

U-238, Th-232 series, and Pu-239 + Pu-240 concentration analysis in biological samples of high natural background radiation residents by alpha spectrometry

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Abstract

The purpose of this research was to determine the isotope concentration of the actinide radioelements ²³⁸U, ²³²Th series, and ²³⁹Pu in biological samples (urine) using the alpha spectroscopy system. The samples were selected from the residence of high natural background radiation (HNBR) area and analyzed using chemical procedure. The separated sample was placed on a membrane filter by the sedimentation method, and it was prepared for counting with the alpha spectrometry system. The average radioactive concentrations found in these samples were as follows: ²³⁴U > ²²⁶Ra > ²³⁰Th > ²²⁸Ra > ²²⁸Th > ²²⁴Ra > ²³²Th > ²²⁴Ra > ²³²Th > ²²⁴Ra > ²³²Th > ²³²Th > ²³²Th > ²³⁴U and ²²⁶Ra were much greater than the other radionuclides.

Keywords Actinide · Alpha spectroscopy · Biological samples · High natural background radiation (HNBR) area

Introduction

Uranium and thorium are radioactive substances that can be found in nature, like soil [1–3], sediment [4, 5], water [6–8], and building materials [9–11]. The U and Th isotopes typically are employed in various tracer investigations to assess their influence on the environment and geology. The major three isotopes of natural uranium are ²³⁸U ($T_{1/2}$ =4.468×10⁹

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a), 235 U (T_{1/2}=704×10⁶ a), and 234 U (T_{1/2}=2.455×10⁵ a), all of which are alpha (α) emitters [12]. Thorium has three primary isotopes, all of which are alpha emitters: 232 Th (T_{1/2}=1.402×10¹⁰ a), 230 Th (T_{1/2}=7.54×10⁴ a) [13], and 228 Th (T_{1/2}=1.9126 a) [14].

Higher levels of radioactivity in the environment are connected with increased radiation damage and risk to humans, as evidenced by kidney damage, mutagenicity, bladder and kidney cancer, leukaemia, testis cancer, and lung cancer [15–17]. There are several residential locations on the earth with high natural background radiation (HNBR), such as Guarapari in Brazil, Yangjiang in China, Kerala in India, and Ramsar in Iran. In HNBRs regions like Ramsar, the yearly effective dosage rate can approach 260 mSv a⁻¹[2].

Uranium and thorium transfer from one environmental compartment to another in biological systems such as vegetables are dependent on the availability of nuclides in the soil as well as the rate of loss from the interior structure of leaves following translocation. This translocation can add the U and Th to the human food chain. Studies on the HNBR area are necessary because clearer findings about the true risks of radiation should obtained and reasonable levels of radiation protection promoted. Although radiobiological and epidemiological studies in HNBR locations have yet to demonstrate any indication of a substantial increase in health damage compared to that in normal background areas, it is necessary to quantify the U and Th concentrations in biological samples [18], particularly if the linear threshold model is used for dose assessments.

Alpha spectrometry could be used to determine the radioactivity of alpha emitter nuclide in environmental samples. Examples include actinide determination on soil samples and analysis of water samples downstream of uranium mining operations for uranium series radionuclides. The technique is also used for internal dosimetry for workers using the analysis of biological samples such as urine [19].

In recent years, natural radionuclides (U and Th) series determination methods in biological samples have been successfully used in various studies around the world [20–27]. There are different methods for the determination of Th series radionuclides in biological samples, but alpha spectrometry is the most accurate and low minimum detection level (MDL) in comparison with other methods like gamma spectrometry and ICP-mass for short-lived radionuclides. However, in the determination of long-lived ²³⁴U, ²³⁵U, and ²³⁸U, the minimum detection level (MDL) of ICP-MS is better than alpha spectrometry.

This study aimed to determine uranium, thorium, radium, and polonium isotopes in biological samples to provide some basic data on residents living in the HNBR area. For these purposes, the activity concentrations of the most important α -emitters, ²³⁸U series (²³⁴U, ²³⁰Th, ²²⁶Ra, and ²¹⁸Po) and ²³²Th series (²³²Th, ²²⁸Th, ²²⁸Ra, and ²²⁴Ra) were measured in biological samples collected in Ramsar, Iran.

Materials and methods

Study area

Ramsar is located at $36^{\circ} 53' \text{ N}$, $50^{\circ} 41' \text{ E}$, and has an average elevation of 20 m a.s.l. This city is situated in the Caspian Sea's southern region. Ramesar area has a high natural background radiation (HNBR) area in the Talesh Mahaleh, Ramak, Sadat Mahale, and Chaparsar districts (Fig. 1) [4]. In 2016, the Ramsar area was home to a population of 80,000 residents. However, only approximately 20,000 of them chose to reside in the HNBR zones within the area.

Samples preparation

The sample preparation includes sampling, dissolution, radiochemical separation, and sample counting [2]. Four urine samples from residency families with ages more than 50 years, each with a volume of 500 ml, were taken from the residence of the HNBR area. After that, 100 ml urine samples were selected to separate possess of each element. ²³²U ($T_{1/2}=68.9$ a), ²²⁹Th ($T_{1/2}=7880$ a), ²⁴²Pu ($T_{1/2}=3.73 \times 10^5$ a) and ²²⁵Ra ($T_{1/2}=14.9$ d) [13] tracer were added to three blank samples and urine samples as a tracer to measure the

separation efficiency. The blank samples were analyzed parallel with urine samples to calculate the difference recovery percentage between blank samples and urine samples. So, there was no significant separation among the alpha-emitting isotopes from the different elements. Also, the concentration of all tracers was corrected for the date of application. In this process, ammonium hydrogen phosphate solution (3 ml 1.3 mol l^{-1}) and 1 ml 1.25 mol l^{-1} calcium nitrate (50 mg Ca) were added to the solution. In the next step, by adding concentrated ammonium hydroxide and creating an alkaline environment ($pH \sim 9$), the actinides in the urine were precipitated as calcium phosphate. The samples were centrifuged at 3000 rpm for ~5 min. The precipitate was centrifuged again at 3000 rpm for about 5 min after being washed once with approximately 20-30 ml of distilled water, and the supernatant was discarded. The obtained sediment, which contained iron phosphate and actinides, was dissolved in concentrated nitric acid (HNO₃ 10 ml 6 mol 1⁻¹) and 10 ml aluminum nitrate $(Al(NO_3)_3 2 \text{ mol } l^{-1})$ solution. The obtained solution was evaporated. Actinide elements were oxidised by adding $0.5 \text{ ml } 1.5 \text{ mol } 1^{-1}$ sodium nitrite (NaNO₂) in an acidic environment. In the oxidation process, the actinides of Pu(III) and U(IV) increased to Pu (IV) and U(VI), respectively.

In the next step, the separation of actinides using a chromatography column containing exchange anionic resin (Dowex 8×1) was carried out. In this process, Th, Pu, and U actinides were separated individually from anionic resin and collected in a separate container. After pre-conditioning the resins with 10 ml of 8 mol 1^{-1} HNO₃, the sample is loaded onto the columns at a flow rate of $\sim 1 \text{ ml min}^{-1}$ to extract and separate Th, Pu, U and Ra. The columns are rinsed with ~ 10 ml of 8 mol 1^{-1} HNO₃ until the effluent turns colorless. The resin in the column was removed, divided into portions, and reloaded into new columns. The resin was blended and then divided equally for individual U, Th Pu and Ra determinations. In addition, the resin fraction was equivalent to about 30 ml of urine. The resin column is rinsed with 10 ml of 12 mol l⁻¹ HCl to remove Th, and then the Pu isotopes are eluted into a 50 ml centrifuge tube using 15 ml of 2 mol l^{-1} HNO₃. The U isotopes are stripped from the resin column with 1.0 ml of 4 mol l^{-1} HCl followed by 15 ml of 0.01 mol l^{-1} HCl. Consequently, the Ra was eluted in 5 ml of 3 mol 1^{-1} HNO₃.

However, the preparation of a good quality α -source which should be thin, weightless, uniform, and homogenous on a suitable flat and smooth substrate, is one of the most important prerequisites to obtain a good quality α -spectrum with high resolution, and a small low energy tail contribution [28]. The method used in this research was to create sediment (5 µm) through deposition of sediment by gravity sedimentation. Th and U as Phosphate and Pu as Fluoride sediment were co-precipitated [29, 30]. It was obtained uniformly on the membrane filters. Actinide filters were counted

Fig. 1 The location of biological sampling in high natural background radiation (HNBR)

area



by alpha spectrometry for around 16 h; however, depending on the required detection limit, shorter count durations of 1 h can also be achieved for emergency response samples utilizing higher-level tracers [31]. Furthermore, the measurement of ²²⁸Ra concentration was conducted to assess the secular equilibrium between ²²⁸Ra and ²²⁸Ac.

Samples counting

The Alpha spectrometry system (ALPHAQUATTRO spectroscopy, SILENA, Italy) was applied with a multi-channel analyzer alpha spectroscopy workstation. It incorporates four independent high-quality modular vacuum chambers equipped with high-performance silicon ion implanted charged particle detectors, their associated electronics, a multichannel pulse-height analyzer, and a test pulser [32]. The effective cross-section of the detector was 450 mm² with an ineffective thickness of 100 μ m. The detector has a resolution of approximately 30 keV (FWHM) and a counting efficiency of 20% for a 5 mm source-detector distance calibrated for 3 to 8 MeV energy. Genie 2000 software was applied for data acquisition and count processing. The counting time was 16 h for each sample, and the working voltage of the detector was + 20 V with 1024 channels. In addition, the uncertainty components to calculate the activity concentration combined uncertainty are presented in Table 1. The system was calibrated for energy, resolution, and counting effectiveness using a standard electrodeposited mixed source

Table 1 Combined standard uncertainties for activity measurements

Uncertainty components	Combined standard uncertainty of sample BS-1 in percentage unit (%)								
	²³⁴ U	²³⁰ Th	²²⁶ Ra	²³² Th	²²⁸ Th	²²⁸ Ra	²²⁴ Ra		
Sample Counts	2	1.3	1.9	5	7	5	5		
Blank Counts	1	1	1	1	1	1	1		
Activity of Tracer	5	5	5	5	5	5	5		
Decay Probability Correction	1	1	1	1	1	1	1		
Sample mass	2	2	2	2	2	2	2		
Time	0.1	0.1	0.1	0.1	0.1	0.1	0.1		
u _c	11.1	10.4	12.0	14.1	16.1	14.1	14.1		

of ²⁴¹Am, ²³⁹Pu, and ²⁴⁴Cm to ensure accurate identification and quantification of alpha-emitting radionuclides. In addition, the sample counting sources and the standard calibration source were physically and chemically identical to ensure the accuracy of the results. The uncertainty of the source activities was 4% at the 95% confidence level.

The activity concentration was calculated by applying the following equation:

$$A_{238} = \frac{\frac{C_{238}V_{232}A_{232}I_{232}}{I_{238}(c_{232}-c_{228})} - \frac{C_{238B}V_{232B}A_{232}I_{232}}{I_{238}(c_{232B}-c_{228B})}}{V}$$
(1)

where C is counts, V is the tracer volume (1), A is the tracer special activity (Bq/l), I is the decay probability, and V is the sample volume (1). The B index was related to the blank sample.

The minimum detection limit (MDL) was determined using the following equation:

$$MDL\left[\frac{Bq}{g}\right]_{238} = \frac{2.71 + 4.66 \times \sigma_B}{T_B \times \epsilon \times m_B \times 2.22}$$
(2)

where T_B is blank counting time (s), ε is measurement system efficiency, m_B is mass of blank sample (g), and σ_B is:

$$\sigma_{BLK} = \frac{\sqrt{C_{238B}} \times m_{238B} \times A_{232} \times I_{232}}{I_{238} \times (C_{232B} - C_{228B})}$$
(3)

where σ_B estimated from the standard deviation of multiple blank measurements, and other parameters were explained in the above section.

The ²²⁸Ra radionuclide was measured by HPGe gamma spectrometry system according to the provirus published paper explained method. The gamma lines of 911.2 keV and 969.0 keV emitted from ²²⁸Ac radionuclide were applied to activity calculation (See the method [10]).

Interferences in the spectrum

The major spectrum interferences for measuring ²²⁴Ra and ²²⁶Ra are predicted to be naturally existing radionuclides in biological samples, such as $^{228}\mathrm{Th},\,^{230}\mathrm{Th},\,\mathrm{and}\,\,^{234}\mathrm{U},\,\mathrm{as}$ well as radium isotope progeny nuclides (²²⁶Ra, and ²²⁴Ra). Dai et al. [33] report revealed that Th and U distinguish in the urine samples process for ²²⁶Ra, employing chromatographic separation stages using coupled cation/anion exchange columns to ensure efficient Th and U interferences eradication. Because the current process used the same cation/anion exchange separation strategy, the spectrum interferences of 230 Th/ 234 U on 226 Ra and 228 Th on 224 Ra were negligible.

Because radium isotopes (²²⁶Ra and ²²⁴Ra) cannot be chemically separated from each other, possible spectrum interferences between these isotopes and their offspring nuclides must be carefully examined to choose adequate regions of interest (ROI) for calculating radium activity. For the 226 Ra peak at 4.78 MeV and the 224 Ra peak at 5.68 MeV in biological samples, high-energy resolution of the FWHM (full-width at half maximum) ranging from 10 to 30 keV was indicated.

Results and discussion

Activity concentration

The activity concentration value of the ²³⁴U, ²³⁰Th, ²²⁶Ra, ²²⁸Th, ²²⁸Ra, ²²⁴Ra and ²³⁹Pu measured in biological (Urine) samples and sample codes area is given in Table 2. The region of interest area technique was applied to measure low-activity concentration radionuclides. The minimum and maximum value of ²³⁴U, ²³⁰Th, ²²⁶Ra, ²³²Th, ²²⁸Th, ²²⁸Ra, 224 Ra and 239 Pu were obtained 8011 to 1079 mBq l⁻¹, 379 to 681 mBq 1⁻¹, 4815 to 8830 mBq 1⁻¹, 4.98 to 7.25 mBq 1⁻¹, 6.27 to 13.15 mBg l^{-1} , 7.18 to 12.17 mBg l^{-1} , 4.25 to 8.95 mBq l^{-1} , and < MDL, respectively. The order of average concentrations of radionuclides recorded in these samples was as follows; ${}^{234}U > {}^{226}Ra > {}^{230}Th > {}^{228}Ra > {}^{228}Th$ $>^{224}$ Ra $>^{232}$ Th $>^{239}$ Pu. 234 U and 226 Ra were significantly higher than the other measured radionuclides, while in the case of ²³⁹Pu, the average concentrations were lower than MDL (= 5 mBq l^{-1}). Table 3 presents the average ratios of ²²⁴Ra/²²⁶Ra, ²²⁸Ra/²²⁶Ra, ²³⁰Th/²³²Th, and ²²⁸Th/²³²Th in biological samples were calculated. The higher average value

Sample	²³⁸ U Series (mBq l ⁻¹)			232Th Series	²³⁹ Pu (mBq l ⁻¹)			
	²³⁴ U	²³⁰ Th	²²⁶ Ra	²³² Th	²²⁸ Th	²²⁸ Ra	²²⁴ Ra	
BS-1	8621±187	438 ± 4	7620 ± 144	5.17 ± 0.01	9.28 ± 0.03	7.18 ± 0.02	8.09 ± 0.01	< MDL *
BS-2	8011 ± 174	379 <u>+</u> 4	4815 ± 91	5.12 ± 0.01	7.42 ± 0.02	12.01 ± 0.03	4.96 ± 0.01	< MDL *
BS-3	9172±199	491 ± 5	6971 ± 131	4.98 ± 0.01	6.27 ± 0.02	12.17 ± 0.03	4.25 ± 0.01	< MDL *
BS-4	$10,079 \pm 219$	681 <u>+</u> 7	8830 ± 166	7.25 ± 0.02	13.15 ± 0.04	10.24 ± 0.02	8.95 ± 0.02	< MDL *
Mean	6720 ± 146	497 ± 5	7059 ± 133	5.63 ± 0.01	9.03 ± 0.03	10.4 ± 0.02	6.56 ± 0.01	_
Min–Max	8011-10079	379–681	4815-8830	4.98–7.25	6.27–13.15	7.18–12.17	4.25-8.95	_

* MDL=5 mBq l^{-1}

was the ²³⁰Th/²³²Th ratio, which shows the difference concentration value of the ²²⁸U series (²³⁰Th) and ²³²Th series. The lower average value was obtained at the ²²⁴Ra/²²⁶Ra ratio that ²²⁴Ra is from the ²³²Th series with low concentration and ²²⁶Ra is from the ²²⁸U series with high concentration in study samples. As per some investigation in the study area, ²³⁸U daughter (²²⁶Ra) concentration was reported to be higher than the worldwide average [34–36].

Table 3 indicated the ratios of ²³⁸U daughters/²³⁸U and ²³²Th daughters/²³²Th to show disequilibrium in the ²³⁸U and ²³²Th decay chains. As demonstrated in Table 3, the average equilibrium ratio in the ²³²Th series and its progeny was found to be greater than one, whereas the equilibrium ratio in the ²³⁸U series and its progeny was less than one. Those results show that the radionuclide retention and excretion biokinetic results from this study are consistent with the ICRP models [37].

The comparison of the measured activity concentration of all radionuclides showed that the activity concentration of 234 U was higher than other radionuclides. The activity concentration of 226 Ra in spring water was reported to be more than 146,000 mBq L⁻¹ in the study area [34]. Therefore, 226 Ra with a high concentration of water can enter the human organs through the food chain.

The activity concentration percentage of ²³⁸U series (²³⁴U, ²²⁸Th, ²²⁶Ra, and ²²⁴Ra) and ²³²Th series (²³²Th, ²²⁸Th, ²²⁸Ra, and ²²⁴Ra) radionuclides in biological samples are presented in Fig. 2-a and Fig. 2b, respectively. As shown in Fig. 2a,

more than 95% of activity concentration was related to two elements of ²³⁴U and ²²⁶Ra. Whiles in the ²³²Th series, the activity concentration percentage was distributed between 15 to 40% in measured radionuclides (Fig. 2b).

Statistical interpretation

To estimate, the correlations between the activity concentration of ²²⁶Ra-²³⁴U, ²²⁶Ra-²²⁸Ra, and ²³²Th-²²⁴Ra, ²³²Th-²²⁸Th, ²³²Th-²³⁰Th in biological samples, Spearman correlation analysis was applied. This kind of investigation is beneficial for determining whether there are any probable links between observed activity concentrations of all primordial radionuclides. According to Spearman's correlation, a strong positive correlation was observed between ²³⁴U and 226 Ra (r=0.977). The correlation coefficient between two sets of variables was determined, and the significance level was decided depending on the sample size. A highly significant correlation was with p < 0.023, whereas, p > 0.05 was not a significant correlation. For the highly significant correlations regression line (with $R^2 = 95.47\%$) and the prediction were plotted in Fig. 3a. While there is no positive correlation between them and ²²⁸Ra and ²²⁶Ra. This correlation is negative (r = -0.501) with a *p*-value of more than 0.05 (*p* = 0.488) and an R-square value of $R^2 = 26.18\%$. The regression and prediction were plotted in Fig. 3b.

Figure 4a shows the relation between ²³²Th concentrations and ²²⁴Ra as a linear model plot. As can be seen from

Table 3 The activity concentration ratio of 224 Ra/ 226 Ra, 228 Ra/ 226 Ra, 230 Th/ 232 Th, 228 Th/ 232 Th; and 230 Th/ 234 U, 226 Ra/ 234 U, 228 Ra/ 232 Th and 224 Ra/ 232 Th in biological samples of study area

Sample	Radionuclide ratios from different chains			Radionuclide ratios from the same chain					
	²²⁴ Ra/ ²²⁶ Ra	²²⁸ Ra/ ²²⁶ Ra	²³⁰ Th/ ²³² Th	²²⁸ Th/ ²³² Th	230Th/234U	²²⁶ Ra/ ²³⁴ U	²²⁸ Ra/ ²³² Th	²²⁴ Ra/ ²³² Th	
BS-1	0.00106 ± 0.00002	0.00123 ± 0.00005	84.72 ± 0.06	1.79 ± 0.07	0.0508 ± 0.0005	0.8839 ± 0.0006	1.39 ± 0.07	1.56 ± 0.08	
BS-2	0.00103 ± 0.00002	0.00151 ± 0.00007	74.02 ± 0.05	1.45 ± 0.04	0.0473 ± 0.0004	0.6010 ± 0.0005	2.34 ± 0.12	0.97 ± 0.05	
BS-3	0.00061 ± 0.00001	0.00093 ± 0.00003	98.59 ± 0.08	1.26 ± 0.03	0.0535 ± 0.0005	0.7600 ± 0.0006	2.44 ± 0.16	0.85 ± 0.04	
BS-4	0.00101 ± 0.00002	0.00154 ± 0.00004	93.93 ± 0.06	1.81 ± 0.08	0.0676 ± 0.0007	0.8761 ± 0.0006	1.41 ± 0.08	1.23 ± 0.06	
Mean	0.00093 ± 0.00002	0.00135 ± 0.00005	87.82 ± 0.06	1.58 ± 0.06	0.0548 ± 0.0005	0.7803 ± 0.0005	1.89 ± 0.09	1.15 ± 0.05	

Fig. 2 The activity concentration percentage of a^{238} U series (²³⁴U, ²³⁰Th, ²²⁶Ra and ²¹⁸Po) and b^{232} Th series (²³²Th, ²²⁸Ra and ²²⁴Ra) in biological samples



the Figure, our results show a direct significant relationship between ²²⁴Ra levels and ²³²Th (r = 0.963, p-value < 0.037). The R-square value (92.77%) and predication plot are shown a strong correlation between ²³²Th and ²²⁴Ra concentration. These results are also not surprising since both radionuclides are part of the same decay series.

The narrow width of the prediction plot shows this relation (see Fig. 4b). This correlation was expected due to the sameness of the radionuclides in the series and the absence of gaseous elements between these two radionuclides in the decay chain.

The correlation between 232 and 230 Th concentrations fitted in the liner model and presented in Fig. 4-c. As can be seen, the measured concentration indicates that there is no significant relation between 232 and 230 Th. A *p*-value of

more than 0.05 (0.395) and r = 65% indicate this weak relation. The correlation between radionuclides is presented in Table 4.

Within the ²³⁸U and ²³²Th decay chains, there exist certain parent and daughter relationships that exhibit relatively weak correlations. Furthermore, there are some surprisingly strong and unexpected correlations between isotopes across the ²³⁸U and ²³²Th chains. These discrepant relationships are generally explained by invoking the rationale of chemical effects within the human body and by local geochemical fractionation processes. The alternative rationale for the presence or lack of correlations among the radionuclides within and across the ²³⁸U and ²³²Th decay chains is due to the limited statistical strength of the study with only n=4test donors. With these interpretations, we conclude that the



Fig. 3 The Spearman's correlation and prediction plot of a^{234} U and 226 Ra, b^{228} Ra and 226 Ra



Fig. 4 The Spearman's correlation and prediction plot of a^{224} Ra and 232 Th, b^{228} Th and 232 Th, c^{230} Th and 232 Th



Relationship between Th-230 and Th-232

Fig. 4 (continued)

Table 4The concentrationcorrelation between measuredradionuclides

	²³⁴ U	²³⁰ Th	²²⁶ Ra	²³² Th	²²⁸ Th	²²⁸ Ra	²²⁴ Ra
²³⁴ U	1.000						
²³⁰ Th	0.891	1.000					
²²⁶ Ra	0.977	0.821	1.000				
²³² Th	0.897	0.395	0.904	1.000			
²²⁸ Th	0.818	0.471	0.845	0.987	1.000		
²²⁸ Ra	-0.330	0.039	-0.501	-0.545	-0.632	1.000	
²²⁴ Ra	0.746	0.373	0.766	0.963	0.991	-0.607	1.000

importance of the first scenario (biokenetik) was more than that of the second scenario (limited statistical strength). The strong correlations found in Figs. 4 and 5 indicate, that the results in Table 3 indicate a significant correlation. In addition, the cluster similarity diagram of ²³⁴U, ²²⁸Th, ²²⁶Ra, ²²⁴Ra, ²³²Th, ²²⁸Th, ²²⁸Ra, ²²⁴Ra and ²³⁹Pu are presented in Fig. 5. The high similarity in concentration trend was observed in two groups of radionuclides. Those radionuclide groups are indicated in Fig. 5.

In addition, Fig. 5 and Table 4 results indicate that ²²⁸Ra was more correlated with the ²³⁸U decay chain than the ²³²Th chain. Because of chemical reactions within the human body

and local geochemical fractionation, this phenomenon can be explained.

Conclusion

In this study, ²³⁸U series (²³⁴U, ²³⁰Th, and ²²⁶Ra), ²³²Th series (²³²Th, ²²⁸Th, ²²⁸Ra, and ²²⁴Ra), and ²³⁹Pu of HNBR (Ramsar) were analyzed in biological samples (urine) by radiochemical separation followed by alpha-spectrometry spectra. The significant concentration of ²³⁴U and ²²⁶Ra





was measured in biological samples (urine), while ²³⁹Pu concentration was not determined and the concentration level was less than MDL. Also, the activity concentration ratio of ²²⁴Ra/²²⁶Ra, ²²⁸Ra/²²⁶Ra, ²³⁰Th/²³²Th and ²²⁸Th/²³²Th was calculated. The significant ratio value was obtained for ²³⁰Th/²³²Th radionuclides. Overall, the analysis of activity concentration ratios of radionuclides in decay series provides valuable information about the processes controlling their distribution in the environment, their mobility, and the timescales involved in their radioactive decay. The statistical analysis was carried out to determine the correlation between radioelements. The correlation results indicate that the radionuclide retention and excretion biokinetic ICRP model is the dominant process of being in the same group in a radioactive decay series. According to the findings of this results living in areas with high natural radiation background (HNRB) can pose unique health and safety considerations for residents. Residents who are living in these areas are suggested to undergo periodic health checks.

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Author contributions AA: experiments, data processing, original draft, and manuscript preparation. FM: lab analysis, manuscript revision. AWA: methodology, software, HMHZ: methodology, software, writing—original draft preparation, writing—review and editing.

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Declarations

Conflict of interest The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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