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地质环境中生物标志物 GDGTs 分析技术研究进展

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摘要: 甘油二烷基甘油四醚脂(GDGTs)是一类来自于微生物细胞膜脂的新兴生物标志物, 广泛存在于海洋、湖泊、土壤、泥炭等环境。在活体细胞中, GDGTs 通常以完整极性膜脂(IPL-GDGTs)的形式存在, 而在地质环境中主要以脱去极性头基的核心脂(CL-GDGTs)的形式存在。CL-GDGTs 结构稳定、不易降解, 并且对环境变化响应敏感, 因此被认为是重建古气候-古环境变化的有力工具。GDGTs 结构复杂、种类繁多, 在环境中的含量通常较低且常与其他化合物共存, 因此分析难度较高, 现有技术和方法在其分离、纯化、定量等方面仍然面临挑战。本文总结了近年来 GDGTs 在分析技术方面的研究进展, 概述了 GDGTs 的分类与结构, 对环境中 IPL-GDGTs 和 CL-GDGTs 的分离、纯化等方法进行总结和比较, 其中 CL-GDGTs 可选择多种提取方法, 而极性较强、热稳定性较差的 IPL-GDGTs 应尽量选取 Bligh-Dyer 提取法。普通的分离、纯化通常采用柱层析法, 而涉及 GDGTs 单体分离时, 一般采用制备液相色谱法。液相色谱-质谱、核磁共振波谱、气相色谱-同位素比值质谱是 GDGTs 含量测定、结构鉴定、同位素分析的主要分析手段。本文评述了现有方法的特点和不足, 并在此基础上, 提出了 GDGTs 分析技术的发展方向, 以期对地质环境中 GDGTs 的分析研究提供启示和参考。

关键词: 甘油二烷基甘油四醚脂 (GDGTs); 提取; 分离纯化; 含量分析; 结构鉴定; 同位素分析

要点:

(1) GDGTs 是来自微生物细胞膜脂的一类新兴的生物标志物, 是当前古气候-古环境重建研究的良好载体和有力工具, 在地质环境中广泛分布。

(2) GDGTs 的结构复杂、种类繁多, 分析难度较高, 现有分析技术在其分离、纯化、准确定量等方面仍面临挑战。

(3) GDGTs 的分析研究重点需要关注方法标准化、分析效率、新组分识别及同位素分析等方面。

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生物标志物, 是从生物体中演化而来的一系列有机分子, 其含量、分布特征及同位素组成能够反映地质历史时期的气候和环境变化^[1-2]。甘油二烷基甘油四醚脂(Glycerol dialkyl glyceryl tetraethers, 简称 GDGTs)是古菌或细菌细胞膜的骨架成分, 也是

二十世纪末发展起来的一类新兴的生物标志物^[3]。GDGTs 在自然界广泛分布, 包括海洋^[4-6]、湖泊^[7-10]、河流^[11]、热泉^[12-13]、土壤^[14-17]、泥炭^[18-22]等环境。GDGTs 的组成对外界环境变化响应敏感, 并且其核心脂结构稳定、不易降解, 能够

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长期保存在沉积环境中,因此被认为是古气候重建研究的良好载体和有力工具^[23-28]。

目前, GDGTs 已被应用于重建早白垩纪以来的气候环境演变历史^[29-30]。例如, Chu 等^[31] 利用湖光岩湖泊沉积物中 GDGTs 重建了末次冰消期以来中国热带地区温度变化, 该结果与格陵兰冰芯记录的温度变化趋势大致相同, 证实高纬度冰盖与热带陆地之间存在耦合作用; Bai 等^[32] 利用土壤中 GDGTs 指标重建了青藏高原南部古海拔变化历史, 为揭示印度板块与欧亚大陆板块的碰撞过程和机制提供了新线索。大量研究表明 GDGTs 在古气候重建中发挥了重要作用, 而进行古气候重建的前提是准确分析 GDGTs^[29-32]。然而, GDGTs 分子结构复杂, 其同系物、异构体理化性质相似, 并且环境中 GDGTs 的含量较低(通常在 ng/g 水平), 又常与其他有机化合物共存, 因此分析难度较高, 现有技术在其分离、纯化、准确定量等方面仍面临挑战^[1,3]。

本文在前人研究基础上, 概述了 GDGTs 的结构与分类, 总结并比较了地质环境样品中 GDGTs 的提取、分离、纯化等前处理方法, 评述了液相色谱-质谱、核磁共振波谱、气相色谱-同位素比值质谱等技术在含量分析、结构鉴定、同位素分析方面的研究进展, 同时提出未来 GDGTs 分析技术的发展方向。

1 GDGTs 的结构与分类

环境中的 GDGTs 种类多样, 通常所说的 GDGTs 是指微生物的核心脂(Core Lipids, CL-GDGTs), 其基本结构包括两个烷基长链和两个甘油分子, 烷基长链末端通过醚键与甘油基团结合形成闭合环状大分子^[33-34]。在活体细胞中, CL-GDGTs 两端的甘油基团上带有由磷酸盐、糖基或葡萄糖醛酸等组成的极性头基, 以完整极性膜类脂(Intact Polar Lipids, IPL-GDGTs)的形式存在(图1), 但随着细胞的死亡降解, IPL-GDGTs 会丢失极性头基而转变成 CL-GDGTs^[16,18,35]。

目前, 研究较多的 CL-GDGTs 主要包括两大类: 类异戊二烯 GDGTs(Isoprenoidal GDGTs, 简称 isoGDGTs)和支链 GDGTs(Branched GDGTs, 简称 brGDGTs)。两者的结构既有相似性, 又有明显差异(图1)。相同点在于: 两者均是由2条长链烷基和2个甘油分子形成的四醚类环状化合物, 并且烷基侧链上均含有一定数量的甲基和五元环或六元环。不同点在于: ①两者结构中甘油的立体构型不同, isoGDGTs 为甘油-1-磷酸(G-1-P)构型, 而 brGDGTs

为甘油-3-磷酸(G-3-P)构型; ②两者烷基侧链的骨架不同, isoGDGTs 的烷基侧链含有40个碳原子且为类异戊二烯结构^[12], 而 brGDGTs 的侧链仅由28个碳原子组成且为支链烷烃结构^[6](图1)。

isoGDGTs 主要包括 GDGT-0 到 GDGT-8(数字代表其结构中包含的环戊烷数量)、泉古菌醇(Crenarchaeol, 简称 Cren)及其立体异构体(Crenarchaeol regioisomer, 简称 Cren')(图1)^[12]。isoGDGTs 的母源微生物是古菌, 包括泉古菌(Crenarchaeota)、奇古菌(Thaumarchaea)、广古门菌(Euryarchaeota)等^[36-37]。其中, 泉古菌主要合成含有0~4个环戊烷的 GDGTs 和 Cren, 氢氧化古菌主要合成 Cren 及 Cren', 而含有4个环戊烷以上的 isoGDGTs(GDGT-5 至 GDGT-8)仅在热泉和嗜热菌的培养液中发现^[38]。

brGDGTs 含有 I、II 和 III 三个系列, 每个系列之间相差一个 CH₂。目前已发现 II 和 III 系列烷基侧链上的甲基可以在 C₅、C₆、C₇ 位置变动, 分别形成 5-甲基、6-甲基和 7-甲基等位置异构体^[8,39-41]。在立体结构方面, brGDGTs 的甘油构型只有一种反平行结构^[42], 这与 Cren 及 Cren' 存在平行和反平行 2 种构型不同^[43], 并且其烷基侧链上的多个手性碳原子的立体构型也是固定的^[44]。目前, brGDGTs 的生物来源还不十分明确, 但其甲基支链结构和甘油立体化学特征指示它们由细菌合成^[8], 其稳定碳、氢同位素表明其来自兼性异养细菌^[21,45]。培养实验也证实酸杆菌可以合成 brGDGTs 个别组分(如 Ia、Ic)^[46-47]。最新研究表明, 除了酸杆菌外, 变形菌、消化螺旋菌、拟杆菌、放线菌和疣菌也可能是 brGDGTs 的潜在母源^[25-27,48]。

2 地质环境样品中 GDGTs 的提取方法与纯化技术

2.1 GDGTs 的提取方法

地质环境样品中 GDGTs 的提取方法主要包括 Bligh-Dyer 法(简称 BD 法)^[12,35,49-50]、超声萃取法^[14,32,51-52]、索氏抽提法^[53-56]、加速溶剂萃取法(ASE)^[31,57-59]和微波辅助提取法(MAE)^[21,59-60]等(表1)。

BD 法最早被应用于提取真核生物和细菌中的细胞膜脂, 以氯仿-甲醇(2:1, V/V)为提取液, 后经不断改良, 如增加萃取液中甲醇的比例, 加入少量的酸(如盐酸、三氯乙酸、磷酸盐缓冲液), 使其对脂类物

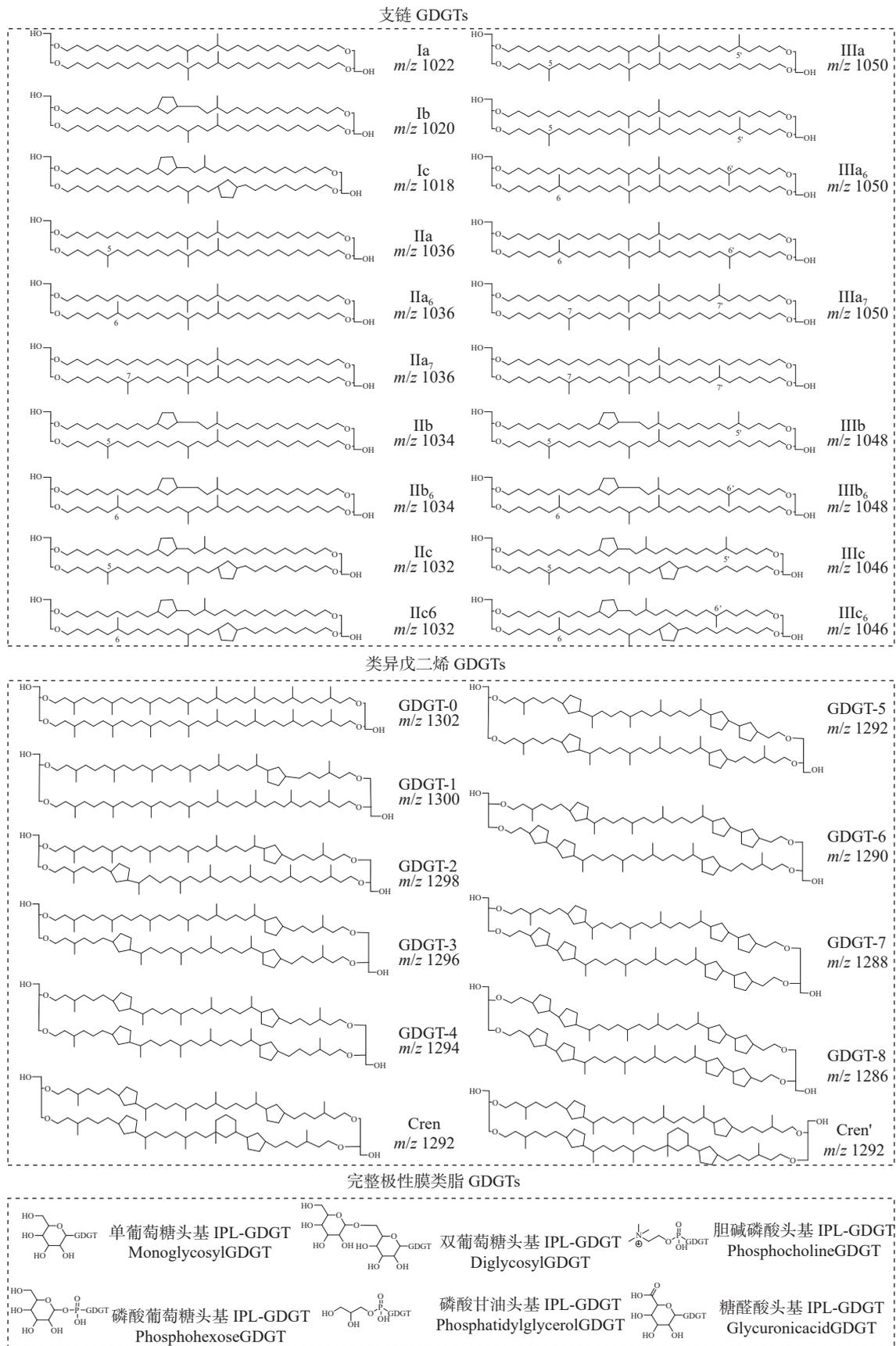


图1 环境中常见的支链 GDGTs、类异戊二烯 GDGTs 和完整极性膜类脂 GDGTs 的分子结构图

Fig. 1 Molecular structures of brGDGTs, isoGDGTs and IPL-GDGTs in the environment.

表 1 地质环境样品中 GDGTs 的不同提取、分离、纯化方法

Table 1 Different extraction, separation and purification methods of GDGTs in geological environment samples.

样品类型	提取方法及提取剂	分离及净化方法	参考文献
土壤	BD 法, 甲醇-二氯甲烷-磷酸缓冲液	硅胶柱, 正己烷/乙酸乙酯、乙酸乙酯、甲醇	Pitcher 等(2009)
土壤	超声萃取法, 甲醇、甲醇-二氯甲烷	硅胶柱, 正己烷/二氯甲烷、二氯甲烷/甲醇	Bai 等(2017)
土壤、湖泊沉积物	BD 法, 甲醇-二氯甲烷-磷酸缓冲液	硅胶柱, 正己烷/乙酸乙酯、甲醇	Buckles 等(2014)
土壤、湖泊沉积物	索氏抽提法, 甲醇-二氯甲烷	Al ₂ O ₃ 柱, 正己烷/二氯甲烷、二氯甲烷/甲醇	Hu 等(2016)
黄土	ASE, 二氯甲烷-甲醇	硅胶柱, 二氯甲烷/乙酸乙酯	Lu 等(2016)
泥炭	超声萃取法, 二氯甲烷、二氯甲烷/甲醇	硅胶柱, 正己烷/甲醇	Zheng 等(2018)
泥炭	MAE, 二氯甲烷-甲醇	硅胶柱, 二氯甲烷/甲醇	Naafs 等(2017)
海洋沉积物	索氏抽提法, 甲醇-二氯甲烷	Al ₂ O ₃ 柱, 正己烷/二氯甲烷、二氯甲烷/甲醇	Liao 等(2020)
海洋沉积物	BD 法, 甲醇-二氯甲烷-磷酸缓冲液	硅胶柱, 正己烷/乙酸乙酯 制备 HPLC, 正己烷/异丙醇	Zhu 等(2013)
湖泊沉积物	ASE 法, 二氯甲烷-甲醇	Al ₂ O ₃ 柱, 二氯甲烷、二氯甲烷/甲醇	Chu 等(2017)
湖泊沉积物	ASE 法, 二氯甲烷-甲醇	Al ₂ O ₃ 柱, 正己烷/二氯甲烷、二氯甲烷/甲醇	Li 等(2023)
湖泊沉积物	ASE 法, 二氯甲烷-甲醇	Al ₂ O ₃ 柱, 二氯甲烷、二氯甲烷/甲醇 制备 HPLC, 正己烷/异丙醇	Weber 等(2015)
海洋、湖泊沉积物	MAE 法, 二氯甲烷-甲醇	Al ₂ O ₃ 柱, 正己烷/二氯甲烷、二氯甲烷/甲醇	Escala 等(2009)
湖泊水体悬浮颗粒物	BD 法, 甲醇-二氯甲烷-磷酸缓冲液	Al ₂ O ₃ 柱, 正己烷/二氯甲烷、二氯甲烷/甲醇	Kumar 等(2019)

质的提取率可达 90% 以上^[3, 59, 61]。超声波萃取法通常采用甲醇、甲醇-二氯甲烷(1 : 1, *V/V*)、二氯甲烷等为提取液, 超声萃取 2 ~ 3 次以保证提取效率^[52, 59]。索氏抽提法通常采用二氯甲烷-甲醇混合溶剂为萃取液, 加热回流提取 24 ~ 72h, 其萃取效率较高, 但溶剂消耗量大且耗时、费力^[53-56]。ASE 法是目前 CL-GDGTs 的主流提取方法, 通常以二氯甲烷-甲醇为提取溶剂, 在温度 100 ~ 120℃、压力 1200 ~ 1500psi 条件下, 静态萃取 5 ~ 10min, 循环萃取 2 ~ 3 次^[57-59]。2020 年 Auderset 等^[62]提出一种 ASE 分步提取法, 通过不同极性溶剂萃取, 从土壤和沉积物中依次分离出正构烷烃、长链烯酮及 GDGTs 三种组分。MAE 法是一种利用微波加热技术对固体样品提取的方法, 通常以二氯甲烷-甲醇为提取溶剂, 在一定温度、压力和微波辐射下, 萃取 10 ~ 20min, 该法操作简单、快速高效、节省溶剂、选择性好, 但由于微波反应器的普及程度不高, 所以目前应用还较少^[28, 59-60]。

不同方法对 GDGTs 的提取效率不尽相同^[50-51, 63-64]。Schouten 等^[51]比较了超声萃取、索氏抽提和 ASE 萃取三种方法对 isoGDGTs 的提取效率, 发现三种方法得到的古气候指标 (TEX₈₆) 值在 ±1σ 误差范围内基本一致。Huguet 等^[59]比较了 BD 法、超声法、索氏抽提法、ASE、MAE 等多种提取方法, 认为索氏提取法对于 IPL-GDGTs 的提取效果最佳, 优于常用的 BD 法。随后, Lengger 等^[63]比较了 BD 法、索氏抽提法和 ASE 法, 发现三种提取方法对

CL-GDGTs 的提取效率基本一致且良好, 但对 IPL-GDGTs 的提取效率则呈现: ASE 法 < 索氏抽提法 < BD 法, 其原因可能是较高的温度 (ASE 法 100℃, 索氏抽提法 65℃) 导致部分 IPL-GDGTs 降解, 而常温下进行的 BD 法更好地保护了 IPL-GDGTs 的极性头基。王欢业等^[50]对比了 BD 法和超声萃取法的提取效率, 指出 BD 法对于 IPL-GDGTs 的提取效果更好, 而超声萃取法对于 CL-GDGTs 的提取效率更高, 但两种方法得到的古气候指标 (TEX₈₆、MBT、CBT) 数值一致, 说明不同提取方法并不影响古气候指标的计算。之后, Yang 等^[64]也证实对于含有极性头基的 OH-GDGTs, 使用 BD 法比超声萃取法的提取效率更高。综上可知, CL-GDGTs 的提取可以选择多种方法, 而对于极性较强、热稳定性较差的 IPL-GDGTs, 应尽量选取 BD 法。

2.2 GDGTs 的分离与纯化方法

在得到总脂类提取物后, 需要从中进一步分离、纯化 GDGTs, 这个过程通常采用柱层析法^[51, 59-61], 少数情况下采用制备液相色谱法^[40, 65-66]。前者适合常规的定量分析, 后者或两者联用适合 GDGTs 的结构鉴定和同位素分析。柱层析法通常采用硅胶、氧化铝 (Al₂O₃) 等吸附剂作为固定相填料^[51, 59-61]。Al₂O₃ 作为固定相时, 通常以正己烷-二氯甲烷、甲醇-二氯甲烷为淋洗液^[31, 51, 57]。硅胶作为固定相时, 通常以正己烷-乙酸乙酯、二氯甲烷-乙酸乙酯等为洗脱液^[58, 66]。收集到的洗脱液通常包括非极性和极性两部分, 其中 GDGTs 存在于极性组分, 将其用氮

气吹干,溶解于合适的溶剂,过滤后即可上机分析^[3,40]。柱层析色谱法的操作性强、分离效果好,但操作繁琐、溶剂消耗量大、容易产生误差。Sanchi等^[67]提出一种采用固相萃取装置进行自动分离、净化的方法,使操作更简单、误差更小、重现性更高。

对于 IPL-GDGTs,其含有的某些极性头基在硅胶柱净化、无水硫酸钠干燥、受热等过程中极易脱落^[63]。为了避免 IPL-GDGTs 在分离、净化过程中损失,Huguet等^[59]提出一种间接测定 IPL-GDGTs 的方法:首先把总脂类提取物平均分为两份,一份用 Al₂O₃ 柱分离、净化,得到 CL-GDGTs 的含量;另一份在酸性条件下水解,使 IPL-GDGTs 丢失极性头基转化为 CL-GDGTs,由此得到的水解产物即是 (IPL+CL)-GDGTs 的总和。最后,用 (IPL+CL)-GDGTs 减去 CL-GDGTs,差值即为 IPL-GDGTs 含量。但需注意这种“差减法”的误差比直接测量法大,尤其当 IPL-GDGTs 的浓度比 CL-GDGTs 低时应谨慎使用^[59]。

制备液相色谱法主要用于分离在柱层析法中不易分离或易损失的组分,如某些 GDGTs 单体、异构体或带有不稳定头基的 IPL-GDGTs^[22,39,66,68]。Weijers等^[18]和 Weber等^[40]利用氨基制备色谱柱,通过制备液相色谱反复分离、纯化,从 brGDGTs 混合物中分离出高纯度、高浓度的 Ib 和 IIIa_{5,6} 组分。Zhu等^[66]采用两根极性正交的半制备色谱柱(一根分离正相体系的二醇基柱,一根分离反相体系的 C₁₈ 柱),通过制备液相色谱,从总脂类提取物中成功分离出 2 种带有糖苷头基的 IPL-GDGTs。

3 地质环境样品中 GDGTs 的分析技术

地质环境中 GDGTs 的分析主要涉及含量分析、结构鉴定、同位素分析三方面,相关分析主要借助于高效液相色谱-质谱仪(HPLC-MS)^[3,51-54]、气相色谱-质谱仪^[3,63,69-70]、核磁共振波谱仪^[18,41,71]和气相色谱-同位素比值质谱仪^[40,65,72]。

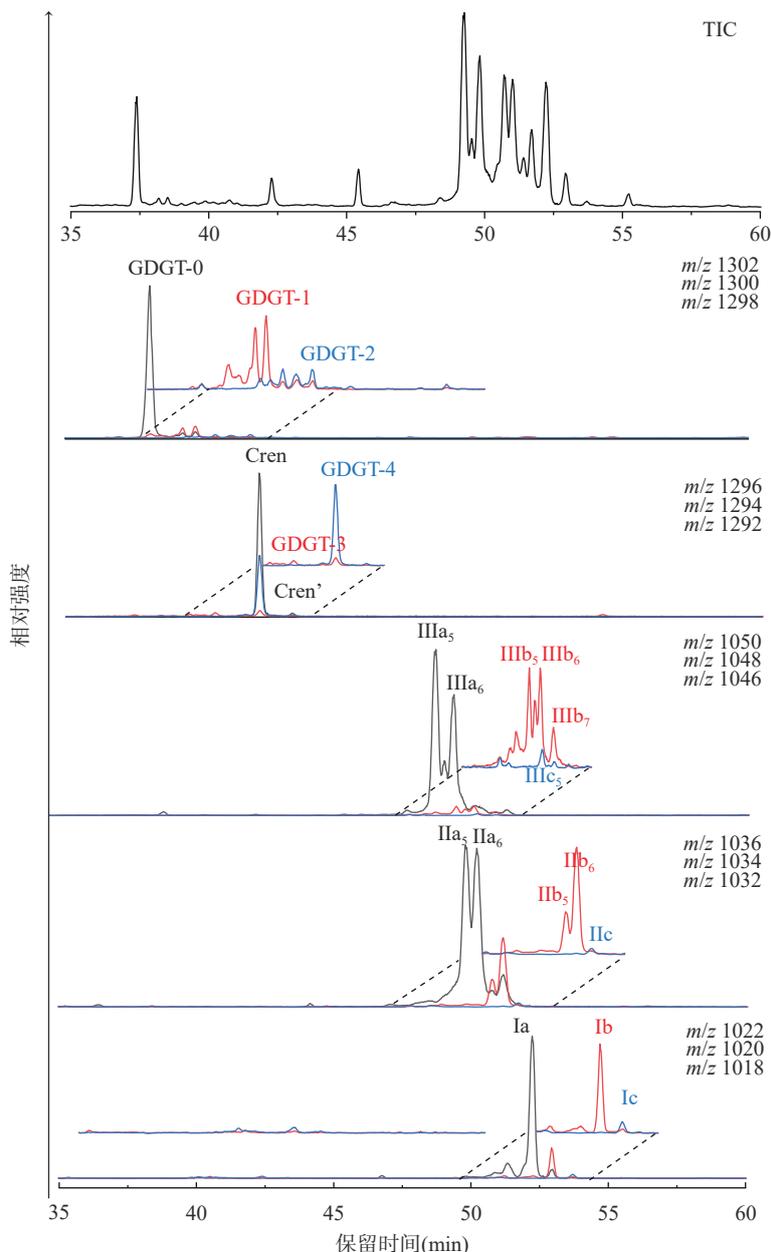
3.1 GDGTs 含量分析

CL-GDGTs 的分析通常采用高效液相色谱-大气压化学电离源-质谱法(HPLC-APCI-MS)。色谱柱一般选择氨基柱、氰基柱、硅胶柱等正相色谱柱,流动相一般选择正己烷-异丙醇^[51,73]或者正己烷-乙酸乙酯^[55,74]。采用 2 根 UPLC 色谱柱或 4 根 HPLC 色谱柱串联,可以使 brGDGTs 的同分异构体(如 III a₅、III a_{5,6}、III a₆ 与 III a₇)达到较好的分离效

果^[8,39-41,73]。在正相液相色谱体系下,GDGTs 各组分的出峰顺序按照其质荷比(m/z)从大到小依次排列(图 2)。isoGDGTs 的 $[M+H]^+$ 范围在 m/z 1286 ~ 1304 之间,出峰顺序通常为:GDGT-0 > GDGT-1 > GDGT-2 > GDGT-3 > GDGT-4 = Cren > Cren'。brGDGTs 的 $[M+H]^+$ 范围在 m/z 1018 ~ 1050 之间,3 个系列的整体出峰顺序为:III > II > I,系列内的出峰顺序为:III a > III b > III c,II a > II b > II c, I a > I b > I c,但相邻系列组分的出峰顺序有时存在交叉的情况^[8]。同分异构体(如 III a、III b)的出峰顺序与其烷基侧链上甲基的位置有关,通常为: C₅ > C₆ > C₇(图 2)。

IPLs-GDGTs 带有极性头基,通常采用反相液相色谱(RP-HPLC)-电喷雾离子源-质谱仪(ESI-MS)分析^[59,66,75]。Zhu等^[66]运用反相超高效液相色谱-四极杆-飞行时间串联质谱,建立了一种可同时分析 IPL-GDGTs 和 CL-GDGTs 的方法,该方法比传统的正相方法(HPLC-APCI-MS)的灵敏度更高、色谱峰形更尖锐,并且能够得到与之一致的古气候指标值。在该方法基础上,Chen等^[75]将 RP-HPLC-ESI-MS 与多反应监测(MRM)技术结合,建立了一种更适合分析 IPL-GDGTs 的方法,其采用的 MRM 模式比常用的选择离子扫描(SIM)模式具有更好的特异性和灵敏度,提高了方法的准确度、稳定性及重现性,并且检出限更低。

在质谱方面,单四极杆质谱和三重四极杆质谱是当前 GDGTs 分析最常用的质谱检测器。定量分析通常在 SIM 模式下进行,该模式通过采集 GDGTs 各组分的特征离子 $[M+H]^+$,使其灵敏度和重现性比全扫模式更好^[51]。但由于四极杆质谱的分辨率不高,SIM 模式会导致 isoGDGTs 和 brGDGTs 产生不同程度(18% ~ 36%)的离子损失,在一定程度上降低了定量结果及古气候指标的准确性^[76]。近年来,功能强大的高分辨率质谱也开始应用于 GDGTs 的分析,如静电场轨道阱质谱(Orbitrap-MS)^[77-78]、傅里叶变换离子回旋共振质谱(FTICR-MS)^[55,79]。高分辨质谱与 HPLC 联用,可以实现样品中 GDGTs 结构和组成的全面表征,并有助于 GDGTs 未知组分的鉴定。在离子源方面,CL-GDGTs 极性较低,所以通常采用常压化学电离源(APCI)。与电喷雾源(ESI)相比,APCI 源的离子产率更高、准分子离子峰的强度更高^[3,51]。近年来,也报道了采用大气压光致电离(APPI 源)成果^[77,79]。APPI 源是一种新电离技术,其电离原理与 APCI 源相似,但其电离能量更低,对弱极性的 CL-GDGTs 的



图中分析条件: 在正己烷-异丙醇梯度洗脱下, 以 2 根 BEH Hilic 色谱柱(150mm×2.1mm×1.7μm)和 1 根硅胶色谱柱(150mm×2.1mm×1.9μm)串联, 经 HPLC-APCI-MS 分析, 实现 GDGTs 各组分良好分离。

Analytical condition: under gradient elution of *n*-hexane/isopropanol, the good separation of GDGTs were achieved by HPLC-APCI-MS method with two BEH Hilic columns (150mm×2.1mm×1.7μm) and one silica column (150mm×2.1mm×1.9μm) in tandem.

图2 内蒙古双沟山天池湖泊沉积物中 GDGTs 的液相色谱总离子流图和提取离子流图

Fig. 2 Total ion chromatogram (TIC) and extracted ion chromatogram (EIC) of GDGTs identified in lake sediment of Lake Shuanggoushan, Inner Mongolia. The chromatogram generated by HPLC-APCI-MS showing the elution order of brGDGTs and isoGDGTs with [M+H]⁺ ions.

电离效果比 APCI 源更好, 因此在未来分析中有很大的潜力 [77, 79]。

目前, GDGTs 定量分析主要采用内标法, 通常以 GTGT-C₄₆ 作为内标 [47], 根据内标与待测组分 [M+H]⁺ 离子的峰面积及相对校正因子来计算

GDGTs 含量。但实际上, C₄₆ 内标与 GDGTs 各组分的理化性质、色谱行为、离子化效率、质谱响应等并不完全相同, 因此根据其计算的 GDGTs 各组分含量并不十分准确。今后, 随着 GDGTs 商用标准物质及同位素内标的出现, 将改善定量分析的准确性。

不同化合物在不同仪器上或同一台仪器在不同时间检测时,会因为仪器的设置参数或状态的变化而导致响应值发生改变,因此开展 GDGTs 分析方法验证十分必要。2009年, Schouten 等^[80]组织了全球 15 个实验室间的比对,采用 HPLC-APCI-MS 法对 2 个海洋沉积物样品进行分析,通过比较 2 种 GDGTs 古气候指标来评估分析方法的准确性和可靠性。比对结果表明, TEX_{86} 指标在实验室内的重复性结果和实验室间的再现性结果均较好,对应的温度误差分别在 1~2℃ 之间和 3~4℃ 之间;而 BIT 指标在同一实验室内的重复性较好,但实验室间的再现性结果不理想,其原因可能是 BIT 指标同时涉及 brGDGTs 和 isoGDGTs,而两者在不同质谱仪上的响应差异较大,导致 BIT 值有较大不同^[80]。2013年,第二轮全球实验室间比对分析表明两种指标的结果均比第一次有所改善, TEX_{86} 指标的再现性温度误差降低至 1.3~3.0℃,与海洋其他古温度指标的误差相近^[81]。但 BIT 指标实验室间的再现性结果依然不理想,这可能与仪器结构、参数设置等有关。Escala 等^[60]和 Schouten 等^[80-81]建议后续可采用已知 BIT 值的样品先对质谱进行校正,再进行 BIT 指标的测定。

3.2 GDGTs 结构鉴定

GDGTs 的结构鉴定通常借助高分辨质谱(HRMS)、气相色谱-质谱(GC-MS)和核磁共振波谱(NMR)^[18, 42, 63, 70]。依据 HRMS 提供的质子化分子离子峰的精确质量数及其同位素峰的相对丰度,可以推测 GDGTs 的分子式^[77, 79]。再根据质谱的一级、二级碎片离子,结合质谱裂解规律,可以推测出 GDGTs 可能分子结构^[41-42]。但是,应用 HRMS 不能揭示同分异构体间的微小结构差异(如 III a₅ 与 III a₆)^[39]。

“醚键裂解法”是将结构复杂的 GDGTs 转化为分子量更小、结构更简单的组分,以便进行更精准的结构解析。该法通过裂解 GDGTs 分子结构中的醚键,将其转化为卤代烃和甘油两部分,再将卤代烃还原为支链烷烃,之后用 GC-MS 对烷烃进行分析。根据特征碎片离子、保留时间、保留指数等信息,结合文献和质谱标准谱库,可以推断出烷烃部分的结构,再结合甘油的构型,最终得到 GDGTs 完整的分子结构^[39-41, 72, 82-83]。在电子轰击电离源下,烷烃的支链节点通常会优先断裂,形成强度较高的特征碎片离子,这一规律可以用来判断烷烃侧链上甲基支链的位置,从而实现 5-甲基、6-甲基等 brGDGTs 位置异

构体的鉴别^[39-41]。上述方法最初用于鉴定 isoGDGTs 的分子结构^[69-70],近来也被用于 brGDGTs 和 OH-GDGTs 等物质的结构解析^[39-41, 82]。

常规气相色谱的工作温度上限在 300~350℃ 之间,难以将 GDGTs 汽化,而高温气相色谱(HT-GC)的工作温度上限达到 400~450℃,可以将 GDGTs 汽化而实现直接分析。Weijers 等^[18]和 Pancost 等^[72]运用配有火焰离子化检测器的 HT-GC,在 140~420℃ 的程序升温下实现 CL-GDGTs 的分析。Sutton 等^[83]利用 HT-GC 结合飞行时间质谱仪,在 20min 内实现 isoGDGTs 的快速分析。

核磁共振氢谱(¹H-NMR)、碳谱(¹³C-NMR)及二维核磁共振技术(如 HSQC、HMBC、COSY)也是解析 GDGTs 结构时常用的分析手段,其提供的大量分子结构信息可与 HPLC-MS、GC-MS 等结果相互验证^[18, 44, 71, 82]。但 NMR 的局限性在于其灵敏度较低且为非选择性分析,故要求测试样品的纯度高(通常>90%),并且待测组分的含量达到 mg 级才能保证信号强度。但是,将 GDGT 单体从结构相似的同系物、异构体中分离、富集并不容易,Liu 等^[82]从 1kg 干重的海洋沉积物中仅分离出 0.6mg 的 GDGT 单体。鉴于此,目前只有个别 GDGTs 组分(如 I a、I b、II a)的结构得到 NMR 验证^[18, 44, 71]。今后,可以尝试应用天然产物、药物代谢等领域常用的 HPLC-NMR 联用技术,简化操作流程,实现更高效、便捷的 GDGTs 结构解析^[84]。

3.3 GDGTs 同位素分析

GDGTs 的同位素组成蕴含着重要的生物和环境信息^[65, 85-90]。在缺氧环境中,可以通过测定沉积物中 isoGDGTs 的 $\delta^{13}C$ 值,揭示是否存在甲烷厌氧氧化古菌^[61]。湖泊沉积物和湖水悬浮颗粒物中 brGDGTs 的 $\delta^{13}C$ 值,可以帮助判断 brGDGTs 的来源及其母源细菌的代谢途径^[87-88]。海洋中 isoGDGTs 的 $\delta^{13}C$ 值可以指示在古海洋生产力的演变历史^[89]。海洋沉积物中 isoGDGTs 的 ¹⁴C 年龄可以揭示其母源微生物的来源^[68],还可以与海洋中其他生物标志物(如浮游有孔虫、长链烯酮)的年龄对比验证^[90-91]。

当前,稳定碳、氢同位素分析主要采用气相色谱-同位素比值质谱(GC-C/TC-IRMS)技术^[40, 61, 65, 87-89]。在分析前, GDGTs 先要经过“醚键裂解”转化为分子量更小、挥发性更强的烷烃,然后再进行 GC-C/TC-IRMS 分析^[61, 69-70, 85]。Pearson 等^[89]利用线轴微燃烧技术建立一种直接测定 isoGDGTs 碳同位素的

方法, 避免了“醚键裂解”过程可能引起的同位素分馏效应, 提高了方法的准确度和精密度(在 1σ 范围内误差仅为 $\pm 0.25\%$), 但该方法由于涉及仪器改装因此并不普及。Lin 等^[86]提出一种测定 IPL-GDGTs 糖苷头基 $\delta^{13}\text{C}$ 的方法, 该法将 IPL-GDGTs 水解得到的糖苷头基衍生、净化, 再由 GC-C-IRMS 测定, 结果表明糖苷头基比其烷烃骨架更富集 $\delta^{13}\text{C}$ (偏移量可达 15%), 这种分子内稳定同位素分测试技术为探索 GDGTs 头基和骨架的代谢提供新的视角和方式。

GDGTs 氢同位素组成可以指示氢的来源、重建降水、海拔等变化趋势, 但目前相关研究较少^[22, 70]。Huguet 等^[22]应用稳定同位素探针技术, 将富含 brGDGTs 的天然泥炭样品在重水和 ^{13}C 标记的碳酸氢钠中培养, 然后利用 IR-GCMS 测定培养后样品 brGDGTs 的 δD 和 $\delta^{13}\text{C}$ 值, 首次量化了泥炭中 brGDGTs 的生产率。Kaneko 等^[70]提出了一种运用 GC-IRMS 测定 isoGDGTs 烷基侧链 δD 值的方法, 该法通过“醚键裂解法”将 isoGDGTs 转化为卤代烃, 利用 H_2/PtO_2 将卤代烃还原为饱和烃, 再通过 GC-IRMS 分析获得氢同位素组成。结果表明, 醚键裂解过程不存在 δD 分馏效应, 而 H_2/PtO_2 氢化过程存在一定程度的 δD 分馏, 但通过质量平衡校正, 可以获得较准确的 δD 值^[70]。2021 年, Lengger 等^[92]利用 HT-GC 结合 IRMS 开发了一种可直接测定 GDGTs 中 δD 值的方法, 该法不需要“醚键裂解”等过程, 操作简单、分析误差小, 但由于 HT-GC 尚未普及, 因此其应用仍然有限。

近年来, 放射性碳同位素技术在 GDGTs 高精度定年、古环境示踪、海洋碳循环等研究中逐渐兴起^[68, 90-91]。放射性碳同位素的分析主要借助加速器质谱仪, 测试样品的最低含碳量约为 $100\mu\text{g C}$ ^[93]。为满足测试要求, 一般需要从大量环境样品中提取、分离、纯化, 才能获取足够量、纯度高的 GDGTs 单体^[18, 90], 如 Haghypour 等^[94]从 3kg 土壤样品中仅提取出 $15\mu\text{g C}$ 。2021 年, Gies 等^[90]提出一种无需分离 GDGTs 单体, 仅按 brGDGTs 和 isoGDGTs 分类即可测定 ^{14}C 的方法, 将初始的取样量降低至 500g,

大幅简化了前处理操作, 并且对于年龄小于 1800y 的样品, 只需 $20\mu\text{g C}$ 的上样量就可以获得准确、精密的 ^{14}C 年龄结果。

4 结语与展望

近二十年来, GDGTs 的分析技术和方法取得长足进步, 这为古气候重建、全球气候变化等研究提供了极大助力。目前, 地质环境中 GDGTs 的样品前处理方法已相对成熟, 以 HPLC-MS、GC-MS、NMR、GC-IRMS 为主的技术手段也成为分析 GDGTs 的重要工具, 但现有方法在分离、纯化、准确定量等方面仍然存在一些问题亟待解决: ①标准化问题。当前已有多种 GDGTs 分析方法, 但尚缺乏测定 GDGTs 的标准化方法和程序, 不同实验室和实验人员使用的前处理方法、仪器方法和数据处理方法的差异, 均会影响结果的比较和重复性的验证。②方法效率问题。现有提取、分离和纯化方法较为复杂, 操作难度较高; HPLC-MS 定量分析用时较长, 分析效率较低。③GDGTs 未知组分问题。环境中仍然存在一些结构未知的 GDGTs 新组分。样品中这些未知组分的存在, 不仅可能影响分析结果的准确性, 还可能给古气候指标带来误差。

因此, 建议今后分析方法的发展应重点考虑以下方面: ①建立 GDGTs 标准分析方法。亟需开展 GDGTs 分析方法标准化研究, 并建立相应的质量控制与评价体系, 从而提高分析结果的可靠性及可比性。②开发便捷、高效的 GDGTs 分析方法。包括: 开发分离性能更好的色谱柱, 以简化现有串连 2~4 根色谱柱的 HPLC 方法; 开发操作更简单、自动化程度更高的提取、分离及净化方法; 运用高精度的 HRMS, 开发更加灵敏、准确的定性及定量方法。③开展环境中 GDGTs 未知组分的识别与结构解析研究。综合运用 HPLC-HRMS、GC-MS、NMR 等技术手段, 进一步识别环境中 GDGTs 未知组分, 从而更好地表征环境中 GDGTs 的分布及组成。④加强 GDGTs 同位素技术方法研究。亟待开发进样量更小、操作更简便的同位素技术及方法, 并拓展其应用。

Research Progress in Analytical Methods of Biomarker GDGTs in Geological Environments

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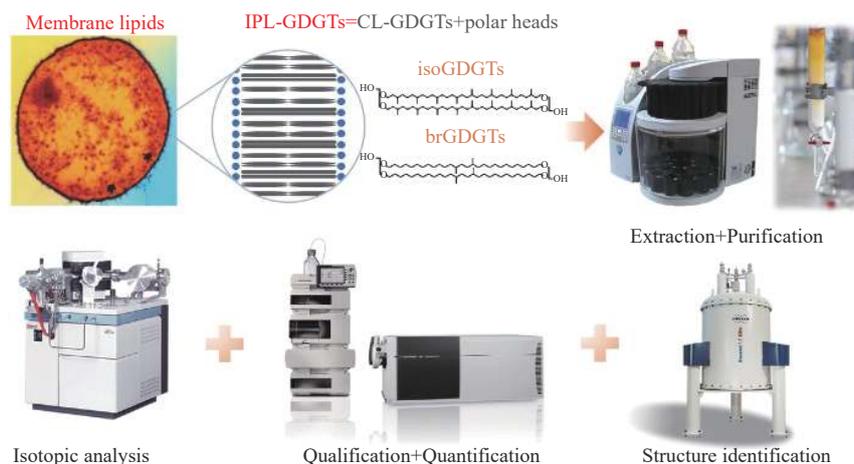
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HIGHLIGHTS

- (1) GDGTs are a new class of biomarkers, which are ubiquitous in geological environments and have unique advantages in paleoclimate reconstruction.
- (2) The analysis of GDGTs is difficult due to their structural diversity, and the existing analytical methods still face challenges in separation, purification, and accurate quantification.
- (3) The future analysis of GDGTs should focus on improving the analytical separation, efficiency, and accuracy and expand to method standardization, new component identification, and isotopic techniques.



ABSTRACT: Glycerol dialkyl glyceryl tetraethers (GDGTs) are a class of environment biomarkers that are widely found in the environment of oceans, lakes, soils, and peat. GDGTs usually exist as intact polar lipids (IPL-GDGTs) in living cells, while they exist as core lipids stripped of polar head groups (CL-GDGTs) in geological environments. CL-GDGTs are structurally stable and sensitive to environmental changes and are considered to be a powerful tool for reconstructing palaeoclimate-palaeoenvironmental changes. GDGTs are structurally complex and diverse, coexisting with other compounds and present low contents, which brings challenges in analysis, especially in separation, purification, and quantification. This article summarizes the classification and structure of GDGTs, and presents a summary and comparison of methods for the separation and purification of IPL-GDGTs and CL-GDGTs in the environment. Multiple extraction methods can be used for CL-GDGTs, while the polar and thermally unstable IPL-GDGTs are preferably extracted using the Bligh-Dyer method. This article reviews the characteristics and limitations of various analysis methods, including liquid chromatography-mass spectrometry, nuclear magnetic resonance spectroscopy, and gas chromatography-isotope ratio mass spectrometry. The BRIEF REPORT is available for this paper at <http://www.ykcs.ac.cn/en/article/doi/10.15898/j.ykcs.202306100077>.

KEY WORDS: glycerol dialkyl glycerol tetraethers (GDGTs); extraction; separation and purification; content analysis; structural identification; isotopic analysis

BRIEF REPORT

Glycerol dialkyl glycerol tetraethers (GDGTs) are lipid biomarkers that are widely distributed in geological environments, including oceans^[4-6], lakes^[7-10], river estuaries^[11], hot springs^[12], soil^[14-17], and peats^[18-22]. They are sensitive to environmental changes and can effectively record paleoclimate information over a range of geological time scales. GDGT-based proxies are widely used in the reconstruction of terrestrial and marine paleoenvironments^[23-28]. However, the accurate analysis of GDGTs is challenging due to their diverse nature and difficulties in their separation, purification, and quantification.

GDGTs in the environment exist in two forms: intact polar lipids (IPL-GDGTs) or as core lipids (CL-GDGTs) (Fig. 1). CL-GDGTs can be further classified into two types based on their structural differences: isoprenoid GDGTs (isoGDGTs) and branched GDGTs (brGDGTs). IsoGDGTs are primarily composed of GDGT-0 to GDGT-8, *Crenarchaeol*, and its stereoisomer. IsoGDGTs are produced by archaea^[36-37], while the biological origin of brGDGTs is still uncertain, although there is evidence to indicate that their potential producers are heterotrophic bacteria, including *Acidobacteria*, *Proteobacteria*, *Nitrospira*, *Bacteroidetes*, *Actinobacteria*, and *Verrucomicrobia*^[25-27, 48]. BrGDGTs consist of three series (I , II , and III), each of which differs by a methylene group. The alkyl side chains of brGDGTs in series II and III may possess various methyl groups at the C₅, C₆, and C₇ positions, leading to the formation of positional isomers, such as 5-methyl, 6-methyl, and 7-methyl brGDGTs^[8, 39-41].

Extraction methods for GDGTs from environmental samples mainly involve the Bligh-Dyer method^[12, 35, 49-50], ultrasonic extraction^[14, 32, 51-52], Soxhlet extraction^[53-56], accelerated solvent extraction (ASE)^[31, 57-59] and microwave-assisted extraction (MAE)^[27, 59-60]. The Bligh-Dyer method was originally used to extract cell membrane lipids from eukaryotes and bacteria, but modifications have been made to increase extraction rates^[3, 59, 61]. Ultrasonic extraction usually uses methanol, methanol-dichloromethane, and dichloromethane as the extraction solution^[52, 59]. Soxhlet extraction is highly efficient but demands a large amount of solvents and is time-consuming^[53-56]. ASE is the most commonly used method for extracting CL-GDGTs, using dichloromethane-methanol as the extraction solvent^[57-59]. MAE is a simple, fast, efficient, and selective method for solid samples, but its application is limited by the lack of widespread microwave reactor availability^[28, 59-60].

Different GDGTs extraction methods have been compared in several studies^[50-51, 63-64]. Schouten et al.^[51] compared the efficiency of three extraction methods: ultrasonic extraction, Soxhlet extraction, and ASE extraction for isoGDGTs. They found that the TEX₈₆ values obtained by the three methods were almost identical within $\pm 1\sigma$ error. Wang et al.^[50] also compared the efficiency of Bligh-Dyer and ultrasonic extraction. They found that the Bligh-Dyer method was more effective in extracting IPL-GDGTs, whereas the ultrasonic extraction method was more efficient in extracting CL-GDGTs. However, the values of palaeoclimatic indicators (TEX₈₆, MBT, and CBT) obtained by both methods were identical, indicating that the different extraction methods did not affect the value of palaeoclimatic proxies^[50]. Yang et al.^[64] later confirmed that the Bligh-Dyer method was more efficient than the ultrasonic extraction in extracting OH-GDGTs that contain polar head groups. In summary, the Bligh-Dyer method is preferable for IPL-GDGTs due to their greater polarity and lower thermal stability, while various extraction methods can be used for CL-GDGTs.

To isolate and purify GDGTs from environmental samples, column chromatography^[51, 59-61] is commonly used along with preparative liquid chromatography (Prep HPLC) in some cases^[40, 65-66]. Prep HPLC is mainly used to separate components that are hard to fractionate or prone to loss in column chromatography, such as certain GDGT

monomers, isomers, or IPL-GDGTs with unstable head groups^[22, 39, 66, 68]. The Ib and IIIa_{5,6} components were successfully isolated from mixtures of brGDGTs through Prep HPLC along with repeated separation and purification^[18, 40]. Silica and alumina (Al₂O₃) are commonly used as stationary phases. The elution process yields nonpolar and polar fractions. GDGTs are usually found in the polar fraction and can be further analysed for structural identification and isotopic analysis using Prep HPLC or a combination of both methods. During the process of separating and purifying GDGTs, some IPL-GDGTs may lose their polar head groups due to poor stability. To address this issue, Huguet et al.^[59] proposed an indirect method for determining IPL-GDGTs. The total lipid extract was divided into two equal parts. The first part was purified through an Al₂O₃ column to obtain the content of CL-GDGTs. The second part underwent acid-hydrolysis, which converted IPL-GDGTs into CL-GDGTs by losing their polar head groups. The hydrolysis product was the sum of (IPL+CL)-GDGTs. The quantity of IPL-GDGTs was determined by subtracting CL-GDGTs from (IPL+CL)-GDGTs. However, the error of this “subtraction method” is greater than that of the direct measurement. Therefore, this method should be used with caution, especially when the concentration of IPL-GDGTs is lower than that of CL-GDGTs^[59].

The analysis of GDGTs in the geological environment mainly involves content determination, structural identification, and isotopic analysis. This is commonly done using high-performance liquid chromatography-mass spectrometry (HPLC-MS)^[3, 51-54], gas chromatography-mass spectrometry (GC-MS)^[3, 63, 69-70], nuclear magnetic resonance spectroscopy (NMR)^[18, 41, 71], and gas chromatography-isotope ratio mass spectrometry (GC-IRMS)^[40, 65, 72]. CL-GDGTs are usually analyzed by HPLC-APCI-MS, while IPLs-GDGTs with polar head groups are usually analyzed by RP-HPLC-ESI-MS. Good separation of 5-, 6-, and 7-methyl isomers of brGDGTs can be achieved by using four HPLC columns or two UPLC columns in tandem^[8, 39-41, 73]. In a normal-phase HPLC system, the peaks of GDGTs are sorted by their mass-to-charge ratios (m/z) from the largest to the smallest, as shown in Fig. 2.

Single and triple quadrupole mass spectrometry are commonly used for the GDGTs analysis through HPLC-MS. SIM mode is preferred for quantification due to its higher sensitivity and reproducibility than the full-scan mode, which captures the characteristic ions $[M+H]^+$ of each GDGT component^[51]. However, low resolution can cause ion loss for isoGDGTs and brGDGTs, which affects the accuracy of quantitative results^[76]. Recently, high-resolution mass spectrometry, such as Orbitrap-MS^[77-78] and FTICR-MS^[55, 79], has been applied to the GDGTs analysis. HPLC-HRMS provides a comprehensive characterization of the structure and composition of GDGTs in environmental samples, offering great potential for future analyses.

Structure identification of GDGTs involves various techniques such as HRMS, GC-MS, and NMR techniques^[18, 42, 63, 70]. The molecular formulae of GDGTs can be determined by HRMS through precise mass measurement of the protonated molecular ion peaks and isotopic peaks evaluation^[77, 79]. Conventional GC has a temperature limit of 300-350°C, which makes it impossible to vaporize GDGTs directly. However, HT-GC has a higher limit of 400-450°C, allowing for direct analysis of GDGTs. HT-GC equipped with a flame ionization detector can be used to analyse CL-GDGTs^[18, 72]. The possible molecular structures of GDGTs can be deduced from analyzing the primary and secondary fragment ions of the mass spectra, along with their fragmentation pathways^[41-42]. NMR techniques provide molecular structural information of GDGTs, which can be verified by other methods such as HPLC-MS and GC-MS^[18, 41, 71, 82]. Nevertheless, NMR has certain limitations and requires high sample purity and content to ensure adequate signal intensity. NMR analysis usually requires high sample purity and milligram injection quantities, which means it is not easy to separate and enrich GDGT monomers from their homologues and isomers, so only a few GDGT components (e.g., Ia, Ib, and IIa) have been verified by NMR^[18, 44, 71]. In the future, HPLC-NMR technology may simplify the process and achieve more efficient and convenient structural analysis of GDGTs^[84].

The isotopic composition of GDGTs provides vital biological and environmental information [65, 85-90]. GC-IRMS is the primary method used for stable carbon and hydrogen isotope analysis of GDGTs [40, 61, 65, 87-89]. Prior to GC-IRMS analysis, GDGTs are converted into smaller and more volatile alkanes through “ether bond cleavage” [61, 69-70, 85]. Accelerator mass spectrometry is used to analyze radiocarbon isotopes for GDGTs, but it requires a carbon content of the sample greater than 100 $\mu\text{g C}$ [93]. To meet this testing requirements, a large amount of environmental samples must be extracted, separated, and purified to obtain an adequate amount of high-purity individual GDGT component [18, 90].

Significant progress has been achieved in the analytical techniques and methods for GDGTs during the last two decades. However, existing approaches still face issues regarding standardization, method efficiency, and novel GDGT components. Future studies should focus on the following aspects: (1) Establishing standard methods and quality control systems for GDGTs. Developing standard analytical procedures for GDGTs and establishing a quality control and evaluation system is crucial to improve the reliability and comparability of the analytical results. (2) Developing convenient and efficient analytical methods. It is necessary to establish simpler and more automated techniques for GDGT extraction, separation, and purification. In addition, developing qualitative and quantitative methods with better sensitivity and accuracy is also important. (3) Identification of unknown components of GDGTs. Using various techniques such as HPLC-HRMS, GC-MS, and NMR to identify the unknown GDGT components, enhancing the understanding of their composition and distribution in the environment. (4) Developing new isotopic techniques and methods for GDGTs with fewer samples and easier procedures, and helping to expand the use of isotopic methods in the future.

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