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## EDITED BY

Alexandra Gogou,  
Hellenic Centre for Marine Research  
(HCMR), Greece

## REVIEWED BY

Giuseppe Francesco Cesare Lama,  
University of Naples Federico II, Italy

## \*CORRESPONDENCE

Khan M. G. Mostofa,  
✉ mostofa@tju.edu.cn

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# The potential nature of “dark” dissolved organic matter in the biosphere

Khan M. G. Mostofa<sup>1\*</sup>, Jie Yuan<sup>2</sup>, Xuemei Yang<sup>1</sup>,  
Mohammad Mohinuzzaman<sup>1</sup>, Cong-Qiang Liu<sup>1</sup>, Nicola Senesi<sup>3</sup>,  
Giorgio S. Senesi<sup>4</sup>, Davide Vione<sup>5,6</sup> and Si-Liang Li<sup>1</sup>

<sup>1</sup>School of Earth System Science, Tianjin University, Tianjin, China, <sup>2</sup>College of Resources and Environment, Xingtai University, Xingtai, Hebei, China, <sup>3</sup>Dip.to di Scienze del Suolo, Della Pianta e Degli Alimenti, Università degli Studi di Bari “Aldo Moro”, Bari, Italy, <sup>4</sup>CNR - Istituto per la Scienza e Tecnologia dei Plasmi (ISTP) - Sede di Bari Via Amendola, Bari, Italy, <sup>5</sup>Università degli Studi di Torino, Dipartimento di Chimica, Torino, Italy, <sup>6</sup>Centro Interdipartimentale NatRisk, Grugliasco, Italy

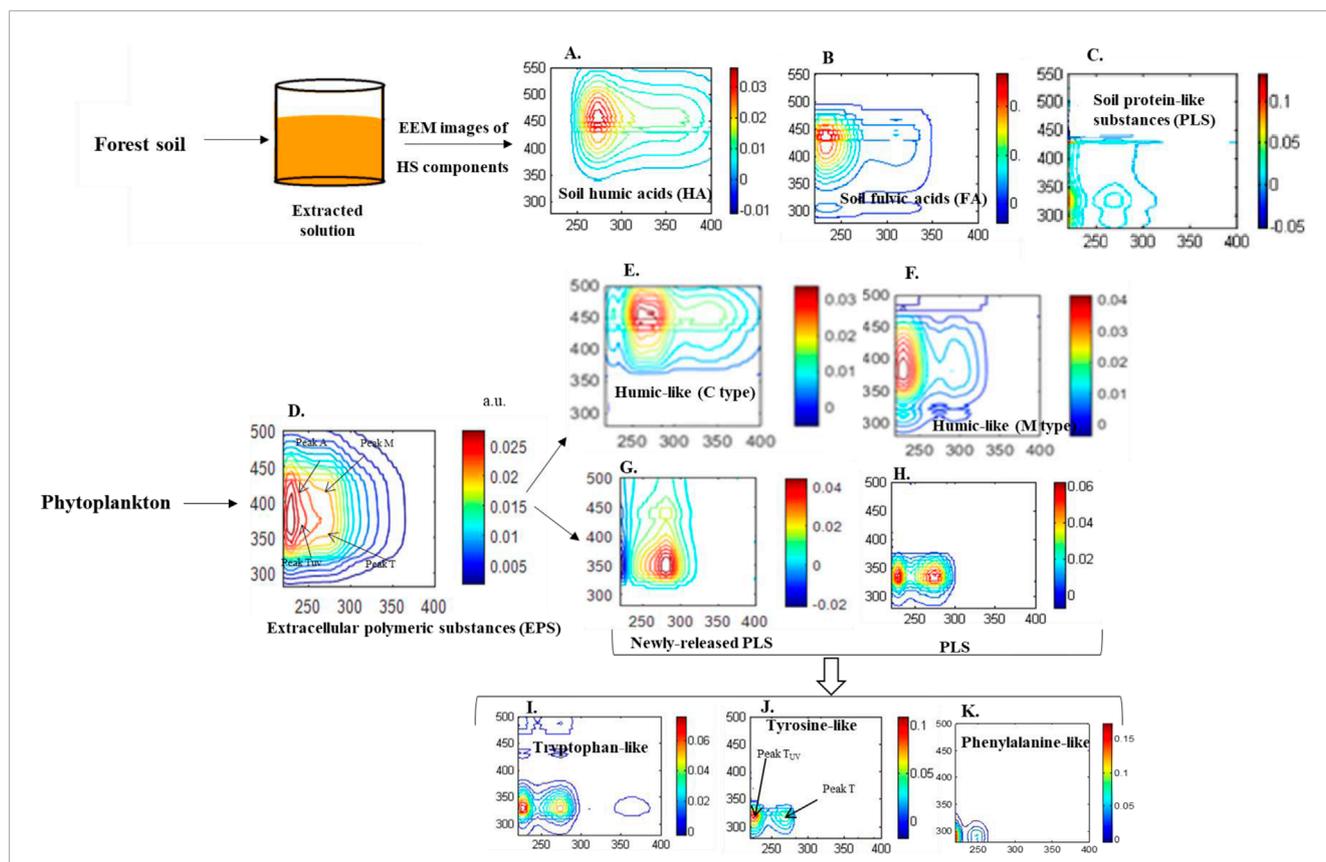
## KEYWORDS

dark matter, dark dissolved organic matter, soil humic acids, autochthonous humiclike substances, soil, surface water

## Introduction

Dark matter, also known as hidden/missing mass or nonluminous matter, is a component of the universe that spans 90 orders of magnitude in mass, ranging from ultralight bosons (often referred to as “fuzzy dark matter” (Hui et al., 2017), to massive primordial black holes (Bertone and Tait, 2018). These concepts have raised renewed interest following the detection of gravitational waves by the Laser Interferometer Gravitational-Wave Observatory (LIGO) and Virgo, which originated from the merging of black holes several tens of times more massive than the Sun (Bertone and Tait, 2018; Bird et al., 2016; Clesse and García-Bellido, 2017), which respond to gravity, and remains invisible to light (Hecht, 2016). Biogeochemical scientists have to tackle a similarly puzzling issue with dark dissolved organic matter (DDOM) in surface waters (Cai et al., 2024; Hu et al., 2023). Recently, Cai et al. (2024) using ultrahigh-resolution Fourier transform ion cyclotron resonance mass spectrometry (FT-ICR MS), identified 9,141 dark DOM molecules that exhibited high-molecular-weight (HMW) and greater diversity than the classical DOM subset, analyzing 38 DOM extracts covering the continuum of the Yangtze River-to-ocean, whereas undetected peaks and bacterial nodes were considered to represent DDOM (Cai et al., 2024; Hu et al., 2023). Notably, the HMW DDOM fraction was found to increase along this river-to-ocean continuum (Bird et al., 2016). Other studies have shown that only 8.7% and 9.6% of the 50,942 and 48,392 m/z peaks of DOM measured by FT-ICR MS in, respectively, sediments and waters of worldwide rivers, could be assigned to identifiable molecular formulae (Toyoda, 2020; Goldman, 2020). However, these undetected peaks derived from DOM remain elusive, primarily due to lack of reference spectra available in current databases (da Silva et al., 2015).

Many of the mentioned studies, however, have not taken into account the authentic sources of DOM, specifically allochthonous (terrestrial) and autochthonous (aquatic) sources, along with their optical and chemical characteristics. This oversight may lead to misconceptions regarding the authenticity of DDOM. In particular, which fractions or components of DOM should be prioritized for consideration as DDOM candidates, and what are



**FIGURE 1**  
Fluorescence (excitation-emission matrix, EEM) spectra of terrestrial humic substances [humic acids-HA (A), fulvic acids-FA (B) and protein-like substances-PLS (C)] extracted from a forest soil, and of extracellular polymeric substances (EPS, (D)) from phytoplankton and their subsequently released autochthonous humic-like substances (C-type, (E) and M-type, (F) and protein-like substances (G, H), which then generate their individual components, i.e., tryptophan-like substances (I), tyrosine-like substances (J) and phenylalanine-like substances (K) in water.

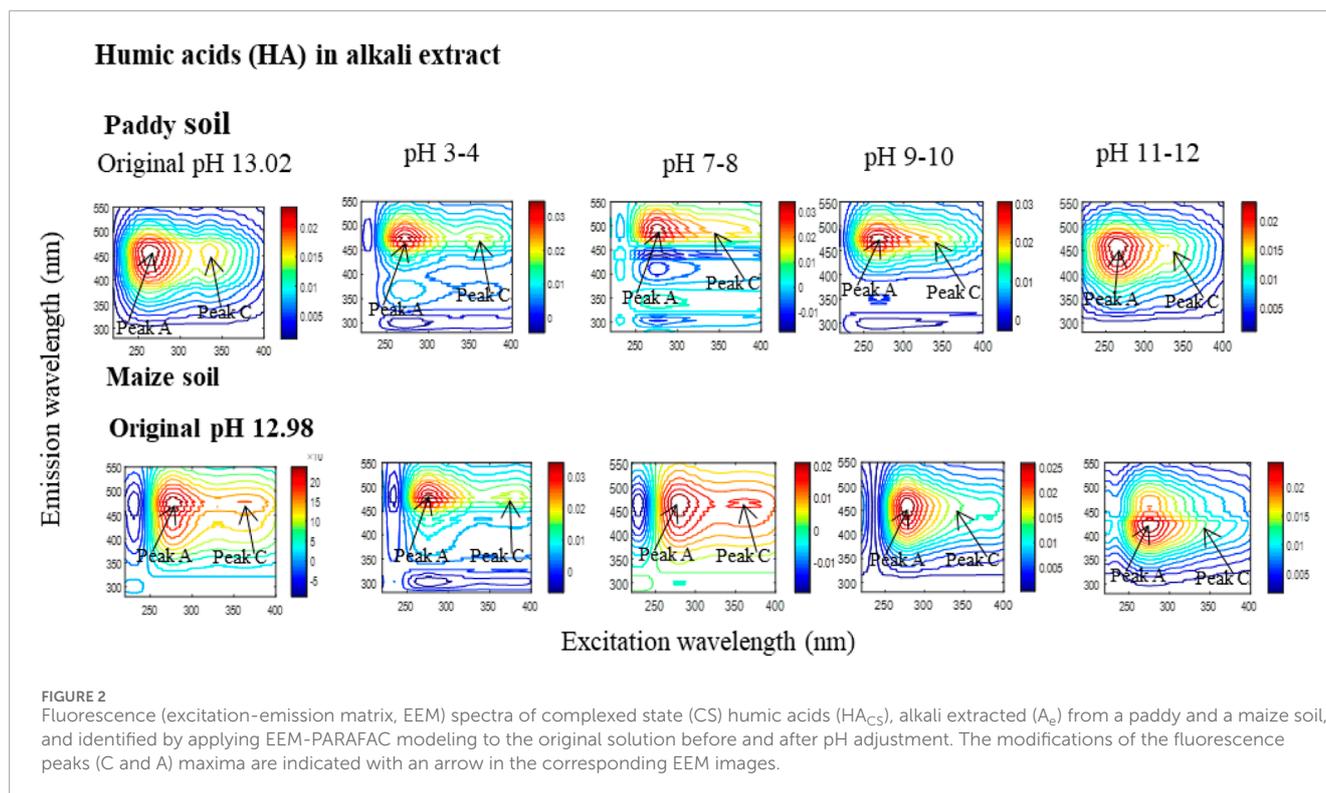
the key fundamental questions regarding DDOM in the biosphere that remain unresolved?

## Source characteristics of DOM and their relevance as dark DOM

In general, allochthonous DOM detected in natural waters is primarily derived from soil containing decaying terrestrial plant materials (Senesi and Loffredo, 1999; Piccolo, 2002), and is then partially transported to surface waters through surface runoff and groundwater leaching (Catalán et al., 2016; Zark and Dittmar, 2018; Mostofa et al., 2018; Yi et al., 2021; Mostofa et al., 2019). Allochthonous DOM is predominantly composed of humic substances (HS), which include humic acids (HA), fulvic acids (FA), and protein-like substances (PLS) (Figure 1A–C) (Gao et al., 2018; Mohinuzzaman et al., 2020; Tadini et al., 2018; Yang et al., 2024).

In contrast, autochthonous DOM in water originates from planktonic photosynthetic organisms (e.g., phytoplankton) via photo/microbial respiration processes (Guidi et al., 2016; Shammi et al., 2017; Yang et al., 2021; Flemming et al., 2016), and is primarily exported as extracellular polymeric substances (EPS) (Figure 1D) (Shammi et al., 2017; Yang et al., 2021; Flemming et al., 2016). The EPS subsequently release various components of DOM,

including autochthonous humic-like substances (C and M types), carbohydrates, lipids, and protein-like substances, which encompass a range of amino acids, including tryptophan-like, tyrosine-like, and phenylalanine-like substances (Figures 1E–G) (Shammi et al., 2017; Yang et al., 2021; Adav et al., 2008; Parlanti et al., 2000; Wei and Jin, 2022). Importantly, all soil FA and PLS fractions, as well as autochthonous DOM exhibit recognized solubility across all pH conditions, and they are highly degradable and undergo modifications when passing from inland to marine waters (Catalán et al., 2016; Zark and Dittmar, 2018; Mostofa et al., 2019; Shammi et al., 2017; Yang et al., 2021; Smith et al., 2017; Zhang et al., 2009; Mostofa et al., 2007; Moran et al., 2000). In contrast, HA possess a macromolecular/supramolecular structure that is chemically, and microbially recalcitrant. They exhibit multifunctional properties, including polyfunctionality, polyelectrolytic, size polydispersity, physical heterogeneity, and structural lability (Senesi and Loffredo, 1999; Piccolo, 2002; Tadini et al., 2018; Schulten and Schnitzer, 1993; Steelink, 2002; Sutton and Sposito, 2005). These characteristics are primarily responsible for the remarkable ability of HA to form organo-mineral complexes, which contribute to the stabilization of organic C (Hemingway et al., 2019; Moore et al., 2023; Zhang et al., 2023) and serve as essential constituents in the continuous supply of nutrients for plant and microorganism growth (Senesi



and Loffredo, 1999; Piccolo, 2002; Sutton and Sposito, 2005; Garcíá et al., 2016; de Melo et al., 2016; Wang et al., 2022; Tiwari et al., 2023).

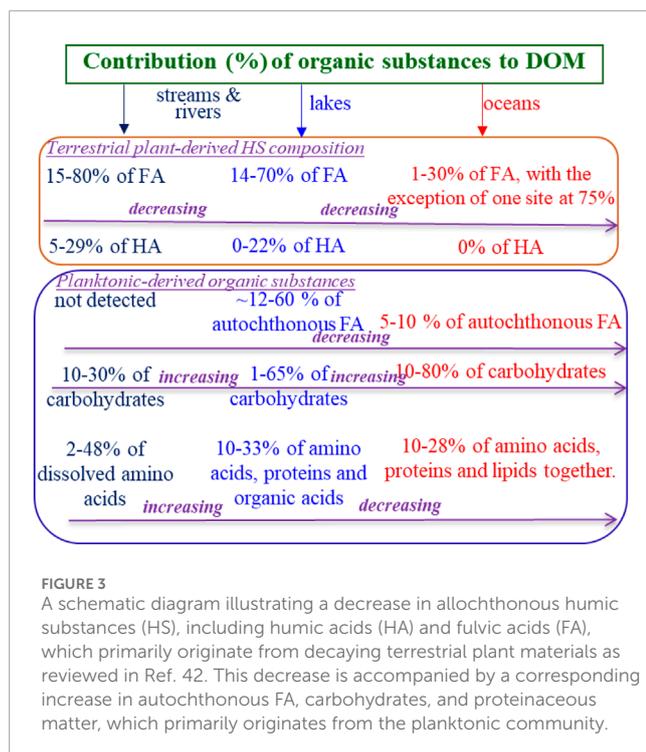
The solubility and/or insolubility characteristics of HA are significant features in their behavior (Senesi and Loffredo, 1999; Piccolo, 2002; Yang et al., 2024). In principle, a decrease in the solution pH enhances the intramolecular forces (IF) of HA by increasing the protonation of their functional groups. This, in turn, reduces their electron-donating capacity in aqueous solutions (Yang et al., 2024). Specifically, as acidity increases, the net IF become predominant, leading to enhanced intramolecular interactions among various functional groups through hydrogen bonding. This interaction can render some fractions or functional groups of HA undetectable (Yang et al., 2024). Ultimately, all functional groups associate, resulting in the precipitation of HA from the solution (Yang et al., 2024). For instance, as pH decreases, the concentration of alkali-extracted HA (dissolved in a 0.1 M NaOH solution at pH ~13.0) gradually diminishes, with a fraction precipitating at pH 6.0, while the remaining HA fractions completely precipitate at pH 1.0. The corresponding pH-dependent changes in fluorescence (excitation-emission matrix, EEM) spectra and their peaks (C and A) are illustrated in Figure 2 (Yang et al., 2024). The alkali-extracted, complexed state dissolved organic carbon ( $DOC_{CS}$ ) is estimated to decrease by approximately 39.1%–46.4% at pH 6 and by 48.1%–53.8% at pH 1. This process is accompanied by a reduction in the intensity of the HA fluorescence peak C by approximately 29.7%–47.0% at pH 6, with a complete disappearance at pH 1–2 (Figure 2) (Yang et al., 2024).

Similarly, the water extracted, labile state DOC ( $DOC_{LS}$ ) decreases by approximately 48.3%–49.2% at pH 6, and completely disappears at pH 1–2 (Figure 2) (Yang et al., 2024). These results

suggest that as pH decreases, the TOC analyzer is unable to detect certain fractions of HA as DOC because these fractions remain in an insoluble state. Consequently, a pH-dependent disappearance of HA from the solution occurs. This missing fraction of DOM/HA likely consists of high-molecular-weight (HMW) DOM, where intramolecular interactions among functional groups impede their identification (Yang et al., 2024). In essence, the pH-dependent behavior of soil HA involves aggregation and precipitation processes that are also influenced by the pH and salinity of seawater. The supersaturation of coastal seawater composition can lead to the settling and storage of HA fractions in the form of organo-mineral complexes, e.g., Fe-(oxy)hydroxide minerals, at coastal seawater sites (Hemingway et al., 2019; Moore et al., 2023; Zhang et al., 2023). Therefore, the pH-dependent soil HA fractions could be recognized as DDOM.

## Discussion

In essence, the solid-phase extraction (SPE) method used by Cai et al. (2024) is unable to recover a significant portion of the hydrophilic DOC fraction, achieving only 3–28% recovery (Grasset et al., 2023). Consequently, major DOM fractions, such as pH-dependent soil HA, are selectively excluded from the subsequent mass spectrometry analysis following SPE (Grasset et al., 2023). In particular, the SPE-based DOM extracts collected along the river-to-ocean continuum exhibit an increasing abundance of HMW components (Cai et al., 2024), which may be associated with the rise of autochthonous protein-like and carbohydrate matter, rather than the decline of terrestrial pH-dependent soil HA fractions



throughout the river-to-ocean continuum (Figure 3) (Catalán et al., 2016; Mostofa et al., 2019; Mostofa et al., 2013).

Notably, HA release sequentially degraded organic molecules that were originally bound to HA through photochemical and microbial processes (Mostofa et al., 2013; Amador et al., 1989). It is highly likely that these SPE-based DOM components are optically active (Mostofa et al., 2019; Zhang et al., 2009; Grasset et al., 2023; Mostofa et al., 2013). Differently, the pH-dependent soil HA are optically inactive in terms of fluorescence intensity, which diminishes with decreasing pH and are not detectable through TOC analysis (Yang et al., 2024). The discussion above suggests that the currently proposed DDOM is not substantiated by the considerations of DOM sources.

Finally, the pH-dependent soil/terrestrial HA, tentatively classified as a DDOM fraction, are prevalent in important soil and sediment environments due to their long-term C stabilization and accumulation through organo-mineral complexes (Hemingway et al., 2019; Moore et al., 2023). This process, in turn, contributes to soil stability and health, promoting sustainable agricultural productivity, as well as providing living habitats for various organisms. Furthermore, a portion of pH-dependent terrestrial HA might be one of the key DOM contributors to the long-term C stability in oceanic environments (Catalán et al., 2016). Additionally, a fraction of pH-dependent soil HA could be recognized as DDOM and would also be optically inactive. Undoubtedly, pH-dependent soil HA cannot be extracted by SPE-based methanol solvents due to their insoluble macromolecular and supramolecular nature (Senesi and Loffredo, 1999; Piccolo, 2002; Yang et al., 2024), leaving them uncharacterized at the molecular level. Therefore, a substantial fraction of pH-dependent soil HA, continuously produced from photosynthetically active terrestrial plants, might be classified as

DDOM, i.e., dark matter in the biosphere. This soil HA fraction fundamentally plays a crucial role in soil's structural framework, serving as a primary building block of the soil matrix across the Earth's crust and facilitating C stabilization by forming organo-mineral complexes. Lastly, the connections between DDOM and the significance of sensitivity analysis in the monitoring and management of natural water resources (Errico et al., 2019; Pirone et al., 2024; Lama and Chirico, 2020) should be the focus of further studies.

## Author contributions

KM: Conceptualization, Funding acquisition, Project administration, Resources, Supervision, Validation, Writing—original draft. JY: Formal Analysis, Methodology, Software, Validation, Writing—review and editing. XY: Formal Analysis, Investigation, Methodology, Validation, Writing—review and editing. MM: Formal Analysis, Investigation, Methodology, Validation, Writing—review and editing. C-QL: Resources, Validation, Writing—review and editing. NS: Validation, Writing—original draft, Resources. GS: Validation, Writing—review and editing. DV: Validation, Writing—review and editing. S-LL: Funding acquisition, Validation, Writing—review and editing.

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## Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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