



Mass Spectrometry of Astrobiologically Relevant Organic Material

Implications on Future Space Missions to Ocean Worlds in the Outer Solar System

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The Solar System's ocean worlds



Steve Vance; NASA/JPL-Caltech

The Solar System's ocean worlds



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Sampling dust and ice particles in space



The Cosmic Dust Analyzer (CDA)



- Sensitive to cations
- Determines impact rates, mass, speed, electric charge, and composition
- Mass resolution: \leq 50 m/ Δ m
- Maximum recorded mass: ca. 200 u



Srama et al. 2004

Analog experiment: IR-FL-MALDI-ToF-MS



- Liquid beam: r = 7.5 μm
- Vaccum chamber: P = 5 x 10⁻⁵ mbar

- Laser: λ = 2480 nm; I ≤ 1152 MW/cm²
- ToF-MS: R = 700 800 m/∆m

Comparison of the impact mechanisms of CDA and the analog experiment





Comparison of the impact mechanisms of CDA and the analog experiment



Future missions and Enceladus Ice Analyzer (ENIA)



Sodium complexation of amino acids



Sodium complexation of amino acids



Detection of two species at the same integer mass



Detection limits of amino acids

Salt poor ma	trix: H ₂ O			Salt ric	h matr	ix: ≥ 0.1 M Na	NaCl		
AMINO ACID	MOLECULAR WEIGHT [u]	DETECTED PEAK	DETECTION LIMIT [PPB MOLAR]	AMINO	ACID	MOLECULAR WEIGHT [u]	DETECTED PEAK	DETECTION LIMIT [PPB MOLAR]	
		M* + 1 u	120	Glyc	ine	75	M + 45 u	4800	
Glycine	75	M – 1 u	25	Serine		105	M + 45 u	340	
		M + 1 u	170	Threo	nine	119	M + 45 u	1500	
Serine	105	M – 1 u	340	Ornithine		132	M + 45 u	1400	
Lysine	146	M + 1 u	0.02	Aspartic acid		133	M + 67 u	70	
Arginine	174	M + 1 u	0.02	Glutamic acid		147	M + 67 u	60	
	132	M + 1 u	0.3	Histidine		155	M + 45 u	230	
Ornithine		M – 1 u	270	Arginine		174	M + 45 u	1000	
		M + 1 u	0.2	Citrul	line	175	M + 45 u	2100	
Histidine	155	M – 1 u	230	Tyros	ine	181	M + 67 u	200	
	175	M + 1 u	20		Amino acids detectable at 5000 x lower				
Citrulline		M – 1 u	100						
	181	M + 1 u	200		in a salt rich matrix				
Tyrosine		M – 1 u	200	L					

Cations detectable at 1000 x lower concentrations than anions

*M: Molecular weight

Amino acid abundances: abiotic/biotic



Amino acids in the Enceladean ocean assuming a chondritic rocky core* and water-rock interaction

* Calculations based on Pizzarello et al. (2012) and Glavin et al. (2010)

ACID	CONCENTRATION [PPM MOLAR]
Glycine	36.0
Alanine	30.3
Valine	11.5
Aminobutyric acid	7.0
Glutamic acid	0.6
Serine	0.5
Aspartic acid	0.3
Valeric acid	10.6
Methylbutyric acid	10.6
Hexanoic acid	4.6
Acetic acid	5.6
Butyric acid	4.1
Heptanoic acid	4.1
Octanoic acid	3.8
Nonanoic acid	3.4
Methylpentanoic acid	3.1
Ethylhexanoic acid	2.5
Benzoic acid	1.5
Formic acid	0.8
Ethylbutyric acid	0.8
Decanoic acid	0.5

Abiotic case

Amino acids in the Enceladean ocean assuming a chondritic rocky core* and water-rock interaction

* Calculations based on Pizzarello et al. (2012) and Glavin et al. (2010)



Amino acids: Biotic case

	ACID	CONCENTRATION [PPM MOLAR * 10]
	Serine	5.1
	Glycine	4.8
1_	Citrulline	2.1
LC. LQ	Tyrosine	2.0
	Aspartic acid	1.4
•	Ornithine	1.4
	Lysine	0.6
2	Arginine	0.5
	Valeric acid	167.6
	Hexanoic acid	93.1
	Octanoic acid	62.5
5 Ar allor	Acetic acid	45.0
	Butyric acid	40.9
Cit noic 10 Cit Association 11 Cit	Nonanoic acid	28.5
	Heptanoic acid	27.7
	Benzoic acid	14.8
	Formic acid	7.8
	Propionic acid	7.7
allan and a state of the state	Decanoic acid	5.2
40 50 60 70 80 90 100 110 120 130 140 150 160 170 180 1	90	
Mass [u]		

Fatty acid abundances: abiotic/biotic



Fatty acids: abiotic case



Fatty acids: biotic case



Fatty acids: biotic case



Summary and outlook

- Analog desorption mass spectra highly comparable to impact ionization mass spectra taken in space
- Most amino acids better detectable as cations, a few amino acids and fatty acids as anions
- Amino acids form sodium complexes M + 45 u and/or M + 67 u in a salt rich matrix (≥ 0.1 M); Fatty acids form complexed dimers
- Peptides have been investigated → Characteristic cleavages lead to characteristic fragmentation patterns diagnostic for its amino acid constituents
- Detection limits of amino acids and fatty acids depend on salt concentration but are in general below 1 ppm; Glu and Asp are the most sensitive amino acids in a salt rich matrix, Arg and Lys in a salt poor matrix
- Peak heights can be well "translated" into concentrations of the organics → abiotic and biotic signatures can be easily distinguished
- By comparing the laboratory results with spacecraft data, we have the ability to recognize and distinguish such signatures on icy moons with a subsurface ocean → Europa Clipper mission + other upcoming missions
- We aim to create a comprehensive spectrum library for in situ space detectors from a wide variety of analog materials in icy grains

Backup

Backup: Which objects harbor liquid oceans?



Backup: Which objects harbor liquid oceans?



Backup: Hypervelocity impact signals (CA)



Figure 6. Hypervelocity impact signals of dust grains onto the big Impact Ionization Target (left) and onto the Chemical Analyzer Target (right). The chemical analyzer is a time-of-flight mass spectrometer and provides the elemental composition of the impacting dust grain. Srama et al. 2004

Backup: Mimicking Type-1 and Type-3 mass spectra



Backup: Some amino acid structures



Davila & McKay, 2014

Backup: Confirmation of Arg Detection



Backup: Amino acids & fatty acids: Biotic case



Backup: Peptides



Backup: Peptides



Backup: Sodium complexation of straight chained carboxylic acids



Backup: Sodium complexation of straight chained carboxylic acids



Backup: Sodium complexation of straight chained carboxylic acids and their detection limits

