An improved FT-TRA system and its application to element analysis of foraminifera

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Element and isotope composition of foraminiferal shells are potentially valuable paleo-proxy for paleoceanography research. However, the standard hand-on batch method for elemental and isotopic analysis often results in substantial loss of sample, as well as contamination by secondary calcite and clay, thus biasing the data toward a more variable and anomalous result. These potentially significant sources of error can be overcome with a flow-through time-resolved (FT-TRA) sequential leaching method that makes time- and labor-intensive pretreatments unnecessary. In this study, we developed an accessible, multifaced FT-TRA module which allows a wide variety of applications can be developed, by customizing key system parameters such as sample cell types. It also offers the choice between on-line analysis or fraction collection for off-line. Also, a kit suite for online addition of internal standards is adopted to improve the accuracy and precision. In addition, a specialized software for FT-TRA data was developed to improve efficiency. To verify the reliability and accuracy of our FT-TRA system, systematic analysis was carried out, including dissolution of foraminifera and standard sample GSR-6 (argillaceous limestone). The result of GSR-6 showed that the reproducibility of Mg/Ca was better than 6 % (1 RSD, n=3), and Sr/Ca was better than 2.1 % (1 RSD, n=3). Moreover, four foraminifera from the same sample were analyzed respectively, and the result showed that the reproducibility of Mg/Ca was better than 8.3 % (1 RSD, n=4), and Sr/Ca was better than 2.9 % (1 RSD, n=4). Furthermore, with gradually increasing and highly regulated acid strength, FT-TRA reliably separated the biogenic calcite (the desire signal) and clay (a contaminant contains Al). The results of the foraminifera also show that FT-TRA technology is highly sensitive, and Mg/Ca, Sr/Ca and Mn/Ca can be obtained with a single foraminiferal shell. However, for rare earth elements and U, the results suggest that larger sample sizes are needed to enhance precision and retrieve reliable results.

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