

A novel model of humin

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Our poor knowledge of the structure and formation of humic substances stems from the lack of suitable tools which enable the study of the molecular structure of these complex, yellow-brownish materials. Here, several findings using a combination of isotopic, pyrolytic, microscopic and gas chromatographic methods lead to the proposal of a novel model of humin structure. The formation of humin should thus involve at least three processes: selective preservation of microbial, straight-chain biopolymers; physical encapsulation of apolar substances by weak forces; and chemical binding by covalent bonds.

Unexpected $^{13}\text{C}/^{12}\text{C}$ ratios

In 1995, while analysing various plant and crop soil materials, we were surprised to find out that soil humus is often enriched in carbon 13 relative to plant matter [1]. This was rather unexpected because the intense biodegradation of plant debris in soils should have led to the selective preservation of the most resistant, ^{13}C -depleted lignin and lipid parts of the plant. We therefore proposed two alternative explanations for the occurrence of ^{13}C -enriched humus. First, our results fit well with the reconcondensation mechanism involving condensation of small molecules such as amino acids and sugars [2], which are indeed ^{13}C -enriched relative to the bulk plant carbon. A such pathway has been nicely evidenced in 1917 by Maillard who compared synthetic and natural humic substances [2]. Second, since highly resistant, aliphatic biopolymers have been recently discovered in microbes and sediments [3], we suspected the occurrence of such biopolymers in soils. Indeed, soil microbes feeding on ^{13}C -enriched amino acids and sugars should biosynthesise ^{13}C -enriched biopolymers. Therefore, several analytical means were used to probe for such aliphatic biopolymers in soils, as described below.

Resistant straight-chain biopolymers

In order to test the biopolymer hypothesis, we watched carefully a sample of HF-treated humin by transmission electron microscopy. Resistant aliphatic biopolymer showing typical

very thin laminae (25 μm thick) were indeed present [4], thus strengthening the biopolymer hypothesis. Further pyrolysis of humin followed by gas chromatography coupled to mass spectrometry (GC-MS) revealed the occurrence of C_{11} – C_{24} linear alkanes and alkene doublets [5,6] which are typical breakdown products of aliphatic biopolymers [3]. Therefore, a part of soil organic matter is indeed composed of straight-chain biopolymers (Fig. 1).

Encapsulated plant wax alkanes

On the other hand, we were puzzled by the additional occurrence in the humin pyrolysate of long-chain C_{27} – C_{33} linear alkanes with the typical plant wax fingerprint [5]. Indeed, since such compounds are not formed by pyrolytic cleavage, and since they are still present in the pyrolysate despite pre-extraction of the soil sample and of the humin, they must somehow have been encapsulated in the humin matrix by

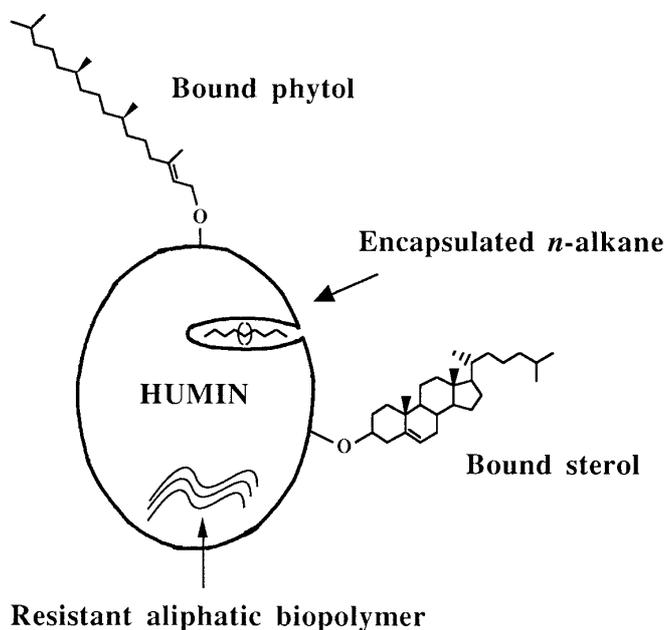


Figure 1. Model of humin explaining the stabilization of soil organic matter 1) by encapsulation of small polar molecules (physical sequestration), 2) by binding of functionalized biomarkers (chemical sequestration), and 3) by selective preservation of aliphatic biopolymers.

non-covalent bonds [7]. However, to confirm this hypothesis, we had to find another way to distinguish "free" soil-extractable substances [8,9] from their humin-bound counterparts. At this point, we thought that trapped molecules should be older than their free counterparts because encapsulation should induce the temporary storage of organic substances. Having previously set up a way to calculate the relative age and turnover of soil molecules by maize labelling [9,10], we measured the $^{13}\text{C}/^{12}\text{C}$ ratio of plant-derived C_{27} – C_{33} linear alkanes both from the soil extract and from the humin pyrolysate. As predicted, the isotopic composition were significantly different, yielding for example $\delta^{13}\text{C}$ values [11] of -28.1 ‰ for the free C_{31} alkane and -29.7 ‰ for the humin-bound homologue [7]. A such difference means that the bound substance is 7 years older than the free substance. These findings clearly demonstrate that apolar organic molecules can be trapped by weak bonds into the humin matrix. Nonetheless, other molecules may also be trapped by strong, covalent bonds, as discussed below.

Biomarkers in humin

Gas chromatography-mass spectrometric analysis of the humin pyrolysate revealed the occurrence of hopanoid and steroid biomarkers [6]. Biomarkers have been widely used to assess the biological sources of dead matter [12]. Here, we found pristene, sterenes, and hopenes in the humin pyrolysate [6]. Pristene is most likely derived from the phytol side chain of chlorophyll. Pristene is thus a marker of photosynthetic activity. Precursors of sterenes include C_{27} algal and C_{29} plant sterols. Hopenes are mainly inherited from bacterial hopanols [13]. Since all those alkenes are products of pyrolytic cleavage, alcohols were most likely linked to humin by ester binding. This finding is in good agreement with the recent studies of the Hamburg group, nicely evidencing the occurrence of ester-bound xenobiotics by Na^{18}OH cleavage of humic compounds [14].

Conclusion

Three processes of humin formation have been assessed using molecular, isotopic and microscopic tools: 1) the selective preservation of microbial, straight-chain biopolymers, 2) the physical encapsulation of apolar substances by weak forces, and 3) the chemical binding by covalent bonds. Noteworthy, the structures of the molecules shown on the

model (Fig. 1) have been unambiguously identified. Moreover, the model does not show links between two or more molecules because, to my best knowledge, such structures have not been identified. Further work at the molecular level is in progress to unravel the humic puzzle. Specifically, the knowledge of bonding forces occurring in humic substances is a prerequisite to understand the fate of xenobiotics in soils, waters and sediments [15].

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