A PROSPECTIVE REMOTE SENSING BIOSIGNATURE FOR PEPTIDES IN ENCELADAN PLUME DEPOSITS Dale P. Winebrenner^{1,2}, Stephen G. Warren², Nathan Oborny³, Charity M. Phillips-Lander⁴, and David J. Anthony⁴ ¹Applied Physics Laboratory, ²Dept. of Earth and Space Sciences, University of Washington, ³Jet Propulsion Laboratory, ⁴Southwest Research Institute **2023** Fall Meeting of the American Geophysical Union, Abstract P43E-3319

Synopsis

- > Enceladan plume surface deposits are likely water-ice 'snow' with traces of organics [1-3]
- > Remote sensing of surface organics informs landing-site selection and context, e.g., for Enceladus Orbilander [4], before and after landing, respectively
- ➢ Nearly all elements needed for carbon-based life present on Enceladus [5] -- but terrestrial amino acids? Nucleic acids?
- Many, varied proteins so essential on Earth [e.g., 6] argue for proteins in any alien carbon-based life in water (even if composed of different amino acids)
- > Peptide bonds between amino acids define proteins –short peptides can occur abiotically [7-10] at low yields, but numerous peptide bonds indicate protein and thus life
- ➤ 'Gatekeeper' calculations support a solar-illumination, 200-350 nm (UV) imaging-spectroscopic signature of proteins (even alien ones) in porous water-ice surface deposits (i.e. 'snow') at concentrations of ppm by weight

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Plume Particles [20]





grains [22]



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Background

Water-ice refractive index [11-13]

Water-ice is very transparent (imaginary part of refractive index ~10⁻⁹) from 190-400 nm wavelength – ice particles scatter almost all and absorb almost no ight in that band

Peptide-Bond Absorption [14-19]

Peptide bonds in aqueous solution (at \sim 300K) display a strong absorption feature centered at ~187 nm and extending to ~220 nm. Moreover, the strength of this absorption for proteins without aromatic side-chains on component amino acids scales linearly with the number of peptide bonds in the molecule. Even with aromatic side chains, peptide bonds are the dominant cause of 187-220 nm absorption.

Molecular/Microbial Entrainment in

 $10^{5} - 10^{7}$ cells/ml [3] 'typical' cell 1 micron size, $7x10^{-13}$ g, 70% water by mass, half of solid mass is protein

Focus on Tiger-Stripes, 'large' ice

Brightness Ratio of Enceladan 'Snow' in Scattered Sunlight at 200 nm vs 350 nm Depends on Protein/Peptide-Bond Concentration at ppm Level



Methods

- measured in aqueous solution [14-19] as on Enceladus. The spectral width of the should be the first target of further investigation.
- distribution of peptides/proteins in spherical ice particles comprising Enceladan 'snow' (and add contributions of ice and protein to the imaginary part of the refractive index). Warren [23, 24] to compute single-particle scattering parameters, and the 2-stream model of Petty [25] to compute albedo of a semi-
- ➢ For initial work, assume homogeneous ➤ Use Mie-scattering code of Wiscombe and infinite snowpack.

> We initially take peptide bond absorption as representative of absorption at ~70-140K in ice absorption at low temperature is critical and

Interferences by Other Molecules?

- > solid NaCl: High, featureless reflectivity irradiation induces a broad, shallow absorption at~230 nm [27]. Hydrated salt data apparently unavailable.
- > solid ammonia: Very low absorption at visible and long UV wavelengths, sharp, strong absorption edge for wavelengths solid ammonia data unavailable
- > amino acid aromatic side chains: Add to peptide absorption at 190-220 nm [17].
- > adenosine in DNA and ATP: Adenosine in cells than protein [30-32].

(i.e., low absorption) of unirradiated solid salt from 500 nm to 200 nm [26], electron <190 or perhaps 200 nm [28,29]. Hydrated

also shows a broad absorption peak at ~ 190 nm, but weaker and with lower abundance