgueil for this study (total mass 1.0 g) which was provided by the Musée  
18 National, Paris (petrographic type 1); less altered CI chondrites (type 2 or type 3) have not  
19 yet been identified. For the CM's, both type 1 and 2 chondrites were analyzed. The CR  
20 chondrites analyzed in this study represent the entire range of aqueous alteration (type 1-3).  
21 Two different pieces of the CM2 Murchison meteorite were provided by the Smithsonian  
22 National Museum of Natural History, Washington, D.C (USNM 6650, total mass 6.3 g;  
23 USNM 5453, total mass 15.2 g). The Antarctic meteorites studied were collected by  
24 Antarctic Search for Meteorites (ANSMET) field teams and are designated by the  
25 geographic area where they were found and a five digit number, the first two digits of which  
26 indicate the collection season year. In addition, the specific daughter split, parent rock  
27 number, and mass of each chip allocated for this study are also shown in parentheses. The  
28 Antarctic CM2 chondrites LEW 90500 (specific 69, parent 1, total mass 5.0 g) and LON  
29 94102 (specific 19, parent 7, total mass 0.6 g), the CR1 chondrite GRO 95577 (specific 46,  
30 parent 10, total mass 0.5 g), CR2 chondrite EET 92042 (specific 58, parent 0, total mass 0.6  
31 g), CR3 chondrite QUE 99177 (specific 29, parent 10, total mass 0.5 g), and the CM1  
32 chondrites MET 01070 (specific 34, parent 0, total mass 0.5 g) and SCO 06043 (specific 8,  
33 parent 0, total mass 0.5 g) were provided by the Antarctic meteorite curator at the NASA  
34 Johnson Space Center in Houston, TX. As controls, a crushed serpentine (a hydrated  
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3 magnesium silicate) sample that had been heated at 500°C in air overnight and amino acid  
4 standards were carried through the identical extraction procedure as the meteorite samples.  
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### 8 9 10 **Chemicals and Reagents**

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12 Most of the chemicals and reagents used in this study were purchased from Sigma-  
13 Aldrich. A stock amino acid solution ( $\sim 10^{-5}$  to  $10^{-6}$  M) was prepared by mixing individual  
14 standards (97-99% purity) in Millipore water (see Fig. 1). The OPA/NAC reagent used for  
15 amino acid derivatization was prepared by dissolving 4 mg OPA in 300  $\mu$ L methanol (Optima  
16 grade), and then adding 685  $\mu$ L 0.1 M sodium borate buffer (pH 9) and 15  $\mu$ L 1 M NAC. A  
17 0.1 M hydrazine ( $\text{NH}_2\text{NH}_2$ ) solution was prepared by vacuum distillation of concentrated  
18 anhydrous hydrazine (98% purity) and subsequent dilution in Millipore water. The HCl was  
19 double-distilled, and the ammonium formate buffer used in the UPLC-FD/ToF-MS analyses  
20 was prepared by ammonia titration (95% purity) of a 50 mM formic acid solution to pH 8. A  
21 10  $\mu$ M phenolphthalein solution in acetonitrile with 0.1% formic acid was used for internal  
22 mass calibration of the ToF-MS.  
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37 A separate standard solution containing all of the acyclic  $\text{C}_5$   $\alpha$ -,  $\beta$ -,  $\gamma$ -, and  $\delta$ -amino  
38 alkanolic acids (see Fig. 2) was prepared by weighing each individual standard using a  
39 microbalance and mixing them in Millipore water at a concentration of  $\sim 10^{-5}$  to  $10^{-6}$  M.  
40 Individual D- and L-2-amino-2-methylbutanoic acid (2-a-2-mba, isovaline) were purchased  
41 from Acros Organics (>99% purity), and a racemic mixture (D = L) was prepared by mixing  
42 the appropriate volumes of each compound. DL-2-aminopentanoic acid (2-apa, norvaline),  
43 DL-2-amino-3-methylbutanoic acid (2-a-3-mba, valine), and 5-aminopentanoic acid (5-apa)  
44 standards were purchased from Sigma-Aldrich (97-99% purity). DL-3-aminopentanoic acid  
45 (3-apa) was from AzaN (97% purity). 3-Amino-3-methylbutanoic acid (3-a-3-mba), 3-amino-  
46 2,2-dimethylpropanoic acid (3-a-2,2-dmpa), DL-3-amino-2-methylbutanoic acid (3-a-2-mba  
47 and allo-3-a-2-mba, 4 stereoisomers), DL-4-aminopentanoic acid (4-apa), DL-4-amino-3-  
48 methylbutanoic acid (4-a-3-mba), L-3-Amino-2-ethylpropanoic acid (3-a-2-epa), DL-4-amino-  
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3 2-methylbutanoic acid (4-a-2-mba) were individually synthesized and provided by S.  
4 Pizzarello (ASU), S. Miller (UCSD), S. Davies (Oxford), and R. Duke (U. Sydney). The L-3-  
5 a-2-epa was mostly racemized by 6 M HCl acid vapor hydrolysis (150°C for 1 week).  
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### 8 9 **Extraction Procedure**

10 All glassware, ceramics, and sample handling tools were rinsed with Millipore Direct Q3  
11 UV (18.2 MΩ, < 5 ppb total organic carbon) ultrapure water, wrapped in aluminum foil and  
12 then heated in a furnace at 500°C overnight. A portion of each powdered meteorite sample  
13 (~ 0.1 to 0.2 g) was flame sealed separately in a glass ampoule with 1 mL of Millipore water  
14 and extracted 100°C for 24 h. Half of the water supernatants were then subjected to a 6 M  
15 HCl acid vapor hydrolysis procedure at 150°C for 3 hours to determine total hydrolyzable  
16 amino acid content (Glavin et al., 2006). The acid-hydrolyzed, hot-water extracts were  
17 desalted by using cation-exchange resin (AG50W-X8, 100-200 mesh, hydrogen form, BIO-  
18 RAD), and the amino acids recovered by elution with 2 M NH<sub>4</sub>OH (prepared from Millipore  
19 water and NH<sub>3</sub>(g) (AirProducts, *in vacuo*). During ion exchange column loading, a DL-  
20 norleucine internal standard was added to each sample to estimate the amino acid  
21 recoveries from desalting and OPA/NAC derivatization. The non-hydrolyzed extracts of the  
22 meteorite samples were taken through the identical desalting procedure in parallel with the  
23 acid-hydrolyzed extracts. After desalting, the amino acids in the NH<sub>4</sub>OH eluates were dried  
24 under vacuum to remove excess ammonia; the residues were then re-dissolved in 100 μL of  
25 Millipore water, transferred to sterile microcentrifuge tubes, and stored at -20°C prior to  
26 analysis. Based on our analysis of amine standards taken through the entire extraction  
27 procedure, it is likely that a significant fraction of simple volatile amines including indigenous  
28 ammonia, methylamine, ethylamine, and propylamines were lost from the meteorite extracts  
29 during evaporation of the NH<sub>4</sub>OH eluates; therefore, these compounds were not quantified.  
30 However, it is important to note that no decomposition or thermal degradation of amino  
31 acids during the extraction procedure was observed after comparison of the peak areas in  
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3 amino acid standards taken through the identical hot water extraction and acid hydrolysis  
4 treatment as the meteorite samples.  
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### 7 8 **UPLC-FD/ToF-MS Amino Acid Analyses** 9

10 Prior to analysis, 10  $\mu\text{L}$  of each meteorite extract, procedural blank, or standard was  
11 added to 10  $\mu\text{L}$  of 0.1 M sodium borate buffer at room temperature and then derivatized by  
12 adding 5  $\mu\text{L}$  of *o*-phthaldialdehyde/*N*-acetyl-L-cysteine (OPA/NAC). The OPA/NAC reaction  
13 was quenched after 1 or 15 minutes with 75  $\mu\text{L}$  of 0.1 M aqueous hydrazine, and 25  $\mu\text{L}$  of  
14 the solution was immediately injected into a Waters ACQUITY UPLC with a fluorescence  
15 detector and Waters LCT Premier time-of-flight mass spectrometer using positive  
16 electrospray ionization. The details of the ToF-MS settings and the amino acid  
17 quantification methods used, and the range of linear response for these analyses are  
18 described elsewhere (Glavin et al., 2006). The remaining derivatized sample (and a parallel  
19 derivatized standard) was stored in a  $-86^\circ\text{C}$  freezer for repeat analyses. We have found  
20 through monitoring standards that OPA/NAC amino acid derivatives are stable for up to 1  
21 week at  $-86^\circ\text{C}$  without significant degradation. Each derivatized sample was analyzed using  
22 our standard tandem LC column conditions for initial two- to ten-carbon ( $\text{C}_2$  -  $\text{C}_{10}$ ) amino acid  
23 separation and characterization (Glavin et al., 2010). The first column was a Waters BEH  
24 C18 column (2.1 x 50 mm, 1.7- $\mu\text{m}$  bead) followed by a second Waters BEH Phenyl-Hexyl  
25 column (2.1 x 150 mm, 1.7- $\mu\text{m}$  bead). The conditions for separation of the OPA/NAC amino  
26 acid derivatives at  $30^\circ\text{C}$  were as follows: flow rate, 150  $\mu\text{L}/\text{min}$ ; solvent A (50 mM  
27 ammonium formate, 8% methanol, pH 8.0); solvent B (methanol); gradient, time in minutes  
28 (%B): 0 (0), 35 (55), 45 (100). The following day a different gradient using the same  
29 columns and buffers optimized specifically for the separation of the five-carbon ( $\text{C}_5$ ) amino  
30 acids was used to analyze the same derivatized extract on the same columns (Glavin and  
31 Dworkin, 2009).  
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58 Amino acid abundances and their enantiomeric ratios in the meteorite extracts were  
59 determined by comparison of the peak areas generated from the UV fluorescence detector  
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3 and ToF-MS of the OPA/NAC amino acid derivatives to the corresponding peak areas of  
4 standards under the same chromatographic conditions on the same day (see Figs. 1-3).  
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6 Amino acid peak identifications are given in Table 1. In addition to identifying the major  
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8 fluorescent peaks present in the UPLC-FD/ToF-MS chromatograms by retention time, we  
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10 also searched for the masses of the OPA/NAC derivatives corresponding to C<sub>2</sub>-C<sub>10</sub> amino  
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12 acids by plotting the mass of each amino acid derivative over the elution time (see Fig. 2).  
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14 The total amino acid concentrations in parts-per-billion (ppb) by bulk sample weight, the  
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16 enantiomeric ratios, and L-isovaline excesses measured for the meteorite extracts reported  
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18 in Tables 2-5 represent the average of between four and twenty three separate analyses.  
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20 The highly sensitive and selective UPLC-FD/ToF-MS technique employed in this study has a  
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22 detection limit for amino acids that is a factor of ~10<sup>3</sup> times lower than for current state-of-  
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24 the-art gas chromatography-mass spectrometer (GC-MS) measurements (Glavin et al.,  
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26 2006). However, due to the extremely low amino acid abundances and limited sample mass  
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28 available for the CM1 and CR1 carbonaceous chondrites analyzed in this study, GC-MS  
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30 analyses and compound specific isotopic measurements for these meteorites were not  
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32 possible.  
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## 38 RESULTS AND DISCUSSION

### 39 Amino Acid Analyses of the Carbonaceous Chondrites

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41 **UPLC-FD measurements and total amino acid abundances.** Typical UPLC-FD  
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43 chromatograms of the 6 M HCl-hydrolyzed, hot-water extracts from the CM1 chondrites  
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45 SCO 06043 and MET 01070, the CR1 chondrite GRO 95577, the serpentine blank, and an  
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47 amino acid standard mixture are shown in Fig. 1. Similar UPLC-FD chromatograms were  
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49 obtained for the other meteorite samples. The amino acid peak numbers correspond to the  
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51 designations given in Table 1. Peaks in the meteorite extracts were only identified as amino  
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53 acids if the retention time of the UV fluorescence peak coincided with the corresponding  
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55 mass peak of the OPA/NAC derivatized amino acid standard (see Figs. 1 and 2). Several  
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57 peaks labeled with an 'X' in the fluorescence chromatograms for the meteorite samples and  
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3 the serpentine blank are unidentified primary amine compounds from either the chemical  
4 workup or analytical artifacts that did not interfere with amino acid identification and  
5 quantification (Fig. 1). All of the identified amino acids and corresponding abundances are  
6 given in Tables 2 and 3. The absolute abundances reported were corrected for the  
7 desalting and derivatization recoveries using the D,L-norleucine internal standard; the  
8 recoveries ranged from 60-80% for the meteorite samples studied. Many other amino acid  
9 peaks were detected in the meteorite extracts (e.g. see Murchison in Fig. 2), but were not  
10 identified due to a lack of amino acid standards. All of the Antarctic meteorite samples  
11 revealed high intensity peaks (see Figs. 1 and 2, peak no. 33) corresponding to the C<sub>6</sub>  
12 amino acid  $\epsilon$ -amino-*n*-caproic acid (EACA). This finding is consistent with a previous  
13 analysis of Antarctic CM2 chondrites and the most likely source of EACA is terrestrial  
14 contamination from the Nylon-6 storage bags used during sample collection and curation of  
15 Antarctic meteorites (Glavin et al., 2006). The abundance of EACA in the Antarctic  
16 meteorites ranged from ~ 400 to 51,000 ppb (Table 2), suggesting that exposure of these  
17 meteorites to Nylon-6 contamination is variable. It should be emphasized that the EACA  
18 contamination in these meteorites did not interfere with the ability to detect trace levels of  
19 other amino acids present in the samples. Since EACA is a potential terrestrial contaminant  
20 and its abundance is highly variable in meteorites, we excluded this amino acid from the  
21 total amino acid abundances and the ratio of free to total amino acids reported in Table 2.  
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45 Much lower abundances ranging from 4 to 295 ppb of the amino acids glycine,  $\alpha$ -  
46 alanine,  $\beta$ -alanine (BALA),  $\alpha$ -,  $\beta$ -, and  $\gamma$ -amino-*n*-butyric acid (ABA), and  $\alpha$ -aminoisobutyric  
47 acid ( $\alpha$ -AIB) were found in the aqueously altered chondrites MET 01070, SCO 06043, and  
48 GRO 95577 (Table 2). The abundances of  $\alpha$ -AIB in these type 1 chondrites (30 to 175  
49 ppb) are 100 to 1000 times lower than the  $\alpha$ -AIB abundances found in the CM2 Murchison  
50 and the CR2 EET 92042 (Table 2). A total amino acid abundance of ~16,000 ppb was  
51 measured in LON 94102, which is in the range of amino acid abundances previously  
52 measured in the CM2 chondrites Murchison and LEW 90500 (Table 2). Previous bulk  
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3 oxygen isotope data (Clayton and Mayeda, 1999), as well as hydrogen, carbon, and  
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5 nitrogen isotopic analysis of the insoluble organic matter of LON 94102 suggest that this  
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7 meteorite is among the least altered of the CM chondrites (Alexander et al., 2010).  
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10 From Table 2, the low total amino acid abundances measured in MET 01070 (~710 ppb)  
11 and GRO 95577 (~1,600 ppb) are consistent with previous amino acid analyses of these  
12 type 1 chondrites (Botta et al., 2007; Martins et al., 2007). Although aspartic acid, glutamic  
13 acid, and serine have all been identified previously in these meteorites, we were unable to  
14 confirm their presence above serpentine blank levels (Fig. 1, Table 2). These amino acids  
15 are difficult to identify by mass due to the low abundances and poor ionization of these  
16 compounds during the UPLC-ToF-MS analysis. The CM1 chondrite SCO 06043 has a very  
17 similar amino acid composition and total abundance (~660 ppb) compared to MET 01070  
18 (Table 2). The total amino acid abundance of the CI1 Orgueil (~ 6,700 ppb) is higher than  
19 the CM1 and CR1 meteorites studied (Table 2). These Orgueil results are consistent with  
20 amino acid abundances measured in a previous HPLC-FD analysis of a different fragment  
21 of this meteorite (Ehrenfreund et al., 2001). Overall, the total amino acid abundances found  
22 in these aqueously altered type 1 chondrites are lower than those found in the less altered  
23 CM and CR type 2 and 3 chondrites which ranged from ~9,400 ppb for the CM2 LEW 90500  
24 up to 320,000 ppb for the CR2 chondrite EET 92042 (Table 2). High amino acid  
25 concentrations ranging from 180,000 to 249,000 ppb have previously been reported for the  
26 CR2 meteorites EET 92042 and Graves Nunataks (GRA) 95229 (Martins et al., 2007;  
27 Pizzarello et al., 2008), but these values did not contain many of the C<sub>5</sub> amino acids that we  
28 included in our total amino acid abundance calculations reported in Tables 2 and 3. We  
29 found that the primitive CR3 chondrite QUE 99177 had a similar distribution of amino acids  
30 compared to EET 92042, but the total amino acid abundance of QUE 99177 (~81,000 ppb)  
31 was lower than in EET 92042 (Table 2).  
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57 **UPLC-FD/ToF-MS measurements of C<sub>2</sub> to C<sub>10</sub> amino acids.** We investigated the  
58 masses corresponding to the OPA/NAC derivatives of the acyclic C<sub>2</sub> to C<sub>10</sub> amino alkanolic  
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3 acids in the sample extracts (see Fig. 2). Only trace levels (< 10 ppb) of glycine (peak 9)  
4 and L-alanine (peak 12) in the serpentine procedural blank were measured by UPLC-  
5 FD/ToF-MS (Fig. 2), which indicates that minimal amino acid contamination of the meteorite  
6 samples occurred during the processing procedure. A few small unidentified non-  
7 fluorescent peaks were detected in both the serpentine blank and meteorite sample extracts  
8 and are marked with an 'X' (Fig. 2). Based on the UPLC-FD/ToF-MS data, the Murchison  
9 meteorite contains a very large diversity of amino acids with evidence for C<sub>2</sub> up to C<sub>10</sub>  
10 acyclic amino alkanolic acid isomers detected above background (Fig. 2). The masses  
11 plotted in Figure 3 represent a 0.02 Da window (the peak width at half maximum) centered  
12 around the corresponding theoretical masses of the mono-protonated positive ions of C<sub>2</sub> to  
13 C<sub>10</sub> OPA/NAC amino acid derivatives, and not the masses of the individual unlabeled amino  
14 acids. Based on the total ToF-MS areas in Figure 3 (data not shown), the C<sub>10</sub> amino acids  
15 in Murchison are a factor of 10-20 times less abundant than the area of each of the C<sub>2</sub>  
16 through C<sub>6</sub> amino acids.  
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34 Amino acids with carbon numbers exceeding C<sub>10</sub> may also be present in Murchison, but  
35 these were not observed with confidence above the detection limit of the UPLC-ToF-MS  
36 instrument under the conditions employed. This may either be a result of meteorite  
37 chemistry or biases in the extraction and workup for the more hydrophilic amino acids.  
38 Diamino acids have also been identified in Murchison meteorite extracts using gas  
39 chromatography mass spectrometry (Meierhenrich et al., 2004). OPA derivatives of diamino  
40 acids have been separated using HPLC (Satyanarayana et al., 2001); however, our  
41 OPA/NAC derivatization and UPLC-FD/ToF-MS analytical technique was not optimized for  
42 the identification of these compounds. Nevertheless, these results are consistent with  
43 previous analyses of the Murchison meteorite that have identified over 80 amino acids,  
44 including both cyclic and acyclic primary  $\alpha$ -amino alkanolic and alkanedioic acids with carbon  
45 numbers ranging from C<sub>2</sub> up to C<sub>8</sub> (Cronin and Chang, 1993; Cronin et al., 1981; Cronin and  
46 Pizzarello, 1983; Cronin and Pizzarello, 1986). To our knowledge, this study is the first  
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3 report of amino acids in the Murchison meteorite that exceed C<sub>8</sub>. Based on our UPLC-  
4 FD/ToF-MS data and the numerous possible structural isomers and enantiomers for amino  
5 acids with carbon numbers up to C<sub>10</sub>, the Murchison meteorite likely contains hundreds of  
6 individual amino acids. Unfortunately, due to their low abundance, incomplete separation,  
7 and a lack of standards, the identity of many specific compounds cannot be confirmed.  
8 Future optimization of our LC separation conditions and synthesis of amino acid standards  
9 that are not commercially available will be required to determine the identities of the  
10 detected C<sub>6</sub> to C<sub>10</sub> amino acids in Murchison and other carbonaceous meteorites.  
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21 Several unidentified peaks above background also appeared in the *m/z* 393.15 mass  
22 window of the Murchison extract, corresponding to the C<sub>6</sub> amino acid isomers (Fig. 2);  
23 however, we only confirmed the identity of two of these peaks:  $\epsilon$ -amino-*n*-caproic acid  
24 (EACA, peak 33) and the norleucine internal standard (I.S.) used to estimate amino acid  
25 desalting recoveries. For EET 92042, the intensity of the EACA peak was so high that the  
26 I.S. peak is present, but not easily visible at the scale shown (Fig. 2). A few of the  
27 unidentified C<sub>6</sub> amino acid peaks were also present in the type 1 chondrites Orgueil and  
28 SCO 06043, however the total number of C<sub>6</sub> isomers is clearly less in these carbonaceous  
29 chondrites compared to Murchison (Fig. 2). In contrast to Murchison, the CR chondrites  
30 GRO 95577 and EET 92042 did not contain any evidence for amino acids with more than  
31 six carbons above serpentine blank levels (Fig. 2). This finding is particularly surprising for  
32 EET 92042, which contains the highest reported amino acid abundance of any meteorite to  
33 date (Martins et al., 2007). The Orgueil and SCO 06043 meteorite extracts contain some  
34 evidence for the presence of C<sub>7</sub> and C<sub>8</sub> amino acids above serpentine background levels;  
35 however, the intensity of these amino acid peaks are reduced compared to Murchison (Fig.  
36 2). These data are consistent with earlier studies showing reduced amino acid abundances  
37 in more aqueously altered type 1 carbonaceous chondrites compared to type 2 chondrites  
38 (Botta et al., 2007; Martins et al., 2007).  
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The UPLC-FD/ToF-MS instrument was specifically optimized for separation of the C<sub>5</sub> acyclic amino alcanoic acids and the retention times for all possible C<sub>5</sub> amino acid isomers and enantiomers were identified based on the analysis of standards (Fig. 3). The UPLC-FD/ToF-MS chromatograms centered at  $m/z$  379.1328  $\pm$  0.02 (the peak width at half maximum) correspond to the C<sub>5</sub> amino acids in the standard, the serpentine blank, and the MET 01070, SCO 06043 and GRO 95577 meteorite extracts (Figure 3). Chromatograms showing the C<sub>5</sub> amino acid distributions in Orgueil, Murchison, LON 94102, EET 92042, and QUE 99177 are not shown here, but have been published previously (Glavin and Dworkin, 2009). Although complete separation of all 23 possible C<sub>5</sub> amino acid isomers and enantiomers could not be achieved under the chromatographic conditions employed, all of the C<sub>5</sub> amino acids were accounted for and we observed no interference or co-elution of the D- and L-enantiomers of isovaline, valine, norvaline, and 3-aminopentanoic acid (3-apa) with other C<sub>5</sub> amino acids. As with a number of rare C<sub>5</sub> amino acids, the D- and L-enantiomers of 3-apa were clearly separated (Fig. 3), however they could not be identified due to the lack of optically pure 3-apa standards.

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A comparison of the individual and total abundances of the C<sub>5</sub> amino acids in the meteorites is shown in Table 3. The only C<sub>5</sub> amino acid that could not be quantified in any of the meteorite extracts due to co-elution was 3-amino-3-methylbutanoic acid (3-a-3-mba); therefore, only upper limits are given for 3-a-3-mba. The total abundances of all other C<sub>5</sub> amino acids were determined by comparing the peak areas with standards (Table 3). The CM1 chondrites MET 01070 and SCO 06043 and the CR1 chondrite GRO 95577 had low total C<sub>5</sub> amino acid abundances ranging from 220 to 500 ppb, much lower than found in the CM2 and CR2/3 chondrites (Table 3). The most likely explanation for the low abundances in type 1 chondrites is decomposition of amino acids during aqueous alteration on the meteorite parent body. This is not surprising, since the decomposition of aliphatic moieties in the free and macromolecular organic matter by low temperature chemical oxidation during aqueous alteration on the parent body could result in the low amino acid levels observed in

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3 the type 1 chondrites (Cody and Alexander, 2005; Sephton et al., 2004). Thermal  
4 decomposition by  $\alpha$ -decarboxylation of  $\alpha$ -amino acids to  $\text{CO}_2$  and the corresponding primary  
5 amine during aqueous alteration on the parent body is another possibility (Simmonds et al.,  
6 1972); however, as noted previously, many of the volatile amines produced from  $\alpha$ -  
7 decarboxylation (including the butylamines from the  $\text{C}_5$  amino acids) were not detected in  
8 our UPLC-FD/ToF-MS analyses of the meteorite extracts and may have been lost during  
9 chemical workup.  
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18 **Presence of free vs. bound amino acids.** It has been shown previously that acid  
19 hydrolysis of the water extracts from carbonaceous meteorites will readily release bound  
20 amino acids. These bound amino acids have their amino group chemically protected and  
21 are not subject to OPA/NAC derivatization or exist as a precursor to the amino acid and  
22 would have a different mass and retention time compared to the OPA/NAC labeled amino  
23 acid. The increase in amino acid concentration in the hydrolyzed extracts observed in many  
24 carbonaceous chondrites is believed to be a result of derivatives and/or acid-labile  
25 precursors that are converted to amino acids after acid hydrolysis. For example, in the  
26 Murchison meteorite most of the acid-labile precursors are low molecular weight derivatives  
27 including mono- and dicarboxylic acid amides, hydroxy acid amides, lactams,  
28 caboxylactams, N-acetylamino acids, and substituted hydantoins (Cooper and Cronin, 1995;  
29 Cronin, 1976a, b).  
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45 Analysis of the non-hydrolyzed water extracts of the CM1s SCO 06043 and MET 01070,  
46 and the CR1 GRO 95577 showed a similar distribution of free amino acids compared to the  
47 total (free + bound) amino acids in the acid-hydrolyzed extracts. From the UPLC-FD/ToF-  
48 MS data (not shown), we calculated the total abundance of free amino acids (not including  
49 EACA) in the non-hydrolyzed extracts (Table 2). The results are similar to previous  
50 analyses of the CM2 chondrites Murchison, LEW 90500, and ALH 83100 (Glavin et al.,  
51 2006) and the CM1 chondrites MET 01070 and ALH 88045 (Botta et al., 2007) which  
52 reported that free amino acids on average represent ~40 to 50% of the total acid-  
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3 hydrolyzable amino acid abundance. In contrast, Pizzarello and coworkers found that the  
4 abundance of amino acids in the relatively unaltered Antarctic CR2 chondrite GRA 95229  
5 hot water extract did not increase substantially after acid hydrolysis, with free amino acids  
6 representing over 70% of the total amino acid abundance on average (Pizzarello et al.,  
7 2008). We also measured a high relative abundance of free amino acids (free/total = 64%)  
8 in another Antarctic CR2 chondrite EET 92042 (Table 2). It is possible that bound amino  
9 acids are more resistant to oxidation during aqueous alteration than free amino acids,  
10 resulting in a lower free/total amino acid ratio in more aqueously altered chondrites.  
11 However, if this were true we would expect a higher ratio of free to total amino acids in the  
12 less altered CR3 chondrite QUE 99177 compared to the CR2 EET 92042, and this was not  
13 observed (Table 2). Amino acid analyses of the non-hydrolyzed water extracts from  
14 additional carbonaceous chondrites that experienced minimal aqueous alteration will be  
15 required to further investigate this hypothesis.  
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### 31 **Unusual Amino Acids and Enantiomeric Measurements**

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33 The presence of non-protein amino acids in meteorites that are not common on Earth,  
34 such as  $\alpha$ -AIB and isovaline, has often been used to argue that these amino acids are  
35 indigenous to the meteorites and not terrestrial contaminants. In fact, the majority of the  
36 80+ amino acids identified in the Murchison meteorite are either rare or nonexistent in  
37 terrestrial proteins (Cronin and Chang, 1993). The high relative abundances of  $\alpha$ -AIB and  
38 isovaline in Murchison and other Antarctic CM2 and CR2 chondrites (Table 2) compared to  
39 the most common protein amino acids provide additional support of an extraterrestrial origin  
40 for these amino acids. For the Antarctic meteorites, the Antarctic ice is a very unlikely  
41 source of these two non-protein amino acids. Amino acid analysis of Antarctic ice from the  
42 Allan Hills region revealed only ppb levels of the protein amino acids glycine, L-aspartic acid,  
43 L-serine, L-glutamic acid, and L-alanine and no evidence for  $\alpha$ -AIB and isovaline above the  
44 2 parts-per-trillion (ppt) detection limit (Bada et al., 1998; McDonald and Bada, 1995). Trace  
45 levels of  $\alpha$ -AIB (46 ppt) were recently identified in an Antarctic ice sample from LaPaz using  
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3 very similar methods to those reported here, although the source of this amino acid remains  
4 uncertain (Glavin et al., 2006). In addition, contamination of the Murchison meteorite by  
5 terrestrial  $\alpha$ -AIB or isovaline from the landing site environment is also unlikely since these  
6 amino acids were not identified above the 10 ppb detection limit in surface and subsurface  
7 (20-30 cm) soil samples collected from the Murchison fall site (Kvenvolden, 2000). For the  
8 Murchison meteorite, stable isotope analyses of  $\alpha$ -AIB and isovaline have shown  
9 enrichments of D,  $^{15}\text{N}$ , and  $^{13}\text{C}$  that fall well outside the range for terrestrial amino acids and  
10 indicate an extraterrestrial origin (Engel and Macko, 1997; Engel et al., 1990; Pizzarello and  
11 Huang, 2005; Pizzarello et al., 2004b).

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13 Although  $\alpha$ -AIB and isovaline are not among the 20 genetically encoded amino acids,  
14 they are present in a variety of terrestrial fungal polypeptides (named peptaibols) that are  
15 biosynthesized through nonribosomal enzymatic pathways (Brückner et al., 2009; Kleinkauf  
16 and Von Dohren, 1996); therefore, presence of these amino acids alone does not prove an  
17 extraterrestrial origin. Additional data, such as the distribution of the entire suite of amino  
18 acids, chirality, or stable isotopes are required to completely rule out a biological source of  
19  $\alpha$ -AIB and isovaline in meteorites. This becomes particularly important (and difficult) for  
20 meteorites with extremely low abundances of  $\alpha$ -AIB and isovaline and where only a limited  
21 mass of sample is available for analysis (e.g. MET 01070, SCO 06043, and GRO 95577).  
22 For these meteorites, compound-specific stable isotopic measurements of  $\alpha$ -AIB and  
23 isovaline could help determine the origin of these amino acids. However, given the low  
24 amino acid abundances in the type 1 chondrites, this analysis is particularly challenging, if  
25 not impossible, with current instrumentation and would require a significant fraction of the  
26 available sample. For example, recent analysis of material returned from the Stardust  
27 mission showed the ability to perform compound-specific carbon isotope analysis from 0.7  
28 nmol of glycine (Elsila et al., 2009). A similar isotopic analysis of D-isovaline would require a  
29 minimum of  $\sim 0.25$  nmol on column. Based on this detection limit and the measured  
30 abundances of D-isovaline in SCO 06043 and GRO 95577 of 3.3 ppb and 14.6 ppb (Table

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3 3), we would require 29 g and 6.3 g of meteorite, respectively. Given the total recovered  
4 mass of 27.6 g for SCO 06043 and 106.2 g of GRO 95577, there is not enough material  
5 available for isotopic measurement of D-isovaline in SCO 06043 and roughly 6% of the total  
6 available mass of GRO 95577 would have to be requested for destructive analysis. Given  
7 that GRO 95577 is the only known CR1 carbonaceous chondrite, obtaining this much  
8 sample appears prohibitive. Therefore, until the sensitivity of analytical techniques for  
9 isotopic measurements can be improved, amino acid enantiomeric measurements and  
10 comparisons of the amino acid distributions with possible terrestrial sources are the only  
11 ways to establish the origin of amino acids in these type 1 chondrites.  
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23 **D/L enantiomeric ratios.** Chirality is another important measurement tool to help  
24 discriminate between biotic and abiotic origins of amino acids in meteorites, even if present  
25 at low abundances. On Earth, biology almost exclusively uses L-amino acids, whereas  
26 abiotic syntheses result in equal mixtures of L- and D-amino acids. Furthermore, protein  
27 amino acids will readily racemize (approach D/L = 1) on geologically short time-scales  
28 (Bada, 1972). Therefore, if we assume that the protein amino acids found in carbonaceous  
29 chondrites were racemic (D/L ~ 1) prior to their fall to Earth, then their current D/L ratios can  
30 be used to determine the relative degree of terrestrial L-amino acid contamination they have  
31 experienced on Earth. The enantiomeric ratios (D/L) for aspartic and glutamic acids, serine,  
32 alanine,  $\beta$ -ABA, valine, norvaline, and isovaline are reported in Table 4 when abundances  
33 permitted quantitation.  
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47 The enantiomeric ratios measured for the protein amino acids (italicized in Table 4) in  
48 the CM chondrites Murchison (USNM 6650), LEW 90500, and LON 94102, and the CR  
49 chondrites EET 92042 and QUE 99177 were mostly found to be racemic (D/L ~ 0.9 to 1.0)  
50 within analytical errors ( $\pm$  0.05 to 0.3, Table 4), indicating that these amino acids are  
51 primarily abiotic in origin and not the result of terrestrial contamination. However, a lower  
52 D/L ratio for glutamic acid of 0.57 measured in Orgueil (Table 4) may indicate that some  
53 terrestrial L-glutamic acid is present in this meteorite. The relatively low D/L ratios of  
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3 aspartic and glutamic acids in Murchison (USNM 5453) of 0.67 and 0.61, indicate that this  
4 fragment has a higher degree of terrestrial L-protein amino acid contamination than the  
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6 fragment has a higher degree of terrestrial L-protein amino acid contamination than the  
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8 other Murchison sample used in this study (USNM 6650) which has much higher D/L ratios  
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10 for aspartic and glutamic acids ( $D/L = 0.91$  and  $0.96$ , respectively) (Table 4). Despite these  
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12 significant differences in D/L aspartic and glutamic acid ratios in the two samples of  
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14 Murchison, the non-protein amino acids  $\beta$ -ABA, norvaline, and isovaline in these samples  
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16 had identical D/L ratios within errors (Table 4), suggesting that terrestrial contamination is  
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18 an unlikely source for these non-protein amino acids. Although aspartic and glutamic acids  
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20 were not detected in MET 01070, SCO 06043, and GRO 95577, the D/L ratio for alanine  
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22 measured in these meteorites was racemic ( $D/L \sim 1$ ) within analytical uncertainties (Table  
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24 4), indicating both an extraterrestrial origin for alanine and low levels of terrestrial L-amino  
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26 acid contamination in these meteorites.  
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29 Although the D- and L-enantiomers of valine were separated using the LC conditions  
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31 employed, there was a relatively large unidentified non-fluorescent co-eluting isobaric  
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33 contaminant at approximately the same retention time as L-valine in some of the analyses  
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35 including the serpentine blank (Fig. 3), which may have led to erroneously high L-valine  
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37 abundances, and hence lower D/L valine ratios ( $\sim 0.2$  to  $0.5$ ) for Orgueil, Murchison (USNM  
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39 5453), SCO 06043 and GRO 95577 (Table 4). Therefore, lower limits were placed on these  
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41 ratios. The non-protein amino acids  $\beta$ -ABA and norvaline detected in these meteorites were  
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43 racemic ( $D/L \sim 1$ ) within analytical error, indicating an abiotic origin for these amino acids  
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45 (Table 4). One notable exception was the enantiomeric ratio of the non-protein amino acid  
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47 isovaline, which ranged from 0.61 to 0.73 in Orgueil, SCO 06043, and Murchison (Table 4).  
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49 Terrestrial isovaline contamination of these meteorites after falling to Earth is highly unlikely  
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51 since no isovaline was detected in any of the relevant collection site environments or any of  
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53 the laboratory procedural blanks. Although some fungal peptide contamination of both D-  
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55 and L-isovaline cannot be completely ruled out, it is important to note that isovaline is most  
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57 commonly found in the D-configuration in fungal peptides (Brückner et al., 2009; Keller et  
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3 al., 1990), which would increase (and not decrease) the D/L isovaline ratio. In addition,  
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5 many common fungal peptides also contain L-alanine (D/L ~ 0.06) at similar abundances to  
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7 isovaline (Brückner et al., 2009); therefore, if any significant fungal peptide contamination of  
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9 the meteorites had occurred, the D/L alanine ratios in these meteorites should have been  
10  
11 affected as well. However, the D/L alanine ratios in all of the meteorites analyzed were  
12  
13 racemic within analytical errors showing no evidence for significant terrestrial L-alanine  
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15 contamination (Table 4). The detection of non-racemic D/L isovaline ratios corresponding  
16  
17 to L-isovaline excesses in Murchison and other carbonaceous chondrites have been  
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19 reported previously (Cronin and Pizzarello, 1997; Glavin and Dworkin, 2009; Pizzarello and  
20  
21 Cronin, 2000; Pizzarello et al., 2008; Pizzarello et al., 2003). Possible mechanisms for the  
22  
23 origin of the L-isovaline asymmetry observed in some carbonaceous chondrites are  
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25 discussed below.  
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### 29 **Variability in Amino Acid Distributions and Formation Pathways**

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32 Comparisons of the relative amino acid abundances between different classes of  
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34 carbonaceous meteorites can be a powerful way to elucidate synthetic pathways for the  
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36 formation of amino acids and to understand the trends between amino acid composition and  
37  
38 degree of aqueous alteration on the meteorite parent body (Botta et al., 2007; Glavin et al.,  
39  
40 2006; Martins et al., 2007). For example, the CI1 chondrites Orgueil and Ivuna were found  
41  
42 to have a relatively simple distribution of amino acids (predominantly  $\beta$ -alanine) compared to  
43  
44 the complex distribution of  $\alpha$ -amino acids (predominantly glycine, isovaline, and  $\alpha$ -AIB)  
45  
46 found in the CM2 meteorites Murchison and Murray believed to have formed by Strecker  
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48 synthesis (Ehrenfreund et al., 2001; Peltzer et al., 1984). Since  $\beta$ -alanine is not produced  
49  
50 directly by the Strecker-cyanohydrin pathway, it was argued that this amino acid must have  
51  
52 formed by an alternate synthetic route (Ehrenfreund et al., 2001), such as Michael addition  
53  
54 of ammonia to cyanoacetylene (Miller, 1957). Based on this amino acid evidence it was  
55  
56 concluded that the Orgueil and Ivuna meteorites must have originated on a chemically  
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58 distinct parent body from the CMs, possibly an extinct comet (Ehrenfreund et al., 2001).  
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4 A subsequent comparison of the relative abundances of  $\beta$ -alanine and  $\alpha$ -AIB in Orgueil  
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6 with four CM2 meteorites showed that the relative abundances of both  $\beta$ -alanine and  $\alpha$ -AIB  
7  
8 tracked with the degree of aqueous alteration of the meteorites, with the more altered  
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10 Orgueil meteorite having the highest  $\beta$ -alanine and lowest  $\alpha$ -AIB relative abundances  
11  
12 (Glavin et al., 2006). Although a similar trend was also observed for the CR1 chondrite  
13  
14 GRO 95577 and the less altered CR2s EET 92042 and GRA 955229 (Martins et al., 2007),  
15  
16 there was no clear trend with aqueous alteration found in relative  $\alpha$ -AIB and  $\beta$ -alanine  
17  
18 abundances in a previous study of CM1 and CM2 chondrites (Botta et al., 2007). One  
19  
20 potential problem with comparing amino acid abundance data from these different studies is  
21  
22 that the meteorites were analyzed in different laboratories at different times using different  
23  
24 instrumentation and different peak integration procedures. In addition, sample  
25  
26 heterogeneity must also be accounted for when drawing comparisons between laboratories.  
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28 Here for the first time we compare the relative abundances of amino acids collected on a  
29  
30 suite of CI, CM, and CR carbonaceous chondrites that cover the entire range of aqueous  
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32 alteration (petrographic types 1-3) using the identical extraction, analytical, and quantitation  
33  
34 techniques in the same laboratory.  
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38 **Relative abundance of  $\beta$ -alanine as an indicator for aqueous alteration.** A  
39  
40 comparison of the relative molar abundances (glycine = 1.0) of  $\alpha$ -alanine (D+L  
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42 enantiomers),  $\beta$ -alanine,  $\alpha$ -AIB, and isovaline (D+L enantiomers) measured in nine different  
43  
44 carbonaceous chondrites is shown in Fig. 4. The order of carbonaceous chondrites in Fig. 4  
45  
46 are based on the approximate degree of aqueous alteration inferred from mineralogical and  
47  
48 isotopic evidence with the most altered CI1 Orgueil and other type 1 chondrites on the left  
49  
50 and the most primitive, least altered CR chondrites on the right (Kallemeyn et al., 1994;  
51  
52 Zolensky and McSween, 1988; Zolensky and Browning, 1994). The relative degree of  
53  
54 aqueous alteration among carbonaceous chondrites within the same group and of the same  
55  
56 petrologic type (e.g. the difference in degree of alteration between the three CM2 chondrites  
57  
58 shown in Fig. 4) is less certain, although recent isotopic analyses of the IOM in LON 94102  
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3 suggests that this meteorite is less altered than Murchison (Alexander et al., 2010) Based  
4 on the data in Fig. 4, there appears to be a correlation between the relative abundance of  $\beta$ -  
5 alanine and degree of aqueous alteration in the meteorites studied. The relative abundance  
6 of  $\beta$ -alanine increases from a  $\beta$ -alanine/glycine ratio of  $\sim 0.1$  to  $0.2$  for the least altered CR  
7 chondrites EET 92042 and QUE 99177 up to  $\sim 2.7$  for the most aqueously altered CI1  
8 Orgueil. We also find that within the corresponding CM and CR chondrite groups, the type 1  
9 chondrites have a much higher relative abundance of  $\beta$ -alanine compared to the less altered  
10 type 2 chondrites (Fig. 4). Thus, in addition to mineralogy and isotopic measurements, the  
11 relative abundance of  $\beta$ -alanine should also be considered when comparing the degree of  
12 aqueous alteration between carbonaceous chondrites.  
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26 In contrast to  $\beta$ -alanine, there do not appear to be any strong trends in relative  $\alpha$ -  
27 alanine,  $\alpha$ -AIB, and isovaline abundances with aqueous alteration (Fig. 4). However, it is  
28 worth noting that the relative abundances of  $\alpha$ -AIB and isovaline tend to be higher in  
29 carbonaceous chondrites that experienced less aqueous alteration (Fig. 4). The relative  
30 abundance of  $\alpha$ -alanine ranged from  $0.2$  for Orgueil up to  $1.3$  for EET 92042. The high  $\alpha$ -  
31 alanine abundance in EET 92042 was not observed in QUE 99177 and cannot be explained  
32 by a large contribution of terrestrial L-alanine contamination since the D/L alanine ratio in  
33 EET 92042 was  $0.97$  (Table 4). Assuming that Strecker synthesis was the primary  
34 mechanism for the formation of  $\alpha$ -alanine in EET 92042 and QUE 99177, it is possible that  
35 the parent body of EET 92042 had a higher ratio of acetaldehyde to formaldehyde (the  
36 Strecker precursors for formation of  $\alpha$ -alanine and glycine, respectively) compared to QUE  
37 99177.  
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53 The weakly altered CM2 chondrite LON 94102 is clearly different compared to the other  
54 more altered CM2 chondrites (or any other meteorite analyzed in this study) and appears to  
55 follow the opposite trend in relative  $\alpha$ -AIB and isovaline abundances, with lower  $\alpha$ -  
56 AIB/glycine and isovaline/glycine ratios compared to Murchison and LEW 90500 (Fig. 4).  
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59  
60 One possibility for the low  $\alpha$ -AIB and isovaline abundances in LON 94102 is that this

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3 meteorite originated from a chemically distinct parent body from other CM chondrites that  
4 was depleted in acetone and 2-butanone, the ketone precursors required for the formation  
5 of  $\alpha$ -AIB and isovaline, respectively, by Strecker-cyanohydrin synthesis. However, based on  
6 a variety of petrologic properties of eleven different CM chondrites that show similar bulk  
7 compositions, it has been suggested that textural differences seen among individual CM  
8 chondrites may reflect progressive alteration of similar hypothetical CM3 starting materials in  
9 different regions of the same parent body (Rubin et al., 2007). Therefore, we cannot rule  
10 out the possibility that the relative abundances of amino acids in LON 94102 are typical for  
11 weakly altered CM chondrites. Comparing the relative distribution of amino acids in LON  
12 94102 to other weakly altered CM2 chondrites, such as QUE 97990 and Yamato (Y) 791198  
13 (Chizmadia and Brearley, 2008; Maeda et al., 2009; Metzler et al., 1992; Rubin et al., 2007),  
14 may help confirm this hypothesis. Previous amino acid analyses found more than ten times  
15 the relative molar abundance of  $\alpha$ -AIB ( $\alpha$ -AIB/glycine = 2.7 to 3.5) in Y-791198 (Shimoyama  
16 et al., 1985; Shimoyama and Ogasawara, 2002) compared to what we observed in LON  
17 94102 in this study (Fig. 4). However, it should be pointed out that we have not yet  
18 analyzed Y-791198 using the same UPLC-FD/ToF-MS technique used for LON 94102. To  
19 our knowledge, no amino acid measurements have been reported yet for QUE 97990.

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41 The similarity in the distributions of  $\alpha$ -alanine,  $\beta$ -alanine,  $\alpha$ -AIB, and isovaline among  
42 type 1 chondrites across three separate carbonaceous chondrite groups (CI, CM, and CR)  
43 that are distinct from the less altered type 2 and type 3 chondrites (Fig. 4), may point toward  
44 degree of aqueous alteration, rather than unique parent body chemical compositions, as an  
45 explanation for the amino acid variations observed in these chondrites. Although the high  
46 relative abundance of  $\beta$ -alanine and the simple distribution of amino acids found in the CI1  
47 chondrites Orgueil and Ivuna has been attributed to a distinct chemical composition on the  
48 CI parent body compared to the CM parent body (Ehrenfreund et al., 2001), the amino acid  
49 data from this study suggests that this might not be the case, and extended aqueous  
50 alteration, rather than differences in the chemical composition of the parent body, may be  
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3 responsible for the high relative abundance of  $\beta$ -alanine over other  $\alpha$ -amino acids in CI1  
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5 chondrites.  
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8 **Possible formation pathways for amino acids found in meteorites.** As stated  
9  
10 above,  $\beta$ -amino acids cannot be formed by the Strecker mechanism, but can be synthesized  
11  
12 by Michael addition of ammonia to  $\alpha,\beta$ -unsaturated nitriles followed by reduction/hydrolysis  
13  
14 in water, a mechanism that was proposed for the formation of  $\beta$ -alanine in spark discharge  
15  
16 experiments (Miller, 1957). For example, the formation of  $\beta$ -alanine by Michael addition of  
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18 ammonia to cyanoacetylene during aqueous alteration on the parent bodies of CI, CM, and  
19  
20 CR chondrites may have occurred. The reaction of HCN,  $\text{NH}_3$ , and carbonyl compounds  
21  
22 (aldehydes and ketones) leads to the formation of  $\alpha$ -amino, imino, and  $\alpha$ -hydroxy acids in  
23  
24 aqueous solution via the Strecker-cyanohydrin pathway and all of these expected Strecker  
25  
26 byproducts have been identified in the Murchison meteorite (Lerner and Cooper, 2005;  
27  
28 Peltzer and Bada, 1978; Peltzer et al., 1984). Alternative sources for the amino acids in  
29  
30 carbonaceous chondrites such as irradiation of ices formed in the interstellar medium (Elsila  
31  
32 et al., 2007) and Fischer-Tropsch Type (FTT) catalysis (Hayatsu et al., 1971) that do not  
33  
34 require aqueous activity for their formation in the parent body have also been proposed.  
35  
36 Several  $\alpha$ -amino acids and one  $\beta$ -amino acid ( $\beta$ -alanine) have been synthesized by Fischer-  
37  
38 Tropsch, however, no  $\gamma$ - or  $\delta$ -amino acids were reported (Yoshino et al., 1971). Additional  
39  
40 studies of the Fischer-Tropsch based synthesis of amino acids are clearly needed using  
41  
42 modern analytical methods to ascertain the distribution of amino acids produced by this  
43  
44 process. Some meteoritic  $\gamma$ - and  $\delta$ -amino acids found in Murchison could be produced from  
45  
46 the hydrolysis of lactams that have been identified in this meteorite (Cooper and Cronin,  
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48 1995; Pizzarello et al., 2006). Although FTT products are expected to be predominantly  
49  
50 straight-chain hydrocarbons (Hayatsu and Anders, 1981), the opposite is found for the  
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52 carbonaceous chondrites where branched-carbon chain amino acid isomers dominate over  
53  
54 straight-carbon chain isomers (e.g. Murchison, LEW 90500, EET 92042, QUE 99177, GRA  
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56 95229) or are approximately equal abundance (Orgueil, MET 01070, SCO 06043, GRO  
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3 95577, LON 94102) (see Fig. 5b and MARTINS et al., 2007). The predominance of branched  
4 over straight-chain amino acid isomers has also been observed in Murchison and suggests  
5 that ion and radical stability was important during carbon chain formation of the interstellar  
6 amino acid precursors (Cronin and Chang, 1993; Herbst, 1995). Identifying variations in the  
7 structure of C<sub>5</sub> amino acids in meteorites has also been used to help constrain parent body  
8 processes for the formation of these compounds (Glavin et al., 2010).  
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### 16 **Variations in C<sub>5</sub> amino acid structure between carbonaceous chondrite groups.**

17 For this study, we calculated the relative molar abundances of the C<sub>5</sub> amino acid isomers in  
18 each meteorite as a function of amine position ( $\alpha$ -,  $\beta$ -,  $\gamma$ , or  $\delta$ -) and valeric acid carbon chain  
19 structure (*n*-, *sec*-, *iso*-, or *tert*-) to determine if any synthetic relationships or chemical  
20 patterns exist between the CI, CM, and CR chondrite groups. These results, normalized to  
21 the total number of possible structural isomers, are illustrated in Fig. 5a and 5b. The  
22 dashed line corresponding to a relative abundance of 1 indicates the expected ratio if there  
23 was an equal probability of forming all of the C<sub>5</sub> amino acid isomers. Therefore, for the  
24 purposes of this discussion, enhanced and depleted relative abundances refer to values that  
25 fall above and below the dashed line, respectively. The high relative abundances of C<sub>5</sub>  $\alpha$ -  
26 amino acids compared to  $\beta$ -,  $\gamma$ -, and  $\delta$ -amino acids in the CM chondrites Murchison and  
27 LEW 90500 and the CR chondrites EET 92042 and QUE 99177 provide strong evidence  
28 that the C<sub>5</sub>  $\alpha$ -amino acids in these meteorites were formed by Strecker-cyanohydrin  
29 synthesis (Fig. 5a). However, the opposite trend is observed for the more aqueously altered  
30 type 1 chondrites Orgueil, SCO 06043, MET 01070 and GRO 95577, which have a much  
31 higher relative abundance of the  $\gamma$ - and  $\delta$ -amino acids compared to the  $\alpha$ - and  $\beta$ -amino  
32 acids (Fig. 5b). One possibility is that more extensive aqueous alteration in the type 1  
33 carbonaceous chondrites results in an increase in the rate of hydrolysis of amino acid  
34 precursors yielding elevated levels of  $\gamma$ - and  $\delta$ -amino acids in these carbonaceous  
35 chondrites. As mentioned earlier, some meteoritic  $\gamma$ - and  $\delta$ -amino acids found in Murchison  
36 could have been produced directly from the hydrolysis of lactams present in the meteorite  
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3 (Cooper and Cronin, 1995; Pizzarello et al., 2006). It is possible that the high abundances  
4 of C<sub>5</sub>  $\gamma$ - and  $\delta$ -amino acids detected in type 1 carbonaceous chondrites could have formed  
5 from lactams by a similar process; however, lactams contain secondary (and not primary)  
6 amines and we are unable to confirm the presence of these compounds in the non-  
7 hydrolyzed meteorite extracts using our analytical method.  
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14 In general, the relative abundances of the C<sub>5</sub>  $\beta$ -amino acids in the type 1 chondrites  
15 seem to be slightly higher than in the type 2 and type 3 chondrites (Fig. 5a); however, unlike  
16 the C<sub>3</sub> amino acid  $\beta$ -alanine, we see no clear trend in C<sub>5</sub>  $\beta$ -amino acid abundance with  
17 degree of alteration in the meteorites analyzed. LON 94102 has a unique C<sub>5</sub> amino acid  
18 distribution compared to the other CM2 chondrites and is only slightly enriched in  $\alpha$ -,  $\gamma$ -, and  
19  $\delta$ -amino acids compared to  $\beta$ -amino acids. One possible explanation for this observation is  
20 that LON 94102 experienced more aqueous alteration than Murchison and LEW 90500  
21 leading to an increased relative abundance of  $\gamma$ - and  $\delta$ -amino acids derived from the  
22 hydrolysis of lactams in LON 94102. However, bulk oxygen isotope data (see Fig. 4 of  
23 CLAYTON and MAYEDA, 1999) and recent isotopic analysis of the insoluble organic matter in  
24 LON 94102 indicates that this meteorite is much less, not more, altered than Murchison and  
25 LEW 90500 (Alexander et al., 2010). Therefore, an alternate explanation is that LON 94102  
26 originated on a chemically distinct parent body from these CM2 chondrites.  
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43 A comparison of the relative abundances of the C<sub>5</sub> amino acids as a function of carbon-  
44 chain (valeric acid) structure yields additional insight into the mechanism of formation of  
45 these amino acids. As illustrated in Fig. 5b, we find a predominance of branched-chain  
46 (*sec*- and *iso*-) over straight-chain (*n*-) C<sub>5</sub> amino acids in the CMs Murchison and LEW  
47 90500 and the CRs EET 92042 and QUE 99177. However, this is not the case for the  
48 aqueously altered type 1 chondrites Orgueil, MET 01070, SCO 06043, and GRO 95577  
49 where branched-chain amino acids are not favored over the straight-chain acids and roughly  
50 equal abundances of the *n*-, *sec*-, and *iso*-valeric acid structures are observed with only a  
51 slight depletion of *tert*-valeric acid (Fig. 5b). Once again, we find that LON 94102 has a  
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3 distinct distribution of C<sub>5</sub> amino acids compared to other CM2 and CR2 chondrites with  
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5 respect to the valeric acid chain structure with a distribution that is very similar to the type 1  
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7 chondrites. Since LON 94102 has been classified as a CM2, the relative distribution of C<sub>5</sub>  
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9 amino acids in this meteorite is puzzling. With the exception of LON 94102, we conclude  
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11 that aqueous alteration on the parent bodies of CM and CR chondrites changes the  
12  
13 distributions of C<sub>5</sub> amino acids from predominately  $\alpha$ -amino acids to predominately  $\gamma$ - and  $\delta$ -  
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15 amino acids with relative enrichments in the *n*- and *tert*-valeric acid structures. Although at  
16  
17 present we have not identified the mechanism(s) responsible for the variations seen in the  
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19 C<sub>5</sub> amino acid distributions for these carbonaceous chondrites, parent body alteration seems  
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21 to be the most reasonable explanation.  
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#### 24 25 **Evidence for Amplification of L-Isovaline Asymmetry during Aqueous Alteration** 26

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28 Although nearly racemic mixtures (D/L ~ 0.9 to 1.0) were measured for several chiral  
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30 protein and non-protein amino acids in at least one pristine sample of Murchison (Table 4),  
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32 small L-enantiomeric excesses ranging from 1.0 to 9.2% have previously been measured for  
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34 several non-protein  $\alpha$ -dialkyl amino acids in the Murchison and Murray meteorites including  
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36 2-amino-2,3-dimethyl-pentanoic acid, isovaline,  $\alpha$ -methylisoleucine,  $\alpha$ -methylnorvaline,  $\alpha$ -  
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38 methylnorleucine, and  $\alpha$ -methylvaline (Cronin and Pizzarello, 1997; Pizzarello and Cronin,  
39  
40 2000; Pizzarello et al., 2003). In contrast to  $\alpha$ -hydrogen amino acids, these  $\alpha$ -dialkyl amino  
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42 acids are highly resistant to racemization (conversion of one enantiomer to the other);  
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44 therefore, small enantiomeric excesses should be preserved for these amino acids since the  
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46 time of formation. However, the detection of enantiomeric enrichments of ~12-14% in the  
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48 C<sub>6</sub> amino acid diastereomers L-isoleucine and D-alloisoleucine in the Antarctic CR meteorite  
49  
50 GRA 95229 that may have derived from asymmetry in their C<sub>5</sub> aldehyde precursor  
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52 (Pizzarello et al., 2008), suggests that aqueous alteration did not lead to extensive  
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54 racemization of the aldehydes on the parent body. L-isovaline excesses ranging from 0 to  
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56 15.2% were measured in several different samples of the Murchison meteorite from the  
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58 Smithsonian Institution (USNM 5341) and Arizona State University (ASU 828 and '70)  
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3 collections using gas chromatography mass spectrometry with significant variations between  
4 meteorite fragments (Pizzarello et al., 2003). The large L-isovaline excesses in Murchison  
5 were later confirmed in our laboratory by LC-FD/ToF-MS analysis of another sample of  
6 Murchison (USNM 5560), and L-isovaline excesses up to 15% were also found for the first  
7 time in the CI Orgueil (Glavin and Dworkin, 2009). The large L-isovaline excesses observed  
8 in Murchison and Orgueil would not likely have obtained their asymmetry by the same  
9 mechanisms as L-isoleucine and D-alloisoleucine, since the Strecker precursor for isovaline  
10 (2-butanone), is achiral.  
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21 It has been proposed that the L-enantiomeric excesses of the  $\alpha$ -dialkyl amino acids  
22 found in Murchison and Murray could be the result of asymmetric photolytic decomposition  
23 of the amino acids or their precursors by UV circularly polarized light (UV-CPL) in the  
24 presolar cloud (Bonner and Rubenstein, 1987). Polarized photons of synchrotron radiation  
25 derived from strong magnetic fields around neutron stars or scattered from interstellar dust  
26 in star forming regions have also been suggested as alternative sources of the amino acid  
27 asymmetry found in meteorites (Bailey et al., 1998; Bonner, 1991; Fukue et al., 2009).  
28 Chiral amino acid symmetry breaking and enantiomeric enrichment of 2-3% by UV CPL  
29 photodestruction of solid leucine has been demonstrated in the laboratory (Flores et al.,  
30 1977). Similar experiments have also shown that small 0.5% excesses of D- and L-alanine  
31 can be produced by R-UV-CPL and L-UV-CPL, respectively (Takano et al., 2007).  
32 However, in order to produce the largest  $\geq 15\%$  left handed excess previously reported for  
33 isovaline in Murchison (Glavin and Dworkin, 2009; Pizzarello et al., 2003), more than 99% of  
34 the compound would have to be destroyed by UV-CPL (Flores et al., 1977). Therefore, UV-  
35 CPL as the sole mechanism for enantiomeric enrichment seems unlikely given the high  
36 concentrations of isovaline found in Murchison and Murray and chemical evidence that  
37 isovaline and other  $\alpha$ -amino acids formed by Strecker-cyanohydrin synthesis during  
38 aqueous alteration in the CM parent body (Peltzer and Bada, 1978), thus shielded from  
39 circularly polarized radiation. It should also be noted that similarities in the carbon isotopic  
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3 variation with chain length in Murchison observed for  $\alpha$ -methyl- $\alpha$ -amino acids including  
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5 isovaline and for  $\alpha$ -hydrogen amino acids provides additional evidence that these amino  
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7 acids did not form by distinct chemical pathways (Pizzarello et al., 2004a). Other possible  
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9 mechanisms for breaking amino acid symmetry include physical processes, such as  
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11 crystallization (Kondepudi et al., 1990). Asymmetric autocatalysis (the Soai reaction)  
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13 provided the first experimental evidence that small initial imbalances can be amplified under  
14  
15 aqueous conditions to produce large enantiomeric excesses up to 90% (Soai et al., 1995).  
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17 Recently, it has also been shown that several protein amino acids that form racemic  
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19 compounds and have nonzero eutectic values can result in significant asymmetric  
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21 amplification from a small initial imbalance due to the equilibrium solid-liquid phase behavior  
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23 of amino acids in solution (Blackmond, 2004; Klussmann et al., 2006). The eutectic value  
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25 of isovaline has not been reported and it remains unclear whether or not this amplification  
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27 mechanism is even relevant under parent body conditions. Nevertheless, since the more  
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29 aqueously altered CM and CI chondrites Murchison and Orgueil have high L-isovaline  
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31 enantiomeric excesses ( $L_{ee} \sim 15$  to 19%), and the most primitive unaltered Antarctic CR  
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33 chondrites EET 92042 and QUE 99177 have very small or no L-isovaline enrichment (Glavin  
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35 and Dworkin, 2009; Pizzarello et al., 2008), supports the idea that aqueous alteration played  
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37 some role in the amplification of L-isovaline excesses on the meteorite parent body.  
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43 Although the current isovaline enantiomeric data for these meteorites suggest that there  
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45 may be a relationship between enantiomeric enrichment and degree of aqueous alteration  
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47 (Glavin and Dworkin, 2009), additional isovaline enantiomeric measurements are needed to  
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49 firmly establish a trend, especially for the type 1 carbonaceous chondrites. In order to  
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51 further investigate the source of L-isovaline excesses in carbonaceous chondrites, we  
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53 expanded our search for isovaline asymmetry to include CM and CR aqueously altered type  
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55 1 chondrites. We found that the CM1 SCO 06043 and the CR1 GRO 95577 carbonaceous  
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57 chondrites both had significant L-isovaline excesses, with values of  $16.5 \pm 7.5\%$  and  $11.0 \pm$   
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59  $7.2\%$ , respectively (Table 5). These excesses agree within uncertainties to previous  
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3 measurements of isovaline in the CI1 Orgueil ( $L_{ee} = 15.2 \pm 4.0\%$ ). Isovaline was not  
4 detected in the CM1 MET 01070 above the 0.5 ppb level (Table 3). Similarly large L-  
5 isovaline excesses of  $17.2 \pm 6.7\%$  and  $18.5 \pm 2.6\%$  were also found for the two Murchison  
6 meteorite samples USNM 5453 and USNM 6650, respectively (Table 5). However, only  
7 slight L-excesses were observed in the CM2 LEW 90500 ( $3.3 \pm 1.8\%$ ), and no detectable L-  
8 isovaline excesses (within analytical uncertainty based on the standard deviation of the  
9 mean) were found in LON 94102 ( $2.4 \pm 4.1\%$ ), EET 92042 ( $-1.0 \pm 4.3\%$ ), and QUE 99177  
10 ( $0.3 \pm 2.1\%$ ) (Glavin and Dworkin, 2009). It should be emphasized that these results do not  
11 rule out circularly polarized radiation as a possible source of a small initial L-isovaline  
12 asymmetry in these weakly altered meteorites that is within our analytical error of a few  
13 percent. The errors in the  $L_{ee}$  values were calculated by standard error propagation of the  
14 absolute uncertainties of the total D- and L-isovaline concentrations reported in Table 3. We  
15 are currently unable to resolve small excesses ( $< 2\text{-}3\%$ ) on complex or dilute samples using  
16 our UPLC-FD/ToF-MS technique with sufficient confidence.  
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34 We have previously shown that the  $\beta$ -alanine/glycine molar ratio (which is approximately  
35 proportional to the  $\beta$ -alanine/ $\alpha$ -AIB ratio presented in (Glavin and Dworkin, 2009) found in  
36 carbonaceous meteorites can be used as an indicator for the extent of parent body aqueous  
37 alteration and that the magnitude of the L-isovaline excesses in CI, CM, and CR meteorites  
38 was correlated with the relative abundance of  $\beta$ -alanine (Glavin and Dworkin, 2009; Glavin  
39 et al., 2006). The  $\beta$ -ala/glycine ratios for the meteorites studied are plotted with the L-  
40 isovaline excesses in Table 5. In general, the more altered meteorites have larger L-  
41 isovaline excesses and higher relative abundances of  $\beta$ -alanine, while the less altered  
42 meteorites have smaller L-isovaline excesses and lower  $\beta$ -alanine abundances. The  
43 exception is the Murchison meteorite which has the highest L-isovaline excess, but is less  
44 altered than the type 1 chondrites. The relative abundance ratios in these meteorites should  
45 be interpreted with caution since they were not corrected for the potential contribution of  
46 terrestrial glycine contamination, which could vary between meteorite samples.  
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The delivery of amino acids by carbonaceous chondrites to the early Earth could have been an important source of the Earth's prebiotic organic inventory (Chyba and Sagan, 1992). Although  $\alpha$ -dialkyl amino acids such as isovaline are not common in the terrestrial biosphere, these amino acids are highly resistant to racemization and may have been well suited as catalysts for transferring their asymmetry to the  $\alpha$ -hydrogen protein amino acids or other prebiotic compounds, such as sugars (Pizzarello and Weber, 2004) common to all terrestrial life. There may have also been primordial chiral asymmetry in the  $\alpha$ -hydrogen amino acids, but all evidence of this would have been long erased by racemization. Based on the results of this study, it seems increasingly likely that most aqueously altered carbonaceous chondrites possess L-isovaline asymmetry. The fact that only L-amino acid excesses (no D-excesses) have been found in carbonaceous meteorites analyzed so far may indicate that the origin of life on Earth and possibly elsewhere in our solar system was biased toward L-amino acid homochirality from the very beginning.

**CONCLUSIONS**

The purpose of this study was to compare the distribution and enantiomeric composition of amino acids found in CI, CM, and CR carbonaceous chondrites that cover a wide range of aqueous alteration states. The type 1 chondrites MET 01070, SCO 06043, and GRO 95577 had not previously been analyzed for amino acids using the highly sensitive and selective liquid chromatography time of flight mass spectrometry technique. For the first time, amino acids were extracted from a set of carbonaceous chondrites and analyzed by UPLC-FD/ToF-MS using the same analytical and quantitation methods in a single laboratory. A comparison of the amino acid abundances and enantiomeric ratios of isovaline in the nine carbonaceous chondrites studied led to the following five observations.

1. The total amino acid abundances found in the type 1 chondrites MET 01070, SCO 06043, and GRO 95577 were much lower than in the corresponding type 2 meteorites from the same carbonaceous chondrite group.

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4 2. The relative abundance of  $\beta$ -alanine in the type 1 chondrites is much higher than in  
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6 less altered type 2 carbonaceous chondrites. The  $\beta$ -alanine/glycine ratio appears to  
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8 be a good chemical indicator to help assess the relative degree of aqueous alteration  
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10 in CI, CM, and CR chondrites.
- 11  
12 3. The relative abundances of the  $C_5$  amino acids show that the less altered type 2  
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14 chondrites are dominated by  $\alpha$ -amino acids with branched-chain isomers favored  
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16 over straight-chain isomers; this is consistent with formation by Strecker synthesis.  
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18 In contrast, the aqueously altered type 1 chondrites are enriched in  $\gamma$ - and  $\delta$ -amino  
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20 acids with no apparent enrichment of branched versus straight-chain isomers.
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22 4. In general, the distribution of amino acids in LON 94102 do not match well with the  
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24 CM2 chondrites Murchison and LEW 90500, possibly indicating that this CM2  
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26 meteorite originated on a chemically distinct parent body.
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28 5. Large enrichments in L-isovaline of up to 11 to 19%, were observed in the aqueously  
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30 altered type 1 chondrites and in Murchison, exceeding those found in other less  
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32 altered type 2 and type 3 chondrites from the respective chondrite groups.  
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39 The differences in the amino acid abundances, distributions, and L-isovaline asymmetry  
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41 found in CI, CM, and CR type carbonaceous chondrites are best explained by differences in  
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43 the degree of aqueous alteration on the respective meteorite parent bodies. In contrast to  
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45 previous studies of CM1 and CM2 chondrites, we observe distinct differences in relative  
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47 amino acid abundances between type 1 and 2 CM chondrites that we believe are best  
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49 explained by aqueous alteration, and are likely not the result of chemical compositional  
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51 differences between distinct CM parent bodies. One notable exception is the weakly altered  
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53 CM2 chondrite LON 94102, which has a  $C_5$  amino acid distribution that does not fit well with  
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55 other type 2 chondrites, but more closely resembles the type 1 chondrites. For LON 94102,  
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57 we cannot rule out the possibility that this meteorite did originate from a parent body that  
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59 was chemically distinct from that of the other CM2s. The finding of large L-isovaline  
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3 excesses in the most altered type 1 chondrites, but not in the most pristine unaltered type 2  
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5 and type 3 CRs provides additional support that amplification of a small L-isovaline excess  
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7 occurred during an aqueous alteration phase on the parent bodies of these carbonaceous  
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9 chondrites. We recognize that aqueous processing on the parent body cannot explain all of  
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11 the compositional differences and patterns we observe in the meteorites. Finally, to gain a  
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13 more thorough understanding for the mechanisms responsible for the differences seen in  
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15 amino acid composition between CI, CM, and CR chondrites, laboratory investigations of the  
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17 synthesis and decomposition of amino acids under realistic parent body conditions should  
18  
19 be carried out.  
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### FIGURE CAPTIONS

**Figure 1.** The 0- to 40-min. region of the LC-FD chromatograms. OPA/NAC derivatization (15 min) of the amino acid standard and of the 6 M HCl-hydrolyzed, hot-water extracts of the serpentine blank and the type 1 chondrites MET 01070, SCO 06043, and GRO 95577 are shown. Similar chromatograms were obtained for the non-hydrolyzed extracts. Peaks in the chromatograms that did not correspond to the same UV fluorescence and mass retention times of the standard amino acids tested were not identified. The identities of the peaks are given in Table 1.

**Figure 2.** The 15- to 40-min. region of the LC-ToF-MS single ion chromatograms ( $C_2$ :  $m/z = 337.09$ ;  $C_3$ :  $m/z = 351.10$ ;  $C_4$ :  $m/z = 365.12$ ;  $C_5$ :  $m/z = 379.13$ ;  $C_6$ :  $m/z = 393.15$ ;  $C_7$ :  $m/z = 407.16$ ;  $C_8$ :  $m/z = 421.18$ ;  $C_9$ :  $m/z = 435.20$ ; and  $C_{10}$ :  $m/z = 449.21$ ) in positive electrospray ionization mode. OPA/NAC derivatization (15 min) of amino acids in the standard mix and of the 6M HCl-hydrolyzed, hot-water extracts of Orgueil, MET 01070, GRO 95577, Murchison (USNM 5453), EET 92042 and the serpentine blank are shown. Similar single ion chromatograms were obtained for the non-hydrolyzed extracts. Peaks were identified by comparison of the retention time and molecular mass to those in amino acid standards run on the same day. Peak identifications are given in Table 1.

**Figure 3.** The 20- to 41-min. region of the LC-ToF-MS single ion chromatograms of the  $C_5$  amino acids ( $m/z = 379.13 \pm 0.02$ ) in positive electrospray ionization mode. OPA/NAC derivatization (15 min) of amino acids in the standard mix and of the 6M HCl-hydrolyzed, hot-water extracts of the serpentine blank and the MET 01070, SCO 06043, and GRO 95577 meteorites are shown. Similar LC-ToF-MS single ion chromatograms were obtained for the non-hydrolyzed extracts. The peaks were identified by comparison of the retention time and exact molecular mass to those in the  $C_5$  amino acid standard run on the same day. Peak identifications are given in Table 1.

**Figure 4.** A comparison of the relative molar abundances (glycine = 1.0) of alanine,  $\beta$ -alanine,  $\alpha$ -aminoisobutyric acid, and isovaline in the 6M HCl-hydrolyzed, hot-water extracts of the carbonaceous meteorites investigated in this study. The relative abundances were

calculated from the data in Table 2 after correcting for the molecular weights of each amino acid. The uncertainties were calculated by standard error propagation of the absolute errors in Table 2. The amino acid data for Murchison (USNM 6650) and LEW 90500 were taken from Glavin *et al.* (2006).

**Figure 5.** A comparison of the relative molar abundances of the C<sub>5</sub> amino acids in CI, CM, and CR carbonaceous chondrites as a function of (a) amine position ( $\alpha$ -,  $\beta$ -,  $\gamma$ -, and  $\delta$ -) and (b) valeric acid carbon chain backbone (*n*-, *sec*-, *iso*-, and *tert*-) normalized to the total number of possible structural isomers. The dashed line corresponds to the expected relative abundance if the amino acids were formed by a completely random synthetic process. The relative abundance data for MET 01070, SCO 06043, and GRO 95577 is from this study. The data for Orgueil, LON 94102, Murchison (USNM 6650), LEW 90500, EET 92042, and QUE 99177 taken from Glavin and Dworkin (2009) are shown for comparison. It is apparent from the data that in most cases there is structural similarity in C<sub>5</sub> amino acid relative abundances within a single meteorite group. However, based on our amino acid relative abundance data, LON 94102 appears to be distinct from other CM2 chondrites.

Figure 1

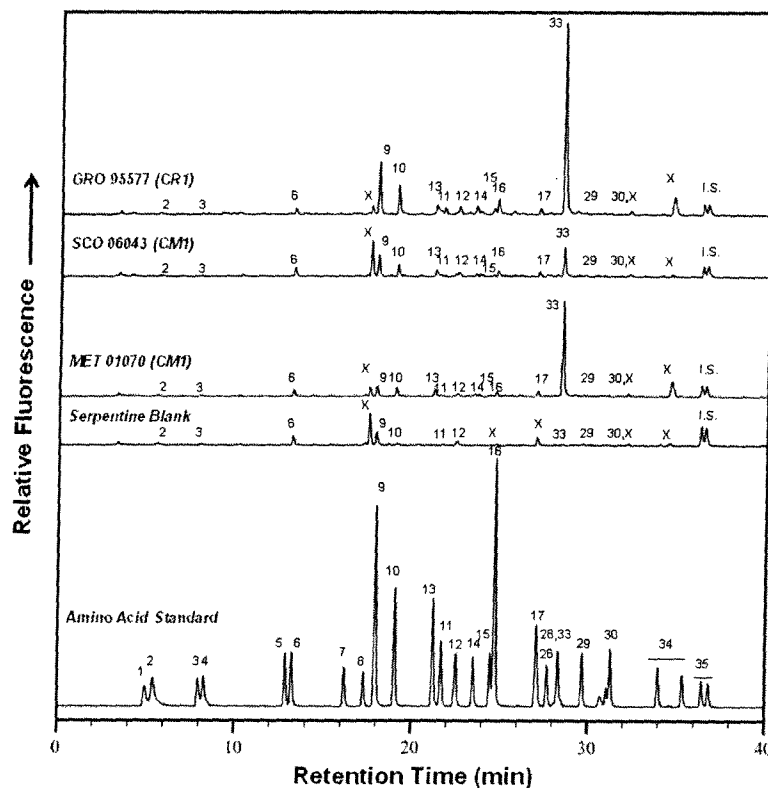
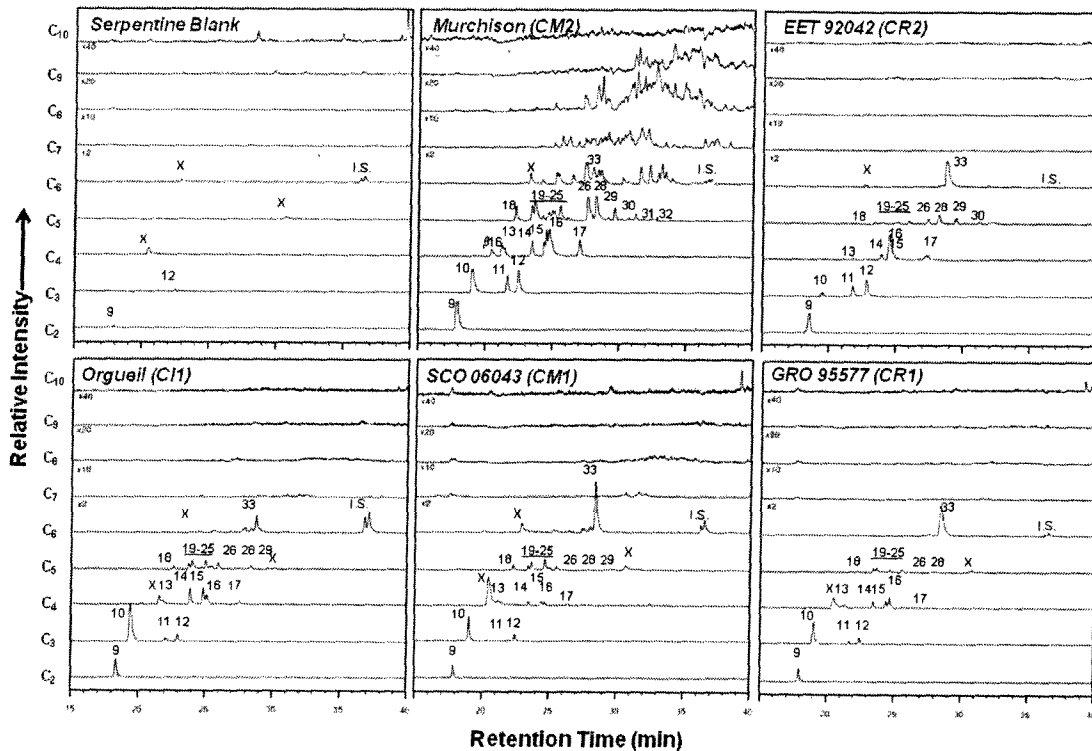
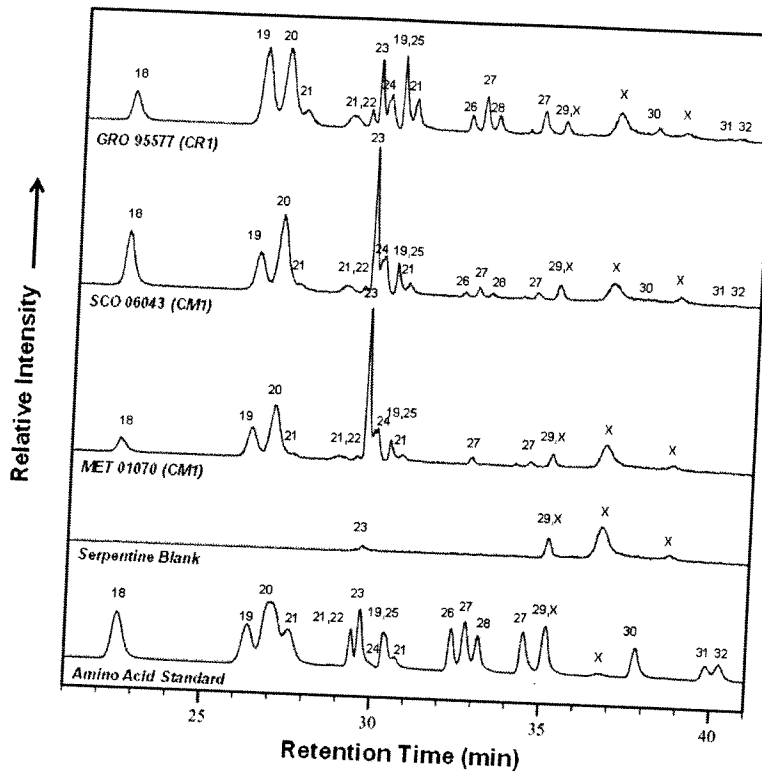


Figure 2



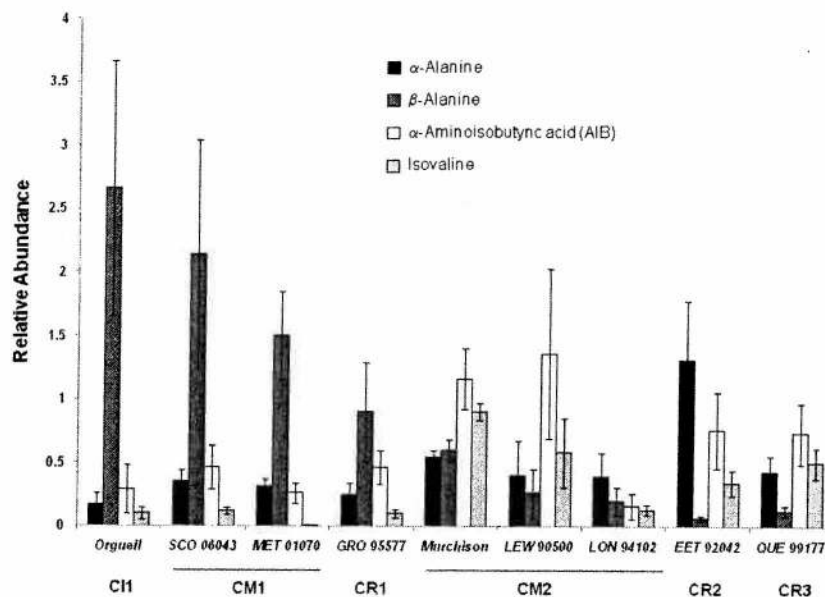
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Figure 3



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Figure 4



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Figure 5a

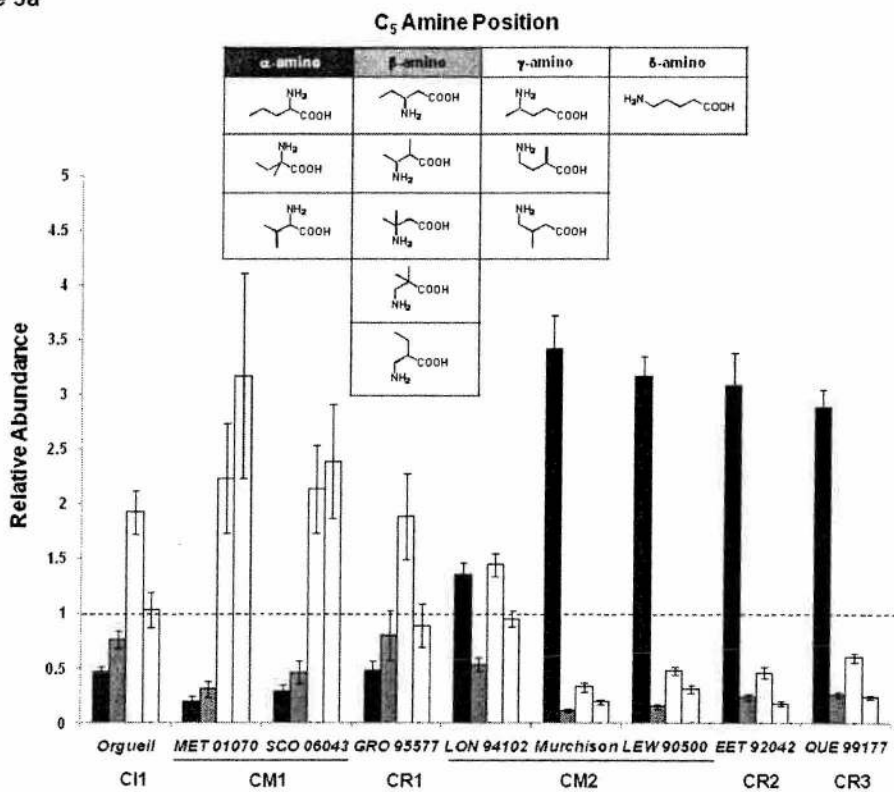
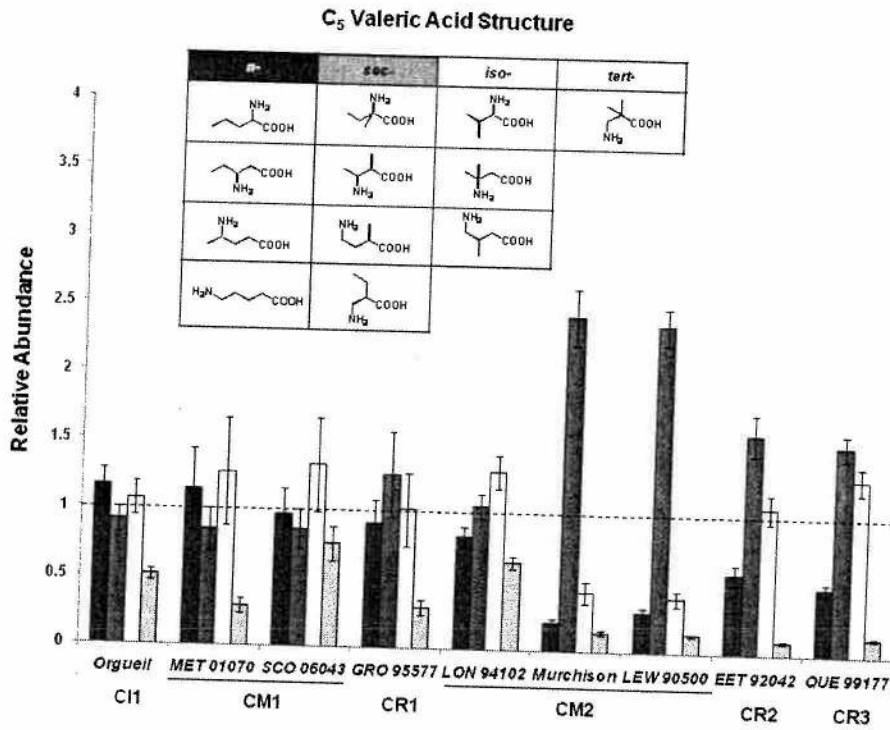


Figure 5b



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**Table 1.** Peak identification numbers and abbreviations for amino acids detected in the chromatograms of the standards, procedural blanks, and meteorite extracts. The number of carbons (#C) for the aliphatic amino acids is also shown.

Peak	Amino Acid	#C
1	D-aspartic acid	
2	L-aspartic acid	
3	L-glutamic acid	
4	D-glutamic acid	
5	D-serine	
6	L-serine	
7	D-threonine	
8	L-threonine	
9	glycine	
10	$\beta$ -alanine (BALA)	2
11	D-alanine	
12	L-alanine	
13	$\gamma$ -amino- <i>n</i> -butyric acid ( $\gamma$ -ABA)	3
14	D- $\beta$ -amino- <i>n</i> -butyric acid (D- $\beta$ -ABA)	
15	L- $\beta$ -amino- <i>n</i> -butyric acid (L- $\beta$ -ABA)	
16	$\alpha$ -aminoisobutyric acid ( $\alpha$ -AIB)	4
17	D,L- $\alpha$ -amino- <i>n</i> -butyric acid (D,L- $\alpha$ -ABA)	
18	3-amino-2,2-dimethylpropanoic acid (3-a-2,2-dmpa)	
19	D,L-4-aminopentanoic acid (D,L-4-apa)	
20	D,L-4-amino-3-methylbutanoic acid (D,L-4-a-3-mba)	
21	D,L-3-amino-2-methylbutanoic acid (D,L-3-a-2-mba)	
22	D,L-3-amino-2-ethylpropanoic acid (D,L-3-a-2-epa)	
23	5-aminopentanoic acid (5-apa)	
24	D,L-4-amino-2-methylbutanoic acid (D,L-4-a-2-mba)	
25	3-amino-3-methylbutanoic acid (3-a-3-mba)	
26	D-2-amino-2-methylbutanoic acid (D-isovaline)	5
27	D,L-3-aminopentanoic acid (D,L-3-apa)	
28	L-2-amino-2-methylbutanoic acid (L-isovaline)	
29	L-2-amino-3-methylbutanoic acid (L-valine)	
30	D-2-amino-3-methylbutanoic acid (D-valine)	
31	D-2-aminopentanoic acid (D-norvaline)	
32	L-2-aminopentanoic acid (L-norvaline)	
33	$\epsilon$ -amino- <i>n</i> -caproic acid (EACA)	
34	D,L-isoleucine	
35	D,L-leucine	
I.S.	D,L-norleucine (internal standard)	6
X	Desalting or non-fluorescent mass artifact	



**Table 2.** Summary of the total (free + bound) abundances of the two- to six-carbon amino acids identified in the acid-hydrolyzed, hot-water extracts of CI, CM, and CR carbonaceous chondrites<sup>a</sup>.

Amino Acids	CI1		CM1		CM2					CR1	CR2	CR3
	Orgueil	MET 01070	SCO 06043	Murchison USNM 5453	Murchison USNM 6650	LEW 90500	LON 94102	GRO 95577	EET 92042	QUE 99177		
	This study		This study									
D-aspartic acid	54 ± 31	< 1	< 1	189 ± 14	120 ± 16	127 ± 24	118 ± 63	< 8	1,028 ± 157			
L-aspartic acid	55 ± 28	< 2	< 2	281 ± 23	132 ± 15	151 ± 73	112 ± 41	< 2	1,016 ± 145		546 ± 214	
D-glutamic acid	47 ± 16	< 2	< 1	503 ± 82	343 ± 44	317 ± 55	534 ± 193	< 9	10,088 ± 1,973		536 ± 170	
L-glutamic acid	83 ± 22	< 2	< 3	819 ± 113	357 ± 42	316 ± 55	552 ± 191	< 1	10,180 ± 1,743		3,517 ± 602	
D-serine	< 1	< 1	< 3	n.d.	48 ± 9	219 ± 151	99 ± 69	< 4	2,584 ± 1,420		3,442 ± 537	
L-serine	< 1	< 2	< 2	n.d.	46 ± 9	235 ± 163	92 ± 30	< 3	2,266 ± 1,237		685 ± 426	
Glycine	865 ± 450	105 ± 15	48 ± 10	2,606 ± 717	1,995 ± 122	1,448 ± 682	3,391 ± 1,029	278 ± 76	54,527 ± 15,365		609 ± 379	
D-alanine	80 ± 17	18 ± 2	10 ± 1	592 ± 99	623 ± 6	343 ± 171	823 ± 295	46 ± 16	40,107 ± 9,275		14,134 ± 3,376	
L-alanine	94 ± 22	21 ± 3	10 ± 2	718 ± 160	659 ± 84	352 ± 161	745 ± 286	35 ± 5	41,326 ± 7,477		3,589 ± 455	
β-alanine	2,732 ± 675	187 ± 34	122 ± 45	1,081 ± 243	1,419 ± 157	442 ± 238	821 ± 343	295 ± 98	4,161 ± 1,182		3,544 ± 594	
D,L-α-ABA*	71 ± 49	23 ± 4	4 ± 1	622 ± 128	403 ± 156	431 ± 159	896 ± 300	45 ± 10	20,859 ± 5,653		1,954 ± 332	
D-β-ABA	221 ± 110	26 ± 4	18 ± 3	349 ± 58	233 ± 17	155 ± 16	279 ± 110	70 ± 10	2,855 ± 580		3,300 ± 637	
L-β-ABA	181 ± 59	25 ± 2	18 ± 2	320 ± 41	256 ± 15	172 ± 40	178 ± 81	60 ± 9	3,160 ± 852		585 ± 127	
γ-ABA	281 ± 131	46 ± 13	54 ± 12	979 ± 97	1,460 ± 213	164 ± 21	470 ± 115	140 ± 12	2,620 ± 982		679 ± 203	
α-AIB	343 ± 140	37 ± 10	30 ± 9	5,124 ± 2,487	3,182 ± 620	2,706 ± 377	753 ± 413	175 ± 15	56,825 ± 15,592		1,322 ± 115	
EACA	108 ± 103	2,014 ± 86	954 ± 147	490 ± 148	268 ± 123	386 ± 169	9,265 ± 370	3,914 ± 573	51,371 ± 13,566		14,246 ± 3,249	
C <sub>5</sub> Amino Acids (from Table 3)	1,600	220	350	7,300	3,100	1,800	5,900	500	65,000		28,000	
<b>Free Amino Acids<sup>†</sup> (% Free)<sup>‡</sup></b>	<b>n.d.</b>	<b>340 (48%)</b>	<b>250 (38%)</b>	<b>11,000 (52%)</b>	<b>7,100 (51%)</b>	<b>4,200 (45%)</b>	<b>8,400 (53%)</b>	<b>830 (52%)</b>	<b>205,000 (64%)</b>		<b>40,000 (49%)</b>	
<b>Total (free + bound) Amino Acids<sup>†</sup></b>	<b>6,700</b>	<b>710</b>	<b>660</b>	<b>21,000</b>	<b>14,000</b>	<b>9,400</b>	<b>16,000</b>	<b>1,600</b>	<b>320,000</b>		<b>81,000</b>	

<sup>a</sup>All values are reported in parts-per-billion (ppb) on a bulk sample basis. Extracts were analyzed by OPANAC derivatization (1 and 15 min.) and HPLC separation with UV fluorescence and time of flight mass spectrometry (ToF-MS) detection. For the LC-ToF-MS data, the mono-isotopic masses of each protonated OPANAC amino acid derivative (M + H<sup>+</sup>) was used for quantification and final peak integrations included background level correction using a serpentine blank and a comparison of the peak areas with those of an amino acid standard run on the same day. The uncertainties (δx) are based on the standard deviation of the average value of 4 to 8 separate measurements (n) with a standard error, δx = σ<sub>x</sub> · (n-1)<sup>-1/2</sup>.  
<sup>†</sup>Enantiomers could not be separated under the chromatographic conditions.  
<sup>‡</sup>Amino acid concentrations rounded to two significant digits. The EACA concentrations were not included in the sum of the individual amino acid concentrations.  
<sup>§</sup>Percent free amino acids in the non-hydrolyzed extracts were calculated by dividing the free amino acid concentration by the total (free + bound) amino acid concentration measured in the acid-hydrolyzed extracts.  
n.d. = not determined.

**Table 3.** Summary of the total (free + bound) abundances of the five-carbon (C<sub>5</sub>) α-, β-, γ-, and δ-amino acids in the acid-hydrolyzed, hot-water extracts of CI, CM, and CR type carbonaceous chondrites<sup>a</sup>.

C <sub>5</sub> Amino Acids	CI1		CM1		CM2				CR1	CR2	CR3
	Orgueil	MET 01070	SCO 06043	Murchison USNM5453	Murchison USNM6650	LEW 90500	LOH 94102	GRO 95577	EET 92042	QUE 99177	
	Glavin and Dworkin (2009)										
	This study										
	13 ± 1	< 0.2	0.6 ± 0.2	59 ± 4	18 ± 2	9 ± 1	259 ± 20	Glavin and Dworkin (2009)			
D-norvaline	14 ± 1	< 0.2	0.8 ± 0.2	59 ± 3	19 ± 2	9 ± 1	263 ± 17	2.2 ± 0.1	3,275 ± 391	960 ± 60	
L-norvaline	36 ± 3	< 0.4	3.3 ± 0.4	1263 ± 125	993 ± 43	652 ± 15	332 ± 19	1.8 ± 0.2	3,231 ± 351	967 ± 68	
D-isovaline	49 ± 2	< 0.5	4.6 ± 0.5	2058 ± 220	1,444 ± 66	696 ± 20	348 ± 21	14.6 ± 1.7	14,446 ± 625	5,534 ± 210	
L-isovaline	22 ± 6	< 0.3	2.6 ± 1.1	160 ± 5	105 ± 15	48 ± 6	357 ± 20	18.2 ± 2.0	14,164 ± 1,038	5,564 ± 92	
D-valine	56 ± 2	< 10 <sup>**</sup>	< 14 <sup>**</sup>	< 330 <sup>**</sup>	111 ± 28	55 ± 8	480 ± 21	5.5 ± 0.1	7,217 ± 427	3,442 ± 208	
D,L-3-apa <sup>†</sup>	181 ± 11	5.9 ± 0.6	10 ± 1	158 ± 19	37 ± 4	36 ± 2	178 ± 9	< 12	7,647 ± 500	3,861 ± 137	
D,L- and allo-3-a-2-mba <sup>†</sup>	64 ± 4	5.2 ± 0.3	9.5 ± 0.6	158 ± 20	26 ± 5	24 ± 4	162 ± 7	41 ± 4	2,077 ± 177	804 ± 25	
3-a-3-mba <sup>††</sup>	< 30	< 3	< 3	< 66	< 14	< 14	< 150	29 ± 2	2,161 ± 102	962 ± 29	
3-a-2,2-dmpa	69 ± 3	5.7 ± 0.5	23 ± 2	195 ± 16	36 ± 3	21 ± 1	320 ± 13	< 4	< 286	< 250	
D,L-3-a-2-epa <sup>*</sup>	171 ± 14	11 ± 1	25 ± 6	470 ± 233	37 ± 2	24 ± 3	534 ± 34	13 ± 1	539 ± 21	315 ± 20	
D,L-4-apa <sup>†</sup>	283 ± 24	21 ± 2	33 ± 2	444 ± 47	84 ± 12	78 ± 9	475 ± 31	88 ± 31	1,393 ± 127	774 ± 59	
D,L-4-a-2-mba <sup>*</sup>	173 ± 11	50 ± 2	61 ± 1	553 ± 58	67 ± 8	75 ± 3	733 ± 38	75 ± 4	2,968 ± 212	1,293 ± 90	
D,L-4-a-3-mba <sup>*</sup>	325 ± 17	62 ± 15	102 ± 21	984 ± 180	103 ± 7	68 ± 2	973 ± 23	62 ± 5	2,396 ± 311	1,485 ± 60	
5-apa	140 ± 19	63 ± 12	73 ± 12	341 ± 59	50 ± 5	48 ± 5	480 ± 24	111 ± 23	2,137 ± 75	1,472 ± 87	
<b>Total C<sub>5</sub> Amino Acids<sup>††</sup></b>	<b>1,600</b>	<b>220</b>	<b>350</b>	<b>7,300</b>	<b>3,100</b>	<b>1,800</b>	<b>5,900</b>	<b>500</b>	<b>65,000</b>	<b>28,000</b>	

<sup>a</sup>All values are reported in parts-per-billion (ppb) on a bulk sample basis. Extracts were analyzed by OPA/NAC derivatization (15 min.) and LC-FD/ToF-MS. For the ToF-MS data, the mono-isotopic masses (m/z 379, 13) of each protonated OPANAC amino acid derivative (M + H<sup>+</sup>) was used for quantification and final peak integrations included background level correction using a procedural blank and a comparison of the peak areas with those of an amino acid standard run on the same day. The uncertainties (δx) are based on the standard deviation of the average value of 4 to 8 separate measurements (n) with a standard error, δx = σ<sub>x</sub> · (n-1)<sup>-1/2</sup>. For isovaline, the reported values represent the average of 8 to 23 separate measurements.

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

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

<sup>††</sup>3-a-3-mba co-elutes with one of the enantiomers of D,L-4-apa, therefore upper limits for 3-a-3-mba were estimated by taking the difference in peak areas of the two D,L-4-apa enantiomers.

<sup>‡</sup>Peak detected above blank levels, however only upper limit reported due to unidentified co-eluting peak X (see Figs. 1 and 2).

<sup>†††</sup>Total (free + bound) amino acid abundances rounded to two significant digits.